

Retroelement Family	database	#databas e	Description	Scientific Name	Max Score	Total Score	Query Cover	E value	Per. Ident	Acc. Len	Accession
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8756	2,99E+08	100%	0.0	100.00%	301019445	NC_029977.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8695	1,86E+08	100%	0.0	99.77%	215701946	NC_029980.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold519	<i>Capisicum annuum</i>	8650	9,15E+03	100%	0.0	99.60%	1101621	NW_015961578.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8634	2,26E+08	100%	0.0	99.47%	261510930	NC_029979.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2298	<i>Capisicum annuum</i>	8632	9,16E+03	100%	0.0	99.52%	1494726	NW_015960454.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8628	1,53E+08	100%	0.0	99.52%	238794889	NC_029985.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8626	1,77E+08	100%	0.0	99.51%	205736368	NC_029986.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8595	1,33E+08	100%	0.0	99.37%	217274494	NC_029981.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8565	1,59E+08	100%	0.0	99.26%	222112641	NC_029983.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8564	1,59E+08	100%	0.0	99.22%	163962470	NC_029978.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8560	1,49E+08	100%	0.0	99.16%	219521584	NC_029982.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1246	<i>Capisicum annuum</i>	8545	2,67E+04	100%	0.0	99.18%	3388803	NW_015960834.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8536	1,23E+08	100%	0.0	99.14%	220335243	NC_029987.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8536	1,66E+08	100%	0.0	99.06%	229934170	NC_029988.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capisicum annuum</i>	8527	8,99E+03	100%	0.0	99.14%	2397553	NW_0159613181.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold181	<i>Capisicum annuum</i>	8523	8,99E+03	100%	0.0	99.03%	2703691	NW_015961338.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold367	<i>Capisicum annuum</i>	8501	9,99E+03	100%	0.0	99.07%	625659	NW_015960847.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2615	<i>Capisicum annuum</i>	8482	9,04E+03	100%	0.0	98.85%	131126	NW_015961557.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8479	8,66E+04	100%	0.0	98.99%	153299543	NC_029984.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2635	<i>Capisicum annuum</i>	8471	1,74E+04	100%	0.0	98.93%	699991	NW_015960547.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold90	<i>Capisicum annuum</i>	8468	9,74E+03	100%	0.0	98.95%	6074945	NW_015960458.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1037	<i>Capisicum annuum</i>	8397	8,87E+03	100%	0.0	98.50%	2186250	NW_015960681.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1191	<i>Capisicum annuum</i>	8272	9,18E+03	100%	0.0	99.01%	145056	NW_015961591.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold301	<i>Capisicum annuum</i>	8248	9,14E+03	100%	0.0	99.80%	1226190	NW_015960697.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4537	<i>Capisicum annuum</i>	8242	9,16E+03	100%	0.0	99.78%	176915	NW_015961775.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold891	<i>Capisicum annuum</i>	8237	1,54E+04	100%	0.0	99.76%	1431046	NW_015960469.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold448	<i>Capisicum annuum</i>	8231	9,19E+03	99%	0.0	99.73%	1323857	NW_015960958.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2719	<i>Capisicum annuum</i>	8187	9,02E+03	99%	0.0	99.64%	277434	NW_015960900.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capisicum annuum</i>	8155	9,04E+03	99%	0.0	99.73%	2470984	NW_015961229.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1874	<i>Capisicum annuum</i>	8150	8,99E+03	99%	0.0	99.66%	719064	NW_015961147.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1360	<i>Capisicum annuum</i>	7974	8,83E+03	96%	0.0	99.72%	360446	NW_015960954.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1399	<i>Capisicum annuum</i>	7904	9,35E+03	100%	0.0	99.86%	1742971	NW_015961630.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold7798	<i>Capisicum annuum</i>	7808	9,30E+03	100%	0.0	99.95%	106903	NW_015961464.1
CopiaCa_01	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2138	<i>Capisicum annuum</i>	6879	7,15E+03	82%	0.0	99.84%	681955	NW_015960370.1
CopiaCa_02	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	10098	1,32E+04	100%	0.0	99.80%	219521584	NC_029982.1
CopiaCa_02	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5062	1,99E+04	100%	0.0	95.23%	261510930	NC_029979.1
CopiaCa_02	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4202	1,23E+04	97%	0.0	95.58%	163962470	NC_029978.1
CopiaCa_02	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	3742	9,12E+03	99%	0.0	92.77%	217274494	NC_029981.1
CopiaCa_02	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	3674	9,57E+03	99%	0.0	94.22%	222112641	NC_029983.1
CopiaCa_02	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	3336	1,56E+04	100%	0.0	94.25%	205736368	NC_029986.1
CopiaCa_02	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	2196	7,83E+03	97%	0.0	87.09%	229934170	NC_029988.1
CopiaCa_03	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	17411	3,23E+08	100%	0.0	100.00%	238794889	NC_029985.1
CopiaCa_03	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	13958	2,98E+08	100%	0.0	93.42%	261510930	NC_029979.1
CopiaCa_03	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	13919	3,24E+08	100%	0.0	93.40%	220335243	NC_029987.1
CopiaCa_03	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	13767	2,72E+08	100%	0.0	93.08%	222112641	NC_029983.1
CopiaCa_03	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	13586	3,48E+08	100%	0.0	92.71%	215701946	NC_029980.1
CopiaCa_03	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	13311	2,43E+08	100%	0.0	92.25%	163962470	NC_029987.1
CopiaCa_03	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	13018	2,20E+08	100%	0.0	91.71%	205736368	NC_029986.1
CopiaCa_03	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold16	<i>Capisicum annuum</i>	11276	1,52E+04	99%	0.0	91.44%	2023789	NW_015960959.1
CopiaCa_03	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold177	<i>Capisicum annuum</i>	7998	1,71E+04	97%	0.0	92.74%	442035	NW_015961759.1
CopiaCa_03	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7949	3,38E+08	100%	0.0	84.22%	301019445	NC_029977.1
CopiaCa_03	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7834	1,21E+08	100%	0.0	83.99%	153299543	NC_029984.1
CopiaCa_03	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7762	2,94E+08	100%	0.0	83.79%	229934170	NC_029988.1
CopiaCa_03	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold376	<i>Capisicum annuum</i>	7158	1,99E+04	100%	0.0	91.89%	776404	NW_015961666.1

CopiaCa_03	refseq_genomes	8	Capisicum annum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	6900	2,80E+08	100%	0.0	84.07%	217274494	NC_029981.1
CopiaCa_03	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold660	<i>Capisicum annuum</i>	5607	7,90E+03	87%	0.0	84.63%	663463	NW_015961299.1
CopiaCa_03	refseq_genomes	8	Capisicum annum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5382	2,83E+08	100%	0.0	84.32%	219521584	NC_029982.1
CopiaCa_03	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold723	<i>Capisicum annuum</i>	2671	5,44E+03	82%	0.0	80.03%	559742	NW_015961191.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	10196	4,94E+08	100%	0.0	99.98%	301019445	NC_029977.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7714	2,62E+08	100%	0.0	91.97%	238794889	NC_029985.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7659	2,51E+08	100%	0.0	91.74%	215701946	NC_029980.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7651	2,74E+08	100%	0.0	91.69%	220335243	NC_029987.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1281	<i>Capisicum annuum</i>	7644	9,64E+03	100%	0.0	91.69%	233328	NW_015961028.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7618	2,33E+08	100%	0.0	91.61%	153299543	NC_029984.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7601	3,29E+08	100%	0.0	91.53%	229934170	NC_029988.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7590	4,08E+08	100%	0.0	91.58%	219521584	NC_029982.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7574	1,69E+08	100%	0.0	91.42%	205736368	NC_029986.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7524	2,95E+08	100%	0.0	91.28%	217274494	NC_029981.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7463	1,77E+08	100%	0.0	91.13%	163962470	NC_029978.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7437	3,67E+08	100%	0.0	91.04%	261510930	NC_029979.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2348	<i>Capisicum annuum</i>	7380	9,27E+03	100%	0.0	90.84%	373378	NW_015960596.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1037	<i>Capisicum annuum</i>	7354	9,24E+03	100%	0.0	90.76%	2186250	NW_015960681.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7341	2,44E+08	100%	0.0	90.73%	222112641	NC_029983.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1422	<i>Capisicum annuum</i>	7252	3,06E+04	100%	0.0	90.71%	802096	NW_015960747.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1633	<i>Capisicum annuum</i>	7160	8,89E+03	99%	0.0	90.38%	1620172	NW_015961041.1
CopiaCa_04	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold592	<i>Capisicum annuum</i>	7068	1,79E+04	100%	0.0	89.84%	1905997	NW_015960988.1
CopiaCa_04	refseq_genomes	8	Solanum pennellii chromosome 11, SPENNv200	<i>Solanum pennellii</i>	6737	5,77E+04	100%	0.0	88.74%	66223686	NC_028647.1
CopiaCa_04	refseq_genomes	8	Solanum pennellii chromosome 4, SPENNv200	<i>Solanum pennellii</i>	6730	1,07E+08	100%	0.0	88.71%	77197300	NC_028640.1
CopiaCa_04	refseq_genomes	8	Solanum pennellii chromosome 12, SPENNv200	<i>Solanum pennellii</i>	6724	1,43E+08	100%	0.0	88.70%	83305730	NC_028648.1
CopiaCa_04	refseq_genomes	8	Solanum pennellii chromosome 6, SPENNv200	<i>Solanum pennellii</i>	6719	4,67E+04	100%	0.0	88.67%	60730942	NC_028642.1
CopiaCa_04	refseq_genomes	8	Solanum pennellii chromosome 1, SPENNv200	<i>Solanum pennellii</i>	6717	1,71E+08	100%	0.0	88.68%	109333515	NC_028637.1
CopiaCa_04	refseq_genomes	8	Solanum pennellii chromosome 3, SPENNv200	<i>Solanum pennellii</i>	6717	9,82E+04	100%	0.0	88.68%	75414019	NC_028639.1
CopiaCa_04	refseq_genomes	8	Solanum pennellii chromosome 10, SPENNv200	<i>Solanum pennellii</i>	6715	1,08E+08	100%	0.0	88.66%	82529941	NC_028646.1
CopiaCa_04	refseq_genomes	8	Solanum pennellii chromosome 5, SPENNv200	<i>Solanum pennellii</i>	6709	1,82E+08	100%	0.0	88.64%	77991103	NC_028641.1
CopiaCa_04	refseq_genomes	8	Solanum pennellii chromosome 2, SPENNv200	<i>Solanum pennellii</i>	6697	5,72E+04	100%	0.0	88.61%	59803892	NC_028638.1
CopiaCa_04	refseq_genomes	8	Solanum pennellii chromosome 8, SPENNv200	<i>Solanum pennellii</i>	6691	1,00E+08	100%	0.0	88.59%	70546659	NC_028644.1
CopiaCa_04	refseq_genomes	8	Solanum pennellii chromosome 9, SPENNv200	<i>Solanum pennellii</i>	6691	7,40E+04	100%	0.0	88.59%	84057508	NC_028645.1
CopiaCa_04	refseq_genomes	8	Solanum pennellii chromosome 7, SPENNv200	<i>Solanum pennellii</i>	6602	5,64E+04	100%	0.0	88.31%	79292169	NC_028643.1
CopiaCa_04	refseq_genomes	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1390	<i>Capisicum annuum</i>	6553	8,02E+03	100%	0.0	88.15%	1730405	NW_015960783.1
CopiaCa_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	<i>Solanum lycopersicum</i>	6540	5,85E+04	100%	0.0	88.12%	56597135	NC_015448.3
CopiaCa_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	<i>Solanum lycopersicum</i>	6516	8,95E+04	100%	0.0	88.01%	72906345	NC_015446.3
CopiaCa_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	<i>Solanum lycopersicum</i>	6466	1,33E+08	100%	0.0	87.97%	65633393	NC_015447.3
CopiaCa_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	<i>Solanum lycopersicum</i>	6438	6,28E+04	100%	0.0	87.74%	68175699	NC_015444.3
CopiaCa_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	<i>Solanum lycopersicum</i>	6423	1,00E+08	100%	0.0	87.83%	66723567	NC_015442.3
CopiaCa_04	refseq_genomes	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold688	<i>Capisicum annuum</i>	6408	9,01E+03	96%	0.0	91.86%	1322070	NW_015960950.1
CopiaCa_04	refseq_genomes	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9	<i>Capisicum annuum</i>	6407	1,33E+04	99%	0.0	87.72%	6567951	NW_015961815.1
CopiaCa_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	<i>Solanum lycopersicum</i>	6405	1,76E+08	100%	0.0	87.68%	98455869	NC_015438.3
CopiaCa_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	<i>Solanum lycopersicum</i>	6372	6,39E+04	100%	0.0	87.63%	66557038	NC_015441.3
CopiaCa_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	<i>Solanum lycopersicum</i>	6370	1,09E+08	100%	0.0	87.56%	68126176	NC_015449.3
CopiaCa_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	<i>Solanum lycopersicum</i>	6368	1,05E+08	100%	0.0	87.67%	65987440	NC_015445.3
CopiaCa_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	<i>Solanum lycopersicum</i>	6277	4,38E+04	100%	0.0	87.45%	55977580	NC_015439.3
CopiaCa_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	<i>Solanum lycopersicum</i>	6277	7,76E+04	100%	0.0	87.39%	72290146	NC_015440.3
CopiaCa_04	refseq_genomes	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold171	<i>Capisicum annuum</i>	6035	7,03E+03	99%	0.0	87.30%	887051	NW_015960473.1
CopiaCa_04	refseq_genomes	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3844	<i>Capisicum annuum</i>	5923	5,92E+03	98%	0.0	86.56%	179318	NW_015961259.1
CopiaCa_04	refseq_genomes	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1341	<i>Capisicum annuum</i>	5875	5,88E+03	99%	0.0	86.08%	1906221	NW_015960665.1
CopiaCa_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	<i>Solanum lycopersicum</i>	5845	4,24E+04	100%	0.0	88.44%	49794276	NC_015443.3
CopiaCa_04	refseq_genomes	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1124	<i>Capisicum annuum</i>	5633	1,07E+04	99%	0.0	85.63%	2110277	NW_015961370.1
CopiaCa_04	refseq_genomes	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold148	<i>Capisicum annuum</i>	5439	6,87E+03	99%	0.0	88.78%	1323090	NW_015961290.1
CopiaCa_04	refseq_genomes	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capisicum annuum</i>	5361	1,91E+04	100%	0.0	90.83%	2397553	NW_015961318.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00015	<i>Solanum tuberosum</i>	5024	5,02E+03	92%	0.0	84.56%	3327864	NW_006238941.1

CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00029	<i>Solanum tuberosum</i>	5018	5,02E+03	82%	0.0	86.55%	2785282	NW_006238955.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00069	<i>Solanum tuberosum</i>	4898	4,96E+03	87%	0.0	85.29%	2089258	NW_006238995.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1973	<i>Capsicum annuum</i>	4859	6,60E+03	100%	0.0	87.22%	1714326	NW_015961422.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00022	<i>Solanum tuberosum</i>	4804	4,80E+03	87%	0.0	84.90%	3032639	NW_006238948.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00120	<i>Solanum tuberosum</i>	4758	4,76E+03	87%	0.0	84.69%	1603502	NW_006239047.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00384	<i>Solanum tuberosum</i>	4710	4,71E+03	81%	0.0	85.77%	565346	NW_006239308.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold6868	<i>Capsicum annuum</i>	4695	5,26E+03	89%	0.0	86.89%	22397	NW_015961342.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00089	<i>Solanum tuberosum</i>	4671	1,14E+04	97%	0.0	84.46%	1856049	NW_006239012.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00589	<i>Solanum tuberosum</i>	4540	4,54E+03	83%	0.0	84.57%	277750	NW_006239511.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1378	<i>Capsicum annuum</i>	4518	9,81E+03	98%	0.0	89.77%	1057095	NW_015961130.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1688	<i>Capsicum annuum</i>	4512	9,06E+03	100%	0.0	89.89%	693367	NW_015961726.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00083	<i>Solanum tuberosum</i>	4503	4,50E+03	80%	0.0	85.28%	1836497	NW_006239015.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1780	<i>Capsicum annuum</i>	4495	4,87E+03	97%	0.0	84.03%	226878	NW_015961280.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4977	<i>Capsicum annuum</i>	4348	7,82E+03	97%	0.0	90.27%	543020	NW_015961055.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold756	<i>Capsicum annuum</i>	4298	8,06E+03	99%	0.0	86.70%	884456	NW_015961254.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00009	<i>Solanum tuberosum</i>	4231	2,22E+04	82%	0.0	86.25%	3955149	NW_006238935.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3027	<i>Capsicum annuum</i>	4080	9,26E+03	98%	0.0	91.73%	193869	NW_015961172.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2239	<i>Capsicum annuum</i>	3978	4,89E+03	93%	0.0	84.42%	615095	NW_015960816.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00086	<i>Solanum tuberosum</i>	3679	6,99E+03	86%	0.0	87.79%	1873685	NW_006239010.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold301	<i>Capsicum annuum</i>	3620	6,71E+03	99%	0.0	86.68%	1226190	NW_015960697.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00606	<i>Solanum tuberosum</i>	3504	4,69E+03	82%	0.0	83.17%	265855	NW_006239521.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold258	<i>Capsicum annuum</i>	3404	1,58E+04	100%	0.0	83.85%	3098259	NW_015961180.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00457	<i>Solanum tuberosum</i>	3339	3,34E+03	84%	0.0	80.10%	449598	NW_006239380.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold619	<i>Capsicum annuum</i>	3317	4,70E+03	83%	0.0	79.89%	2512642	NW_015960498.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold179	<i>Capsicum annuum</i>	3282	5,41E+03	85%	0.0	89.27%	1172749	NW_015961157.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1795	<i>Capsicum annuum</i>	3282	7,53E+03	86%	0.0	91.38%	162617	NW_015961207.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3478	<i>Capsicum annuum</i>	3236	7,81E+03	86%	0.0	85.10%	211087	NW_015961510.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold969	<i>Capsicum annuum</i>	3229	5,60E+03	88%	0.0	86.36%	1557263	NW_015961740.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3037	<i>Capsicum annuum</i>	3142	5,79E+03	91%	0.0	87.17%	1979607	NW_015960688.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1741	<i>Capsicum annuum</i>	3103	4,67E+03	86%	0.0	85.65%	1897442	NW_015960687.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00137	<i>Solanum tuberosum</i>	2952	4,70E+03	83%	0.0	87.24%	1472262	NW_006239066.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3079	<i>Capsicum annuum</i>	2918	7,77E+03	97%	0.0	88.36%	299956	NW_015961555.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00165	<i>Solanum tuberosum</i>	2870	4,41E+03	98%	0.0	82.86%	1320774	NW_006239090.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00471	<i>Solanum tuberosum</i>	2761	7,68E+03	87%	0.0	83.63%	432374	NW_006239389.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00033	<i>Solanum tuberosum</i>	2497	1,04E+04	82%	0.0	81.90%	2695005	NW_006238959.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00036	<i>Solanum tuberosum</i>	2481	1,30E+04	85%	0.0	85.98%	2637368	NW_006238960.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold536	<i>Capsicum annuum</i>	2451	8,20E+03	85%	0.0	92.31%	55870	NW_015961681.1
CopiaCa_04	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold181	<i>Capsicum annuum</i>	2403	1,05E+04	93%	0.0	92.69%	2703691	NW_015961388.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00475	<i>Solanum tuberosum</i>	2379	5,09E+03	85%	0.0	87.81%	401808	NW_006239407.1
CopiaCa_04	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00307	<i>Solanum tuberosum</i>	2161	5,09E+03	82%	0.0	87.79%	776345	NW_006239230.1
CopiaCa_05	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9225	7,18E+04	100%	0.0	100.00%	215701946	NC_029980.1
CopiaCa_05	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5127	1,28E+08	100%	0.0	85.65%	219521584	NC_029982.1
CopiaCa_05	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5114	9,36E+04	100%	0.0	95.51%	222112641	NC_029983.1
CopiaCa_05	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4518	7,65E+04	99%	0.0	91.28%	163962470	NC_029978.1
CopiaCa_05	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4329	1,23E+08	99%	0.0	88.13%	301019445	NC_029977.1
CopiaCa_05	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3323	7,18E+04	99%	0.0	79.12%	220335243	NC_029987.1
CopiaCa_05	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3142	8,44E+04	99%	0.0	89.08%	229934170	NC_029988.1
CopiaCa_05	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2673	1,17E+08	100%	0.0	95.54%	261510930	NC_029979.1
CopiaCa_05	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2649	6,54E+04	100%	0.0	87.45%	238794889	NC_029985.1
CopiaCa_05	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2091	6,40E+04	99%	0.0	88.76%	205736368	NC_029986.1
CopiaCa_06	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8696	3,30E+04	100%	0.0	99.98%	301019445	NC_029977.1
CopiaCa_06	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8615	1,49E+04	100%	0.0	99.68%	215701946	NC_029980.1
CopiaCa_06	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8610	1,05E+04	100%	0.0	99.66%	163962470	NC_029978.1
CopiaCa_06	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8593	2,18E+04	100%	0.0	99.60%	219521584	NC_029982.1
CopiaCa_06	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8558	1,50E+04	100%	0.0	99.45%	205736368	NC_029986.1
CopiaCa_06	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8556	3,37E+04	100%	0.0	99.47%	238794889	NC_029985.1

CopiaCa_06	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8549	1,51E+04	100%	0.0	99.43%	229934170	NC_029988.1
CopiaCa_06	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8528	1,02E+04	100%	0.0	99.36%	220335243	NC_029987.1
CopiaCa_06	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8521	2,81E+04	100%	0.0	99.36%	222112641	NC_029983.1
CopiaCa_06	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00196	<i>Solanum tuberosum</i>	5513	5,51E+03	98%	0.0	88.13%	1164816	NW_006239123.1
CopiaCa_06	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00063	<i>Solanum tuberosum</i>	5326	5,33E+03	98%	0.0	87.51%	2166112	NW_006238988.1
CopiaCa_06	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00187	<i>Solanum tuberosum</i>	5155	6,31E+03	98%	0.0	89.28%	1214092	NW_006239111.1
CopiaCa_06	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	4863	4,86E+03	99%	0.0	85.59%	56862	NW_017675902.1
CopiaCa_06	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold68506	<i>Nicotiana tabacum</i>	4674	4,67E+03	95%	0.0	85.50%	46075	NW_015929298.1
CopiaCa_06	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold9198	<i>Nicotiana sylvestris</i>	4663	4,66E+03	95%	0.0	85.46%	119532	NW_009583808.1
CopiaCa_06	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 7, NIATTR2	<i>Nicotiana attenuata</i>	4639	4,64E+03	96%	0.0	85.27%	114765206	NC_031995.1
CopiaCa_06	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	4634	4,63E+03	98%	0.0	84.80%	346969	NW_017671151.1
CopiaCa_06	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00467	<i>Solanum tuberosum</i>	4595	6,11E+03	98%	0.0	88.99%	436040	NW_006239388.1
CopiaCa_06	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold37785	<i>Nicotiana sylvestris</i>	4560	4,56E+03	97%	0.0	84.63%	30223	NW_009523592.1
CopiaCa_06	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold97453	<i>Nicotiana tabacum</i>	4549	4,55E+03	99%	0.0	84.29%	28591	NW_015953462.1
CopiaCa_06	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold9493	<i>Nicotiana sylvestris</i>	4503	4,50E+03	97%	0.0	84.48%	212305	NW_009587086.1
CopiaCa_06	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold56083	<i>Nicotiana tabacum</i>	4501	4,50E+03	98%	0.0	84.35%	71758	NW_015918293.1
CopiaCa_06	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	<i>Solanum pennellii</i>	4473	5,50E+03	98%	0.0	84.39%	82529941	NC_028646.1
CopiaCa_06	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 12, NIATTR2	<i>Nicotiana attenuata</i>	4440	4,44E+03	99%	0.0	84.06%	53491057	NC_032000.1
CopiaCa_06	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold21214	<i>Nicotiana tomentosiformis</i>	3954	4,96E+03	94%	0.0	87.57%	216106	NW_008900503.1
CopiaCa_06	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold34809	<i>Nicotiana tabacum</i>	3910	4,92E+03	94%	0.0	87.33%	228018	NW_015895033.1
CopiaCa_06	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	<i>Solanum pennellii</i>	3871	4,81E+03	98%	0.0	84.11%	109333515	NC_028637.1
CopiaCa_06	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	<i>Solanum pennellii</i>	3849	5,37E+03	95%	0.0	83.92%	70546659	NC_028644.1
CopiaCa_06	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	<i>Solanum pennellii</i>	3836	3,84E+03	98%	0.0	81.84%	60730942	NC_028642.1
CopiaCa_06	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	<i>Solanum lycopersicum</i>	3773	3,84E+03	87%	0.0	83.53%	65987440	NC_015445.3
CopiaCa_06	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold6365	<i>Nicotiana tabacum</i>	3766	3,77E+03	87%	0.0	83.34%	84648	NW_015925051.1
CopiaCa_06	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold11333	<i>Nicotiana sylvestris</i>	3766	3,77E+03	87%	0.0	83.34%	286430	NW_009353615.1
CopiaCa_06	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00264	<i>Solanum tuberosum</i>	3589	4,60E+03	88%	0.0	86.70%	901441	NW_006239193.1
CopiaCa_06	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3517	4,09E+03	82%	0.0	86.71%	128824	NW_017673572.1
CopiaCa_06	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	<i>Solanum pennellii</i>	3410	3,78E+04	98%	0.0	82.91%	83305730	NC_028648.1
CopiaCa_06	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	<i>Solanum lycopersicum</i>	3230	4,93E+03	91%	0.0	82.97%	98455869	NC_015438.3
CopiaCa_06	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00291	<i>Solanum tuberosum</i>	3088	5,70E+03	98%	0.0	87.99%	820352	NW_006239218.1
CopiaCa_06	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7671	<i>Nicotiana tabacum</i>	3070	3,58E+03	83%	0.0	82.87%	222948	NW_015936362.1
CopiaCa_06	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold11003	<i>Nicotiana tomentosiformis</i>	3070	3,58E+03	83%	0.0	82.89%	364001	NW_008839637.1
CopiaCa_06	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold756	<i>Capisicum annuum</i>	2418	4,30E+03	80%	0.0	85.95%	884456	NW_015961254.1
CopiaCa_06	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	<i>Solanum lycopersicum</i>	2279	6,08E+03	98%	0.0	83.95%	66723567	NC_015442.3
CopiaCa_06	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	<i>Solanum pennellii</i>	1977	7,14E+03	98%	0.0	84.70%	77991103	NC_028641.1
CopiaCa_07	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	10532	1,81E+08	100%	0.0	100.00%	261510930	NC_029979.1
CopiaCa_07	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	6946	1,07E+08	100%	0.0	93.72%	217274494	NC_029981.1
CopiaCa_07	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	6427	1,21E+08	100%	0.0	91.14%	215701946	NC_029980.1
CopiaCa_07	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	6204	1,12E+08	100%	0.0	90.00%	205736368	NC_029986.1
CopiaCa_07	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5965	1,37E+08	100%	0.0	92.10%	229934170	NC_029988.1
CopiaCa_07	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5716	1,06E+08	100%	0.0	88.34%	219521584	NC_029982.1
CopiaCa_07	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5692	8,15E+04	100%	0.0	91.09%	238794889	NC_029985.1
CopiaCa_07	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5603	1,55E+08	100%	0.0	90.72%	301019445	NC_029977.1
CopiaCa_07	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold519	<i>Capisicum annuum</i>	5598	7,61E+03	86%	0.0	90.62%	1101621	NW_015961578.1
CopiaCa_07	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capisicum annuum</i>	5367	1,10E+04	93%	0.0	89.68%	7059680	NW_015960855.1
CopiaCa_07	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5352	2,57E+04	100%	0.0	89.73%	153299543	NC_029984.1
CopiaCa_07	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5276	8,88E+04	100%	0.0	83.75%	220335243	NC_029987.1
CopiaCa_07	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2615	<i>Capisicum annuum</i>	5094	6,19E+03	86%	0.0	88.52%	131126	NW_015961557.1
CopiaCa_07	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4876	1,30E+08	100%	0.0	90.93%	222112641	NC_029983.1
CopiaCa_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00123	<i>Solanum tuberosum</i>	4542	4,54E+03	82%	0.0	84.34%	1602102	NW_006239048.1
CopiaCa_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00099	<i>Solanum tuberosum</i>	4530	9,24E+03	86%	0.0	86.21%	3955149	NW_006238935.1
CopiaCa_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00245	<i>Solanum tuberosum</i>	4464	4,90E+03	82%	0.0	85.92%	1006898	NW_006239167.1
CopiaCa_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00074	<i>Solanum tuberosum</i>	4423	4,82E+03	82%	0.0	85.78%	2045796	NW_006238998.1
CopiaCa_07	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	<i>Solanum pennellii</i>	4377	1,24E+04	83%	0.0	85.53%	59803892	NC_028638.1
CopiaCa_07	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	<i>Solanum pennellii</i>	4185	1,42E+04	80%	0.0	84.77%	66223686	NC_028647.1

CopiaCa_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	<i>Solanum lycopersicum</i>	4161	1,01E+08	86%	0.0	84.65%	98455869	NC_015438.3
CopiaCa_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00016	<i>Solanum tuberosum</i>	4085	7,76E+03	82%	0.0	82.59%	3295453	NW_006238942.1
CopiaCa_07	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	<i>Solanum pennellii</i>	4034	1,17E+04	87%	0.0	84.12%	84057508	NC_028645.1
CopiaCa_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	<i>Solanum lycopersicum</i>	4034	1,51E+04	86%	0.0	84.18%	68126176	NC_015449.3
CopiaCa_07	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	<i>Solanum pennellii</i>	4026	2,62E+04	82%	0.0	84.15%	60730942	NC_028642.1
CopiaCa_07	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	<i>Solanum pennellii</i>	4023	2,04E+04	85%	0.0	84.13%	77197300	NC_028640.1
CopiaCa_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00157	<i>Solanum tuberosum</i>	4019	4,80E+03	85%	0.0	85.39%	1381298	NW_006239084.1
CopiaCa_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	<i>Solanum lycopersicum</i>	3986	1,27E+04	82%	0.0	83.96%	55977580	NC_015439.3
CopiaCa_07	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 12, NIATTr2	<i>Nicotiana attenuata</i>	3962	1,95E+04	84%	0.0	83.73%	53491057	NC_032000.1
CopiaCa_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00019	<i>Solanum tuberosum</i>	3923	1,20E+04	82%	0.0	86.72%	3257788	NW_006238945.1
CopiaCa_07	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capicum annuum</i>	3910	5,02E+04	89%	0.0	89.51%	163962470	NC_029978.1
CopiaCa_07	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 10, NIATTr2	<i>Nicotiana attenuata</i>	3888	1,55E+04	82%	0.0	83.38%	51082206	NC_031998.1
CopiaCa_07	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	<i>Nicotiana attenuata</i>	3873	4,35E+03	86%	0.0	83.43%	317299	NW_017671305.1
CopiaCa_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	<i>Solanum lycopersicum</i>	3819	1,67E+04	82%	0.0	83.36%	66557038	NC_015441.3
CopiaCa_07	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	<i>Solanum pennellii</i>	3812	1,27E+08	86%	0.0	83.69%	109333515	NC_028637.1
CopiaCa_07	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9282	<i>Nicotiana tabacum</i>	3808	3,81E+03	83%	0.0	81.43%	72030	NW_015949806.1
CopiaCa_07	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsy Nsy_scaffold23224	<i>Nicotiana sylvestris</i>	3808	3,81E+03	83%	0.0	81.43%	209544	NW_009485737.1
CopiaCa_07	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	<i>Solanum pennellii</i>	3751	7,37E+03	82%	0.0	82.93%	79292169	NC_028643.1
CopiaCa_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00235	<i>Solanum tuberosum</i>	3747	4,48E+03	80%	0.0	84.97%	1041465	NW_006239159.1
CopiaCa_07	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold107691	<i>Nicotiana tabacum</i>	3736	3,74E+03	83%	0.0	81.16%	10487	NW_015793168.1
CopiaCa_07	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsy Nsy_scaffold123674	<i>Nicotiana sylvestris</i>	3736	3,74E+03	83%	0.0	81.17%	11578	NW_009365108.1
CopiaCa_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	<i>Solanum lycopersicum</i>	3705	8,75E+03	82%	0.0	82.80%	49794276	NC_015443.3
CopiaCa_07	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 7, NIATTr2	<i>Nicotiana attenuata</i>	3687	2,44E+04	80%	0.0	83.36%	114765206	NC_031995.1
CopiaCa_07	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold30239	<i>Nicotiana tabacum</i>	3675	3,96E+03	82%	0.0	82.55%	58947	NW_015888271.1
CopiaCa_07	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold23275	<i>Nicotiana tomentosiformis</i>	3648	3,65E+03	83%	0.0	80.75%	44328	NW_008902792.1
CopiaCa_07	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13198	<i>Nicotiana tabacum</i>	3611	3,61E+03	83%	0.0	80.62%	39825	NW_015810878.1
CopiaCa_07	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsy Nsy_scaffold31696	<i>Nicotiana sylvestris</i>	3602	3,88E+03	83%	0.0	82.24%	130121	NW_009516827.1
CopiaCa_07	refseq_genomes	8	Capicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capicum annuum</i>	3544	6,12E+03	86%	0.0	88.15%	2397553	NW_015961318.1
CopiaCa_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	<i>Solanum lycopersicum</i>	3467	3,07E+04	86%	0.0	83.79%	65633393	NC_015447.3
CopiaCa_07	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	<i>Solanum pennellii</i>	3446	3,12E+04	86%	0.0	86.15%	82529941	NC_028646.1
CopiaCa_07	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	<i>Solanum pennellii</i>	3277	3,13E+04	82%	0.0	81.04%	77991103	NC_028641.1
CopiaCa_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00255	<i>tuberosum Nicotiana</i>	2265	4,67E+03	81%	0.0	84.93%	929374	NW_006239184.1
CopiaCa_07	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	<i>attenuata Nicotiana</i>	1995	4,42E+03	85%	0.0	85.22%	296448	NW_017671438.1
CopiaCa_07	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsy Nsy_scaffold41756	<i>sylvestris Capsicum</i>	1855	5,33E+03	82%	0.0	84.20%	83875	NW_009528005.1
CopiaCa_08	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	9214	1,01E+04	100%	0.0	100.00%	220335243	NC_029987.1
CopiaCa_08	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	4966	8,68E+03	100%	0.0	93.65%	261510930	NC_029979.1
CopiaCa_08	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	4170	1,24E+04	95%	0.0	94.51%	215701946	NC_029980.1
CopiaCa_08	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	4126	4,20E+03	92%	0.0	83.26%	222112641	NC_029983.1
CopiaCa_08	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	3779	7,63E+03	97%	0.0	91.53%	301019445	NC_029977.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	11315	1,08E+09	100%	0.0	100.00%	163962470	NC_029978.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold985	<i>annuum Capsicum</i>	8237	3,30E+04	100%	0.0	90.97%	1349800	NW_015961816.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	8189	1,10E+09	100%	0.0	90.83%	215701946	NC_029980.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	8135	1,09E+09	100%	0.0	90.71%	205736368	NC_029986.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	8046	1,54E+09	100%	0.0	90.44%	261510930	NC_029979.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	8024	1,39E+09	100%	0.0	90.44%	219521584	NC_029982.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	7938	2,00E+09	100%	0.0	90.11%	301019445	NC_029977.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	7928	5,75E+08	100%	0.0	90.14%	153299543	NC_029984.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	7912	1,28E+09	100%	0.0	90.02%	217274494	NC_029981.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	7906	1,16E+09	100%	0.0	90.01%	220335243	NC_029987.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	7814	1,19E+09	100%	0.0	89.73%	222112641	NC_029983.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1741	<i>annuum Capsicum</i>	7594	1,27E+04	99%	0.0	89.24%	1897442	NW_015960687.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5846	<i>annuum Capsicum</i>	7559	2,05E+04	99%	0.0	89.15%	430778	NW_015961483.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	7526	1,10E+09	100%	0.0	88.90%	238794889	NC_029985.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1894	<i>annuum Capsicum</i>	7454	3,89E+04	99%	0.0	88.80%	2334823	NW_015961490.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>annuum Capsicum</i>	7448	1,41E+09	100%	0.0	88.74%	229934170	NC_029988.1
CopiaCa_09	refseq_genomes	8	Capicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold188	<i>annuum</i>	7372	8,98E+03	100%	0.0	88.48%	980346	NW_015960557.1

CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1107	<i>Capsicum annuum</i>	6556	7,91E+03	99%	0.0	86.19%	318496	NW_015961239.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1289	<i>Capsicum annuum</i>	6270	7,30E+03	96%	0.0	87.76%	151797	NW_015961729.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold181	<i>Capsicum annuum</i>	6192	4,37E+04	100%	0.0	87.29%	2703691	NW_015961388.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capsicum annuum</i>	6082	2,97E+04	98%	0.0	87.01%	2397553	NW_015961318.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold547	<i>Capsicum annuum</i>	6063	6,06E+03	88%	0.0	87.09%	243438	NW_015961245.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	5978	7,42E+03	88%	0.0	86.81%	3311724	NW_015960635.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold949	<i>Capsicum annuum</i>	5960	4,04E+04	99%	0.0	86.77%	3128562	NW_015961487.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold15236	<i>Capsicum annuum</i>	5934	6,22E+03	87%	0.0	86.98%	79606	NW_015961305.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold90	<i>Capsicum annuum</i>	5879	1,18E+04	96%	0.0	86.49%	6074945	NW_015960458.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold561	<i>Capsicum annuum</i>	5869	1,74E+04	96%	0.0	86.63%	1145873	NW_015961012.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3822	<i>Capsicum annuum</i>	5840	7,26E+03	99%	0.0	86.05%	410103	NW_015960760.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1037	<i>Capsicum annuum</i>	5777	6,70E+04	100%	0.0	85.89%	2186250	NW_015960681.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold937	<i>Capsicum annuum</i>	5775	1,57E+04	96%	0.0	86.28%	1465884	NW_015961782.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1246	<i>Capsicum annuum</i>	5722	3,41E+04	98%	0.0	86.00%	3388803	NW_015960834.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2375	<i>Capsicum annuum</i>	5616	5,62E+03	88%	0.0	85.47%	1171512	NW_015961637.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold688	<i>Capsicum annuum</i>	5589	1,82E+04	96%	0.0	86.33%	1322070	NW_015960950.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5785	<i>Capsicum annuum</i>	5581	5,92E+03	94%	0.0	85.65%	79892	NW_015961741.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold46	<i>Capsicum annuum</i>	5561	9,13E+03	98%	0.0	88.07%	941867	NW_015960787.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3083	<i>Capsicum annuum</i>	5542	2,41E+04	95%	0.0	85.54%	288042	NW_015960510.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold592	<i>Capsicum annuum</i>	5515	2,59E+04	88%	0.0	87.17%	1905997	NW_015960988.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2563	<i>Capsicum annuum</i>	5491	1,17E+04	98%	0.0	85.13%	1037879	NW_015961158.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1633	<i>Capsicum annuum</i>	5474	3,08E+04	97%	0.0	85.37%	1620172	NW_015961041.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2298	<i>Capsicum annuum</i>	5426	9,17E+03	85%	0.0	85.95%	1494726	NW_015960454.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1354	<i>Capsicum annuum</i>	5289	3,82E+04	97%	0.0	90.48%	1998780	NW_015960394.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold16	<i>Capsicum annuum</i>	5238	5,24E+03	80%	0.0	85.92%	2023789	NW_015960959.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold258	<i>Capsicum annuum</i>	5197	4,14E+04	98%	0.0	84.69%	3098259	NW_015961180.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold171	<i>Capsicum annuum</i>	5171	1,43E+04	98%	0.0	84.22%	887051	NW_015960473.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4695	<i>Capsicum annuum</i>	5169	1,17E+04	82%	0.0	88.89%	183029	NW_015961566.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold15320	<i>Capsicum annuum</i>	5162	6,01E+03	83%	0.0	88.51%	66261	NW_015961648.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold924	<i>Capsicum annuum</i>	5097	1,04E+04	100%	0.0	85.61%	1342734	NW_015960575.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold891	<i>Capsicum annuum</i>	5079	1,69E+04	91%	0.0	84.34%	1431046	NW_015960469.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3911	<i>Capsicum annuum</i>	4992	1,25E+04	100%	0.0	88.64%	141203	NW_015961599.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3070	<i>Capsicum annuum</i>	4975	6,87E+03	94%	0.0	87.38%	174894	NW_015960750.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold367	<i>Capsicum annuum</i>	4957	1,62E+04	94%	0.0	83.86%	625659	NW_015960847.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	4926	1,77E+04	95%	0.0	86.03%	3493078	NW_015960517.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2901	<i>Capsicum annuum</i>	4909	1,95E+04	99%	0.0	88.82%	1161691	NW_015961380.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold376	<i>Capsicum annuum</i>	4907	4,91E+03	81%	0.0	84.51%	776404	NW_015961666.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2157	<i>Capsicum annuum</i>	4876	7,07E+03	85%	0.0	86.06%	158560	NW_015960692.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3037	<i>Capsicum annuum</i>	4867	1,47E+04	96%	0.0	83.25%	1979607	NW_015960688.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3946	<i>Capsicum annuum</i>	4855	4,86E+03	85%	0.0	83.62%	591642	NW_015960367.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3714	<i>Capsicum annuum</i>	4815	1,57E+04	96%	0.0	84.90%	935490	NW_015961128.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold919	<i>Capsicum annuum</i>	4774	9,30E+03	89%	0.0	86.28%	689688	NW_015961602.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3292	<i>Capsicum annuum</i>	4732	6,36E+03	89%	0.0	88.84%	271392	NW_015960604.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold8858	<i>Capsicum annuum</i>	4728	9,78E+03	98%	0.0	89.71%	16280	NW_015960392.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1174	<i>Capsicum annuum</i>	4630	5,27E+03	88%	0.0	83.70%	656414	NW_015961459.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold659	<i>Capsicum annuum</i>	4604	9,00E+03	87%	0.0	87.45%	613124	NW_015961271.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4495	<i>Capsicum annuum</i>	4534	1,58E+04	100%	0.0	88.41%	406145	NW_015960864.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold979	<i>Capsicum annuum</i>	4481	1,41E+04	99%	0.0	89.67%	1172749	NW_015961157.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold756	<i>Capsicum annuum</i>	4397	1,12E+04	95%	0.0	85.91%	884456	NW_015961254.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3682	<i>Capsicum annuum</i>	4372	9,53E+03	97%	0.0	84.41%	568842	NW_015961135.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1129	<i>Capsicum annuum</i>	4359	1,32E+04	95%	0.0	82.14%	912839	NW_015961757.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold680	<i>Capsicum annuum</i>	4313	1,32E+04	99%	0.0	81.83%	788567	NW_015961649.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold53	<i>Capsicum annuum</i>	4265	1,33E+04	95%	0.0	85.97%	1681775	NW_015961414.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3313	<i>Capsicum annuum</i>	4257	4,26E+03	87%	0.0	81.18%	770237	NW_015961029.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold6648	<i>Capsicum annuum</i>	4181	4,18E+03	88%	0.0	80.86%	204650	NW_015961623.1
CopiaCa_09	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3740	<i>Capsicum annuum</i>	4065	9,46E+03	98%	0.0	88.90%	22371	NW_015960529.1

CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1390	<i>Capisicum annuum</i>	4037	5,23E+03	85%	0.0	80.79%	1730405	NW_015960783.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9	<i>Capisicum annuum</i>	3989	1,70E+04	95%	0.0	80.12%	6567951	NW_015961815.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold899	<i>Capisicum annuum</i>	3978	8,78E+03	97%	0.0	86.12%	265334	NW_015961225.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold127	<i>Capisicum annuum</i>	3969	5,96E+03	99%	0.0	83.55%	289836	NW_015960887.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold555	<i>Capisicum annuum</i>	3964	1,00E+04	99%	0.0	86.65%	1396435	NW_015960461.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold742	<i>Capisicum annuum</i>	3847	7,33E+03	91%	0.0	80.44%	2382255	NW_015961471.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3224	<i>Capisicum annuum</i>	3784	7,27E+03	97%	0.0	84.12%	891864	NW_015961692.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1874	<i>Capisicum annuum</i>	3725	1,57E+04	97%	0.0	88.39%	719064	NW_015961147.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2523	<i>Capisicum annuum</i>	3723	7,90E+03	89%	0.0	84.13%	735434	NW_015960464.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capisicum annuum</i>	3677	6,10E+04	92%	0.0	83.25%	7059680	NW_015960855.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capisicum annuum</i>	3668	1,83E+04	99%	0.0	88.07%	2470984	NW_015961229.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1191	<i>Capisicum annuum</i>	3651	5,93E+03	85%	0.0	86.46%	145056	NW_015961591.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold152	<i>Capisicum annuum</i>	3618	5,58E+03	86%	0.0	84.46%	270641	NW_015961614.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1973	<i>Capisicum annuum</i>	3528	1,34E+04	95%	0.0	84.85%	1714326	NW_015961422.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4311	<i>Capisicum annuum</i>	3465	9,14E+03	84%	0.0	85.89%	294642	NW_015961514.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold6075	<i>Capisicum annuum</i>	3417	6,97E+03	84%	0.0	90.14%	316563	NW_015960475.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4195	<i>Capisicum annuum</i>	3395	6,82E+03	80%	0.0	85.21%	98679	NW_015961226.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold516	<i>Capisicum annuum</i>	3382	8,86E+03	97%	0.0	82.68%	4299619	NW_015961344.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1620	<i>Capisicum annuum</i>	3347	1,16E+04	86%	0.0	87.15%	500299	NW_015961331.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold306	<i>Capisicum annuum</i>	3323	7,46E+03	84%	0.0	81.88%	749978	NW_015961154.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold11031	<i>Capisicum annuum</i>	3321	5,55E+03	92%	0.0	83.83%	96667	NW_015961583.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold8872	<i>Capisicum annuum</i>	3299	4,72E+03	88%	0.0	84.89%	40610	NW_015961697.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold148	<i>Capisicum annuum</i>	3221	1,29E+04	89%	0.0	78.93%	1323090	NW_015961290.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1110	<i>Capisicum annuum</i>	3195	7,05E+03	88%	0.0	82.29%	740557	NW_015961474.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold170	<i>Capisicum annuum</i>	3121	8,58E+03	89%	0.0	83.44%	2217302	NW_015960390.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1913	<i>Capisicum annuum</i>	3079	7,68E+03	94%	0.0	84.40%	65133	NW_015961762.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3931	<i>Capisicum annuum</i>	3035	4,82E+03	99%	0.0	83.79%	84366	NW_015960476.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold920	<i>Capisicum annuum</i>	2870	9,09E+03	99%	0.0	84.95%	197492	NW_015961621.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold438	<i>Capisicum annuum</i>	2824	7,07E+03	98%	0.0	79.60%	1337825	NW_015961507.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1399	<i>Capisicum annuum</i>	2800	8,94E+03	87%	0.0	89.18%	1742971	NW_015961630.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4129	<i>Capisicum annuum</i>	2682	9,36E+03	85%	0.0	82.89%	328737	NW_015961069.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold24	<i>Capisicum annuum</i>	2560	8,42E+03	94%	0.0	86.28%	1258456	NW_015961700.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5097	<i>Capisicum annuum</i>	2531	6,45E+03	89%	0.0	82.71%	302211	NW_015961801.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold603	<i>Capisicum annuum</i>	2344	5,04E+03	81%	0.0	87.31%	1800753	NW_015960521.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1908	<i>Capisicum annuum</i>	2206	6,62E+03	83%	0.0	79.33%	666372	NW_015961321.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold7937	<i>Capisicum annuum</i>	2189	5,75E+03	80%	0.0	88.23%	90413	NW_015961057.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1652	<i>Capisicum annuum</i>	2135	4,31E+03	85%	0.0	83.20%	205168	NW_015961309.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4197	<i>Capisicum annuum</i>	2008	3,91E+03	82%	0.0	82.29%	182937	NW_015961354.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1373	<i>Capisicum annuum</i>	1960	5,91E+03	80%	0.0	87.46%	351742	NW_015960632.1
CopiaCa_09	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5380	<i>Capisicum annuum</i>	1786	5,56E+03	83%	0.0	88.31%	27469	NW_015961313.1
CopiaCa_10	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8458	9,88E+04	100%	0.0	100.00%	217274494	NC_029981.1
CopiaCa_10	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	6024	8,32E+04	100%	0.0	90.58%	215701946	NC_029980.1
CopiaCa_10	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	6010	8,65E+04	100%	0.0	90.92%	301019445	NC_029977.1
CopiaCa_10	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5851	8,85E+04	100%	0.0	89.86%	220335243	NC_029987.1
CopiaCa_10	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5688	4,95E+04	100%	0.0	89.52%	219521584	NC_029982.1
CopiaCa_10	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5620	6,00E+04	100%	0.0	89.09%	205736368	NC_029986.1
CopiaCa_10	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5563	5,31E+04	100%	0.0	90.95%	261510930	NC_029979.1
CopiaCa_10	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5448	3,48E+04	100%	0.0	89.68%	163962470	NC_029978.1
CopiaCa_10	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5432	7,17E+04	100%	0.0	88.60%	229934170	NC_029988.1
CopiaCa_10	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1399	<i>Capisicum annuum</i>	5400	1,34E+04	100%	0.0	89.36%	1742971	NW_015961630.1
CopiaCa_10	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold920	<i>Capisicum annuum</i>	5380	5,83E+03	100%	0.0	89.43%	197492	NW_015961621.1
CopiaCa_10	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5321	8,74E+04	100%	0.0	88.05%	222112641	NC_029983.1
CopiaCa_10	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold148	<i>Capisicum annuum</i>	5260	5,56E+03	94%	0.0	90.20%	1323090	NW_015961290.1
CopiaCa_10	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5101	1,97E+04	100%	0.0	88.14%	238794889	NC_029985.1
CopiaCa_10	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2138	<i>Capisicum annuum</i>	5097	5,45E+03	99%	0.0	87.96%	681955	NW_015960370.1
CopiaCa_10	refseq_genomes	8	Capisicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold688	<i>Capisicum annuum</i>	4920	1,06E+04	100%	0.0	87.40%	1322070	NW_015960950.1

CopiaCa_10	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold171	<i>Capsicum annuum</i>	4765	4,85E+03	97%	0.0	86.05%	887051	NW_015960473.1
CopiaCa_10	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	<i>Solanum pennellii</i>	4250	2,75E+08	98%	0.0	84.73%	109333515	NC_028637.1
CopiaCa_10	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	<i>Solanum pennellii</i>	4241	1,39E+08	98%	0.0	84.54%	77991103	NC_028641.1
CopiaCa_10	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	<i>Solanum pennellii</i>	4229	1,53E+08	98%	0.0	84.66%	83305730	NC_028648.1
CopiaCa_10	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	<i>Solanum pennellii</i>	4220	1,49E+08	98%	0.0	84.62%	79292169	NC_028643.1
CopiaCa_10	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	<i>Solanum pennellii</i>	4217	1,13E+08	98%	0.0	85.14%	75414019	NC_028639.1
CopiaCa_10	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	<i>Solanum pennellii</i>	4215	1,30E+08	98%	0.0	84.60%	60730942	NC_028642.1
CopiaCa_10	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	<i>Solanum pennellii</i>	4211	2,37E+08	98%	0.0	84.22%	82529941	NC_028646.1
CopiaCa_10	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	<i>Solanum pennellii</i>	4202	1,68E+08	98%	0.0	84.54%	84057508	NC_028645.1
CopiaCa_10	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	<i>Solanum pennellii</i>	4198	8,57E+04	97%	0.0	84.53%	66223686	NC_028647.1
CopiaCa_10	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	<i>Solanum pennellii</i>	4196	1,04E+08	97%	0.0	84.74%	70546659	NC_028644.1
CopiaCa_10	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	<i>Solanum pennellii</i>	4194	1,17E+08	97%	0.0	84.72%	77197300	NC_028640.1
CopiaCa_10	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	<i>Solanum pennellii</i>	4191	7,94E+04	96%	0.0	84.50%	59803892	NC_028638.1
CopiaCa_10	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold258	<i>Capsicum annuum</i>	4060	4,48E+03	100%	0.0	84.98%	3098259	NW_015961180.1
CopiaCa_10	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	<i>Solanum lycopersicum</i>	4037	1,74E+08	98%	0.0	83.88%	65633393	NC_015447.3
CopiaCa_10	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	<i>Solanum lycopersicum</i>	3975	1,22E+08	97%	0.0	83.33%	49794276	NC_015443.3
CopiaCa_10	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	<i>Solanum lycopersicum</i>	3971	1,41E+08	98%	0.0	83.82%	72290146	NC_015440.3
CopiaCa_10	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	<i>Solanum lycopersicum</i>	3949	1,18E+08	98%	0.0	83.46%	66723567	NC_015442.3
CopiaCa_10	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3940	1,51E+04	100%	0.0	90.29%	153299543	NC_029984.1
CopiaCa_10	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	<i>Solanum lycopersicum</i>	3890	1,33E+08	98%	0.0	82.63%	66557038	NC_015441.3
CopiaCa_10	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	<i>Solanum lycopersicum</i>	3886	1,88E+08	98%	0.0	83.23%	72906345	NC_015446.3
CopiaCa_10	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	<i>Solanum lycopersicum</i>	3884	2,34E+08	98%	0.0	83.48%	98455869	NC_015438.3
CopiaCa_10	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	<i>Solanum lycopersicum</i>	3877	1,47E+08	98%	0.0	82.95%	68126176	NC_015449.3
CopiaCa_10	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	<i>Solanum lycopersicum</i>	3868	1,20E+08	98%	0.0	82.55%	65987440	NC_015445.3
CopiaCa_10	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	<i>Solanum lycopersicum</i>	3832	1,75E+08	98%	0.0	83.24%	68175699	NC_015444.3
CopiaCa_10	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	<i>Solanum lycopersicum</i>	3827	6,49E+04	97%	0.0	83.00%	56597135	NC_015448.3
CopiaCa_10	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	<i>Solanum lycopersicum</i>	3757	8,93E+04	97%	0.0	82.95%	55977580	NC_015439.3
CopiaCa_10	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1647	<i>Capsicum annuum</i>	3561	5,47E+03	99%	0.0	86.70%	1362880	NW_015960892.1
CopiaCa_10	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC00000001	<i>Solanum lycopersicum</i>	3557	6,45E+03	92%	0.0	82.07%	2014460	NW_020442482.1
CopiaCa_10	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC00000003	<i>Solanum lycopersicum</i>	3542	3,54E+03	90%	0.0	82.43%	1066869	NW_020442484.1
CopiaCa_10	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC00000009	<i>Solanum lycopersicum</i>	3334	3,33E+03	96%	0.0	80.62%	797759	NW_020442490.1
CopiaCa_10	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold17904	<i>Capsicum annuum</i>	3330	4,33E+03	85%	0.0	86.53%	12803	NW_015960964.1
CopiaCa_10	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC00000002	<i>Solanum lycopersicum</i>	3238	5,82E+03	92%	0.0	80.80%	177393	NW_020442483.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10763	7,13E+04	100%	0.0	100.00%	238794889	NC_029985.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8818	1,67E+08	100%	0.0	94.00%	301019445	NC_029977.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8674	1,18E+08	100%	0.0	93.53%	215701946	NC_029980.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8447	1,12E+08	100%	0.0	92.97%	219521584	NC_029982.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold7059	<i>Capsicum annuum</i>	7557	8,52E+03	99%	0.0	91.24%	199068	NW_015961188.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7467	1,11E+08	100%	0.0	90.02%	217274494	NC_029981.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7448	1,43E+08	100%	0.0	90.03%	229934170	NC_029988.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7384	1,61E+08	100%	0.0	90.23%	163962470	NC_029978.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7321	1,76E+08	100%	0.0	89.54%	261510930	NC_029979.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6937	1,34E+08	100%	0.0	90.27%	220335243	NC_029987.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6824	1,41E+08	100%	0.0	92.38%	205736368	NC_029986.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2400	<i>Capsicum annuum</i>	6789	8,16E+03	97%	0.0	90.21%	1353284	NW_015960968.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6538	4,26E+04	100%	0.0	89.26%	153299543	NC_029984.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1894	<i>Capsicum annuum</i>	6453	8,32E+03	99%	0.0	88.69%	2334823	NW_015961490.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6268	9,30E+04	100%	0.0	89.09%	222112641	NC_029983.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5971	<i>Capsicum annuum</i>	6056	7,94E+03	97%	0.0	90.85%	156035	NW_015961177.1
CopiaCa_11	refseq-genome	8	Solanum pennellii chromosome 1, SPENN200	<i>Solanum pennellii</i>	5651	7,57E+04	99%	0.0	86.79%	109333515	NC_028637.1
CopiaCa_11	refseq-genome	8	Solanum pennellii chromosome 9, SPENN200	<i>Solanum pennellii</i>	5646	3,92E+04	94%	0.0	86.76%	84057508	NC_028645.1
CopiaCa_11	refseq-genome	8	Solanum pennellii chromosome 8, SPENN200	<i>Solanum pennellii</i>	5622	1,23E+04	94%	0.0	87.09%	70546659	NC_028644.1
CopiaCa_11	refseq-genome	8	Solanum pennellii chromosome 5, SPENN200	<i>Solanum pennellii</i>	5620	4,32E+04	99%	0.0	86.68%	77991103	NC_028641.1
CopiaCa_11	refseq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	<i>Solanum lycopersicum</i>	5483	2,69E+04	94%	0.0	86.22%	55977580	NC_015439.3
CopiaCa_11	refseq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00147	<i>Solanum tuberosum</i>	5445	6,41E+03	90%	0.0	86.43%	1440303	NW_006239070.1
CopiaCa_11	refseq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	<i>Solanum lycopersicum</i>	5408	5,01E+04	94%	0.0	85.94%	65633393	NC_015447.3

CopiaCa_11	refseq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	<i>Solanum lycopersicum</i>	5397	3,79E+04	96%	0.0	85.90%	66723567	NC_015442.3
CopiaCa_11	refseq-genome	8	Solanum pennellii chromosome 3, SPENN200	<i>Solanum pennellii</i>	5371	2,97E+04	99%	0.0	86.02%	75414019	NC_028639.1
CopiaCa_11	refseq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	<i>Solanum lycopersicum</i>	5365	3,35E+04	94%	0.0	85.81%	68126176	NC_015449.3
CopiaCa_11	refseq-genome	8	Solanum pennellii chromosome 11, SPENN200	<i>Solanum pennellii</i>	5315	2,23E+04	94%	0.0	85.82%	66223686	NC_028647.1
CopiaCa_11	refseq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	<i>Solanum lycopersicum</i>	5293	1,80E+04	94%	0.0	85.56%	56597135	NC_015448.3
CopiaCa_11	refseq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00493	<i>Solanum tuberosum</i>	5249	6,24E+03	92%	0.0	85.55%	370066	NW_006239430.1
CopiaCa_11	refseq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	<i>Solanum lycopersicum</i>	5212	3,12E+04	94%	0.0	85.31%	72290146	NC_015440.3
CopiaCa_11	refseq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00161	<i>Solanum tuberosum</i>	5186	5,85E+03	93%	0.0	85.29%	1344915	NW_006239086.1
CopiaCa_11	refseq-genome	8	Solanum pennellii chromosome 10, SPENN200	<i>Solanum pennellii</i>	5147	2,18E+04	98%	0.0	85.05%	82529941	NC_028646.1
CopiaCa_11	refseq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00281	<i>Solanum tuberosum</i>	5138	6,12E+03	90%	0.0	85.40%	861293	NW_006239206.1
CopiaCa_11	refseq-genome	8	Solanum pennellii chromosome 7, SPENN200	<i>Solanum pennellii</i>	5134	3,26E+04	94%	0.0	85.15%	79292169	NC_028643.1
CopiaCa_11	refseq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	<i>Solanum lycopersicum</i>	5081	1,51E+04	94%	0.0	85.12%	65987440	NC_015445.3
CopiaCa_11	refseq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00046	<i>Solanum tuberosum</i>	4887	6,42E+03	96%	0.0	86.30%	2411080	NW_006238971.1
CopiaCa_11	refseq-genome	8	Solanum pennellii chromosome 4, SPENN200	<i>Solanum pennellii</i>	4883	3,07E+04	96%	0.0	84.14%	77197300	NC_028640.1
CopiaCa_11	refseq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold561	<i>Capsicum annuum</i>	4658	8,67E+03	100%	0.0	89.98%	1145873	NW_015961012.1
CopiaCa_11	refseq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	<i>Solanum lycopersicum</i>	4617	2,81E+04	94%	0.0	83.26%	68175699	NC_015444.3
CopiaCa_11	refseq-genome	8	Solanum pennellii chromosome 12, SPENN200	<i>Solanum pennellii</i>	4518	2,22E+04	94%	0.0	86.73%	83305730	NC_028648.1
CopiaCa_11	refseq-genome	8	Solanum pennellii chromosome 2, SPENN200	<i>Solanum pennellii</i>	4475	1,27E+04	94%	0.0	84.86%	59803892	NC_028638.1
CopiaCa_11	refseq-genome	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5309	<i>Capsicum annum</i>	4420	8,06E+03	92%	0.0	92.00%	181449	NW_015960649.1
CopiaCa_11	refseq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	<i>Solanum lycopersicum</i>	4420	1,94E+04	94%	0.0	83.03%	72906345	NC_015446.3
CopiaCa_11	refseq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00430	<i>Solanum tuberosum</i>	4283	5,53E+03	95%	0.0	84.68%	474478	NW_006239366.1
CopiaCa_11	refseq-genome	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold985	<i>Capsicum annum</i>	4259	9,02E+03	83%	0.0	93.48%	1349800	NW_015961816.1
CopiaCa_11	refseq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold36352	<i>Nicotiana tabacum</i>	3951	4,22E+03	80%	0.0	82.89%	39048	NW_015900127.1
CopiaCa_11	refseq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold18553	<i>Nicotiana sylvestris</i>	3951	4,22E+03	80%	0.0	82.89%	42023	NW_009433836.1
CopiaCa_11	refseq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	<i>Solanum lycopersicum</i>	3919	2,46E+04	99%	0.0	84.68%	98455869	NC_015438.3
CopiaCa_11	refseq-genome	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold979	<i>Capsicum annum</i>	3869	7,94E+03	94%	0.0	91.73%	1172749	NW_015961157.1
CopiaCa_11	refseq-genome	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3904	<i>Capsicum annum</i>	3768	8,69E+03	99%	0.0	90.35%	169740	NW_015960928.1
CopiaCa_11	refseq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 contig00003	<i>Solanum tuberosum</i>	3581	6,17E+03	99%	0.0	85.59%	28034	NW_006239975.1
CopiaCa_11	refseq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00073	<i>Solanum tuberosum</i>	3542	6,18E+03	98%	0.0	83.99%	2002897	NW_006239002.1
CopiaCa_11	refseq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00667	<i>Solanum tuberosum</i>	3308	6,05E+03	95%	0.0	85.99%	208703	NW_006239588.1
CopiaCa_11	refseq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTTr2	<i>Nicotiana attenuata</i>	3208	4,91E+03	84%	0.0	82.84%	837735	NW_017670380.1
CopiaCa_11	refseq-genome	8	Solanum pennellii chromosome 6, SPENN200	<i>Solanum pennellii</i>	3169	6,19E+03	95%	0.0	82.01%	60730942	NC_028642.1
CopiaCa_11	refseq-genome	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold170	<i>Capsicum annum</i>	2994	1,01E+04	88%	0.0	88.22%	2217302	NW_015960390.1
CopiaCa_11	refseq-genome	8	Nicotiana attenuata strain UT chromosome 12, NIATTTr2	<i>Nicotiana attenuata</i>	2957	4,44E+03	84%	0.0	83.17%	53491057	NC_032000.1
CopiaCa_11	refseq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	<i>Solanum lycopersicum</i>	2926	2,82E+04	96%	0.0	83.23%	66557038	NC_015441.3
CopiaCa_11	refseq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold62305	<i>Nicotiana tabacum</i>	2922	4,24E+03	86%	0.0	81.08%	97576	NW_015923851.1
CopiaCa_11	refseq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00060	<i>Solanum tuberosum</i>	2706	5,93E+03	97%	0.0	84.39%	2184951	NW_006238985.1
CopiaCa_11	refseq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00279	<i>Solanum tuberosum</i>	2691	6,34E+03	98%	0.0	84.15%	864930	NW_006239204.1
CopiaCa_11	refseq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	<i>Solanum lycopersicum</i>	2516	5,01E+03	95%	0.0	80.86%	49794276	NC_015443.3
CopiaCa_11	refseq-genome	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2138	<i>Capsicum annum</i>	2326	9,22E+03	99%	0.0	89.14%	681955	NW_015960370.1
CopiaCa_11	refseq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold5195	<i>Nicotiana sylvestris</i>	2222	5,04E+03	85%	0.0	82.07%	112661	NW_009539331.1
CopiaCa_11	refseq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold24535	<i>Nicotiana tomentosiformis</i>	2154	5,38E+03	83%	0.0	84.44%	79504	NW_008904192.1
CopiaCa_11	refseq-genome	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold592	<i>Capsicum annum</i>	2037	7,27E+03	80%	0.0	92.78%	1905997	NW_015960988.1
CopiaCa_11	refseq-genome	8	Nicotiana attenuata strain UT chromosome 5, NIATTTr2	<i>Nicotiana attenuata</i>	1949	5,51E+03	87%	0.0	83.60%	57012777	NC_031993.1
CopiaCa_11	refseq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold26753	<i>Nicotiana sylvestris</i>	1718	5,10E+03	87%	0.0	83.38%	138211	NW_009511335.1
CopiaCa_11	refseq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTTr2	<i>Nicotiana attenuata</i>	1639	4,75E+03	85%	0.0	83.61%	923158	NW_017670348.1
CopiaCa_12	refseq-genome	8	Capsicum annum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annum</i>	10532	1,40E+08	100%	0.0	100.00%	215701946	NC_029980.1
CopiaCa_12	refseq-genome	8	Capsicum annum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annum</i>	7806	6,07E+04	100%	0.0	91.73%	153299543	NC_029984.1
CopiaCa_12	refseq-genome	8	Capsicum annum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annum</i>	6861	1,77E+08	100%	0.0	88.96%	261510930	NC_029979.1
CopiaCa_12	refseq-genome	8	Capsicum annum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annum</i>	6505	8,92E+04	100%	0.0	88.84%	219521584	NC_029982.1
CopiaCa_12	refseq-genome	8	Capsicum annum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annum</i>	6444	1,21E+08	99%	0.0	89.18%	229934170	NC_029988.1
CopiaCa_12	refseq-genome	8	Capsicum annum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annum</i>	5638	9,06E+04	100%	0.0	91.72%	222112641	NC_029983.1
CopiaCa_12	refseq-genome	8	Capsicum annum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annum</i>	5520	9,97E+04	100%	0.0	89.23%	217274494	NC_029981.1
CopiaCa_12	refseq-genome	8	Capsicum annum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annum</i>	5437	1,19E+08	100%	0.0	89.68%	205736368	NC_029986.1
CopiaCa_12	refseq-genome	8	Capsicum annum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annum</i>	4584	6,52E+04	100%	0.0	88.20%	163962470	NC_029978.1
CopiaCa_12	refseq-genome	8	Capsicum annum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annum</i>	4486	1,42E+08	99%	0.0	88.51%	301019445	NC_029977.1

CopiaCa_12	refseq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1307	<i>Capisicum annuum</i>	4353	5,65E+03	82%	0.0	87.91%	1549638	NW_015960466.1
CopiaCa_12	refseq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4198	7,91E+04	99%	0.0	87.17%	238794889	NC_029985.1
CopiaCa_12	refseq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold11001	<i>Capisicum annuum</i>	3633	8,10E+03	82%	0.0	85.04%	37862	NW_015961788.1
CopiaCa_12	refseq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	3445	7,68E+04	99%	0.0	86.39%	220335243	NC_029987.1
CopiaCa_12	refseq-genome	8	Nicotiana attenuata strain UT chromosome 12, NIATTR2	<i>Nicotiana attenuata</i>	3147	2,12E+04	84%	0.0	81.57%	53491057	NC_032000.1
CopiaCa_12	refseq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3068	3,85E+03	82%	0.0	81.21%	189028	NW_017672480.1
CopiaCa_12	refseq-genome	8	Nicotiana attenuata strain UT chromosome 6, NIATTR2	<i>Nicotiana attenuata</i>	3066	1,96E+04	85%	0.0	81.20%	97571839	NC_031994.1
CopiaCa_12	refseq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3066	3,79E+03	82%	0.0	81.20%	418277	NW_017670890.1
CopiaCa_12	refseq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3025	3,82E+03	81%	0.0	81.00%	435747	NW_017670841.1
CopiaCa_12	refseq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3013	1,34E+04	84%	0.0	81.05%	392266	NW_017670978.1
CopiaCa_12	refseq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold438	<i>Capisicum annuum</i>	2963	1,01E+04	87%	0.0	89.37%	1337825	NW_015961507.1
CopiaCa_12	refseq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold54772	<i>Nicotiana sylvestris</i>	2959	3,70E+03	82%	0.0	80.77%	63529	NW_009542467.1
CopiaCa_12	refseq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8858	<i>Nicotiana tabacum</i>	2952	3,79E+03	84%	0.0	80.73%	107390	NW_015946302.1
CopiaCa_12	refseq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold65329	<i>Nicotiana sylvestris</i>	2950	3,97E+03	81%	0.0	80.69%	37459	NW_009554197.1
CopiaCa_12	refseq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold145645	<i>Nicotiana tabacum</i>	2944	3,97E+03	81%	0.0	80.66%	34136	NW_015820110.1
CopiaCa_12	refseq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold19190	<i>Nicotiana tabacum</i>	2924	3,75E+03	80%	0.0	80.63%	71204	NW_015847341.1
CopiaCa_12	refseq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold44111	<i>Nicotiana sylvestris</i>	2918	3,74E+03	80%	0.0	80.62%	44903	NW_009530623.1
CopiaCa_12	refseq-genome	8	Nicotiana attenuata strain UT chromosome 5, NIATTR2	<i>Nicotiana attenuata</i>	2902	2,10E+04	85%	0.0	80.56%	57012777	NC_031993.1
CopiaCa_12	refseq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold8899	<i>Nicotiana sylvestris</i>	2896	3,61E+03	81%	0.0	80.49%	268616	NW_009580485.1
CopiaCa_12	refseq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold21160	<i>Nicotiana sylvestris</i>	2892	3,61E+03	82%	0.0	80.41%	45124	NW_009462804.1
CopiaCa_12	refseq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold86202	<i>Nicotiana tabacum</i>	2887	3,61E+03	82%	0.0	80.39%	42827	NW_015944353.1
CopiaCa_12	refseq-genome	8	Nicotiana attenuata strain UT chromosome 2, NIATTR2	<i>Nicotiana attenuata</i>	2878	4,76E+04	84%	0.0	80.44%	96359640	NC_031990.1
CopiaCa_12	refseq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	2870	6,83E+03	81%	0.0	80.51%	424392	NW_017670869.1
CopiaCa_12	refseq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	2863	3,57E+03	81%	0.0	80.39%	226049	NW_017672020.1
CopiaCa_12	refseq-genome	8	Solanum pennellii chromosome 7, SPENNIV200	<i>Solanum pennellii</i>	2830	1,52E+04	81%	0.0	81.30%	79292169	NC_028643.1
CopiaCa_12	refseq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	2789	3,61E+03	83%	0.0	79.97%	545044	NW_017670607.1
CopiaCa_12	refseq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold42894	<i>Nicotiana tabacum</i>	2713	3,73E+03	80%	0.0	80.65%	69027	NW_015906203.1
CopiaCa_12	refseq-genome	8	Nicotiana attenuata strain UT chromosome 1, NIATTR2	<i>Nicotiana attenuata</i>	2691	2,83E+04	81%	0.0	80.71%	110538649	NC_031989.1
CopiaCa_12	refseq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold38642	<i>Nicotiana sylvestris</i>	2675	3,70E+03	80%	0.0	80.46%	45223	NW_009524545.1
CopiaCa_12	refseq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1894	<i>Capisicum annuum</i>	2213	5,20E+03	82%	0.0	87.58%	2334823	NW_015961490.1
CopiaCa_12	refseq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold61724	<i>Nicotiana tabacum</i>	2076	3,47E+03	80%	0.0	83.02%	21877	NW_015923322.1
CopiaCa_12	refseq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31406	<i>Nicotiana tomentosiformis</i>	2076	3,49E+03	80%	0.0	83.02%	35658	NW_008911827.1
CopiaCa_13	refseq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	9371	1,07E+04	100%	0.0	99.55%	229934170	NC_029988.1
CopiaCa_13	refseq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1422	<i>Capisicum annuum</i>	4673	3,40E+04	100%	0.0	94.48%	802096	NW_015960747.1
CopiaCa_13	refseq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4567	8,69E+03	99%	0.0	93.75%	163962470	NC_029987.1
CopiaCa_13	refseq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4523	8,34E+03	99%	0.0	93.58%	220335243	NC_029987.1
CopiaCa_13	refseq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4298	1,61E+04	99%	0.0	92.28%	217274494	NC_029981.1
CopiaCa_13	refseq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4292	8,15E+03	99%	0.0	92.23%	301019445	NC_029977.1
CopiaCa_13	refseq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1795	<i>Capisicum annuum</i>	3223	8,67E+03	99%	0.0	94.11%	162617	NW_015961207.1
CopiaCa_13	refseq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	3142	1,64E+04	99%	0.0	93.45%	219521584	NC_029982.1
CopiaCa_13	refseq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold7565	<i>Capisicum annuum</i>	2394	7,89E+03	87%	0.0	94.35%	48932	NW_015960657.1
CopiaCa_14	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	9049	3,03E+04	100%	0.0	100.00%	238794889	NC_029985.1
CopiaCa_14	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4809	5,43E+04	100%	0.0	87.31%	163962470	NC_029978.1
CopiaCa_14	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4647	4,76E+04	100%	0.0	86.79%	261510930	NC_029979.1
CopiaCa_14	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4599	3,89E+04	99%	0.0	86.53%	215701946	NC_029980.1
CopiaCa_14	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4569	6,30E+04	100%	0.0	86.41%	217274494	NC_029981.1
CopiaCa_14	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4479	5,10E+04	99%	0.0	87.39%	205736368	NC_029986.1
CopiaCa_14	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4460	6,00E+04	100%	0.0	87.29%	229934170	NC_029988.1
CopiaCa_14	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4455	2,83E+04	99%	0.0	85.96%	219521584	NC_029982.1
CopiaCa_14	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4357	2,79E+04	100%	0.0	86.03%	153299543	NC_029984.1
CopiaCa_14	refeq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2938	<i>Capisicum annuum</i>	4300	5,49E+03	98%	0.0	86.82%	524377	NW_015960410.1
CopiaCa_14	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4231	4,14E+04	99%	0.0	85.14%	301019445	NC_029977.1
CopiaCa_14	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4226	6,35E+04	100%	0.0	85.56%	222112641	NC_029983.1
CopiaCa_14	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00221	<i>Solanum tuberosum</i>	3336	3,34E+03	80%	0.0	82.17%	1069689	NW_006239151.1
CopiaCa_14	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf0024	<i>Solanum tuberosum</i>	3288	5,07E+03	80%	0.0	81.94%	2984907	NW_006238950.1
CopiaCa_14	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	3099	1,13E+04	88%	0.0	81.07%	220335243	NC_029987.1

CopiaCa_14	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1037	<i>Capsicum annuum</i>	2957	1,03E+04	98%	0.0	83.86%	2186250	NW_015960681.1
CopiaCa_14	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold775	<i>Capsicum annuum</i>	2922	5,20E+03	95%	0.0	88.76%	696435	NW_015961719.1
CopiaCa_14	refeq-genome	8	Nicotiana attenuata strain UT chromosome 6, NIATT2	<i>Nicotiana attenuata</i>	2660	3,44E+03	87%	0.0	81.13%	97571839	NC_031994.1
CopiaCa_14	refeq-genome	8	Solanum pennellii chromosome 7, SPENNV200	<i>Solanum pennellii</i>	2641	2,64E+03	80%	0.0	79.21%	79292169	NC_028643.1
CopiaCa_14	refeq-genome	8	Solanum pennellii chromosome 1, SPENNV200	<i>Solanum pennellii</i>	2451	3,00E+03	80%	0.0	78.14%	109333515	NC_028637.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11819	1,02E+08	100%	0.0	100.00%	220335243	NC_029987.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6445	9,30E+04	94%	0.0	93.02%	205736368	NC_029986.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6434	2,44E+08	100%	0.0	92.99%	301019445	NC_029977.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6433	2,00E+08	95%	0.0	92.50%	261510930	NC_029979.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6346	1,20E+08	95%	0.0	92.02%	229934170	NC_029988.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6342	9,64E+04	94%	0.0	92.03%	219521584	NC_029982.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6253	1,11E+08	94%	0.0	91.65%	217274494	NC_029981.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6218	7,52E+04	88%	0.0	91.96%	222112641	NC_029983.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold603	<i>Capsicum annuum</i>	6207	7,94E+03	92%	0.0	91.92%	1800753	NW_015960521.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6185	6,66E+04	93%	0.0	91.90%	215701946	NC_029980.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6141	1,26E+08	97%	0.0	91.19%	163962470	NC_029978.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6102	1,10E+08	93%	0.0	91.50%	238794889	NC_029985.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6030	6,73E+04	93%	0.0	91.05%	153299543	NC_029984.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2274	<i>Capsicum annuum</i>	6028	7,48E+03	80%	0.0	91.51%	1395869	NW_015961152.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	5646	6,68E+03	91%	0.0	90.14%	3311724	NW_015960635.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4706	<i>Capsicum annuum</i>	5384	1,03E+04	92%	0.0	93.22%	267933	NW_015961096.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold258	<i>Capsicum annuum</i>	5083	7,13E+03	81%	0.0	92.33%	3098259	NW_015961180.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold680	<i>Capsicum annuum</i>	4929	9,86E+03	92%	0.0	92.53%	788567	NW_015961649.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold181	<i>Capsicum annuum</i>	4763	6,98E+03	81%	0.0	91.97%	2703691	NW_015961388.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold592	<i>Capsicum annuum</i>	4549	1,39E+04	91%	0.0	90.77%	1905997	NW_015960988.1
CopiaCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1653	<i>Capsicum annuum</i>	3448	1,10E+04	86%	0.0	91.21%	913988	NW_015961387.1
CopiaCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8654	8,27E+04	100%	0.0	100.00%	215701946	NC_029980.1
CopiaCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7247	4,52E+04	100%	0.0	94.59%	220335243	NC_029987.1
CopiaCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7020	9,42E+04	100%	0.0	93.76%	301019445	NC_029977.1
CopiaCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5718	6,31E+04	100%	0.0	94.95%	219521584	NC_029982.1
CopiaCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5670	6,33E+04	100%	0.0	94.54%	217274494	NC_029981.1
CopiaCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5269	6,68E+04	100%	0.0	91.80%	205736368	NC_029986.1
CopiaCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4543	4,32E+04	96%	0.0	90.62%	238794889	NC_029985.1
CopiaCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4327	4,47E+04	100%	0.0	93.67%	163962470	NC_029978.1
CopiaCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3869	4,80E+04	100%	0.0	85.11%	229934170	NC_029988.1
CopiaCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3515	6,40E+04	100%	0.0	90.56%	261510930	NC_029979.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9450	5,63E+04	100%	0.0	100.00%	220335243	NC_029987.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6109	5,73E+04	100%	0.0	88.35%	261510930	NC_029979.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5199	4,45E+04	95%	0.0	88.29%	301019445	NC_029977.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4964	3,45E+04	93%	0.0	87.31%	205736368	NC_029986.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold181	<i>Capsicum annuum</i>	4959	5,75E+03	90%	0.0	87.47%	2703691	NW_015961388.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4833	4,55E+04	98%	0.0	86.73%	219521584	NC_029982.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4828	2,25E+04	95%	0.0	86.82%	163962470	NC_029978.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4820	5,61E+04	100%	0.0	86.76%	215701946	NC_029980.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4687	4,92E+04	91%	0.0	86.14%	229934170	NC_029988.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1647	<i>Capsicum annuum</i>	4547	4,55E+03	84%	0.0	85.68%	1362880	NW_015960892.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4499	1,73E+04	95%	0.0	88.50%	153299543	NC_029984.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4492	2,53E+04	94%	0.0	85.79%	238794889	NC_029985.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold406	<i>Capsicum annuum</i>	4401	4,40E+03	85%	0.0	85.10%	80370	NW_015961461.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4244	5,52E+04	99%	0.0	85.03%	217274494	NC_029981.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4204	3,75E+04	94%	0.0	84.06%	222112641	NC_029983.1
CopiaCa_17	refeq-genome	8	Solanum pennellii chromosome 4, SPENNV200	<i>Solanum pennellii</i>	3951	9,25E+04	93%	0.0	84.22%	77197300	NC_028640.1
CopiaCa_17	refeq-genome	8	Solanum pennellii chromosome 5, SPENNV200	<i>Solanum pennellii</i>	3951	5,58E+04	93%	0.0	84.22%	77991103	NC_028641.1
CopiaCa_17	refeq-genome	8	Solanum pennellii chromosome 6, SPENNV200	<i>Solanum pennellii</i>	3945	1,14E+08	94%	0.0	84.19%	60730942	NC_028642.1
CopiaCa_17	refeq-genome	8	Solanum pennellii chromosome 1, SPENNV200	<i>Solanum pennellii</i>	3941	1,06E+08	93%	0.0	84.29%	109333515	NC_028637.1
CopiaCa_17	refeq-genome	8	Solanum pennellii chromosome 3, SPENNV200	<i>Solanum pennellii</i>	3940	7,77E+04	92%	0.0	84.16%	75414019	NC_028639.1

CopiaCa_17	refeq-genome	8	Solanum pennellii chromosome 10, SPENN200	<i>Solanum pennellii</i>	3934	1,04E+08	93%	0.0	84.26%	82529941	NC_028646.1
CopiaCa_17	refeq-genome	8	Solanum pennellii chromosome 8, SPENN200	<i>Solanum pennellii</i>	3928	6,04E+04	93%	0.0	84.11%	70546659	NC_028644.1
CopiaCa_17	refeq-genome	8	Solanum pennellii chromosome 9, SPENN200	<i>Solanum pennellii</i>	3928	7,00E+04	93%	0.0	84.11%	84057508	NC_028645.1
CopiaCa_17	refeq-genome	8	Solanum pennellii chromosome 2, SPENN200	<i>Solanum pennellii</i>	3925	3,63E+04	93%	0.0	84.10%	59803892	NC_028638.1
CopiaCa_17	refeq-genome	8	Solanum pennellii chromosome 7, SPENN200	<i>Solanum pennellii</i>	3923	5,54E+04	92%	0.0	84.21%	79292169	NC_028643.1
CopiaCa_17	refeq-genome	8	Solanum pennellii chromosome 12, SPENN200	<i>Solanum pennellii</i>	3923	4,83E+04	91%	0.0	84.08%	83305730	NC_028648.1
CopiaCa_17	refeq-genome	8	Solanum pennellii chromosome 11, SPENN200	<i>Solanum pennellii</i>	3919	5,73E+04	93%	0.0	84.07%	66223686	NC_028647.1
CopiaCa_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	<i>Solanum lycopersicum</i>	3786	7,25E+04	93%	0.0	83.48%	72906345	NC_015446.3
CopiaCa_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	<i>Solanum lycopersicum</i>	3733	4,18E+04	91%	0.0	83.23%	66557038	NC_015441.3
CopiaCa_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	<i>Solanum lycopersicum</i>	3703	5,98E+04	91%	0.0	83.12%	98455869	NC_015438.3
CopiaCa_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	<i>Solanum lycopersicum</i>	3701	5,93E+04	92%	0.0	83.22%	49794276	NC_015443.3
CopiaCa_17	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00337	<i>Solanum tuberosum</i>	3675	7,82E+03	88%	0.0	82.95%	664521	NW_006239266.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1754	<i>Capsicum annuum</i>	3651	5,27E+03	85%	0.0	81.96%	1029277	NW_015960393.1
CopiaCa_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	<i>Solanum lycopersicum</i>	3651	6,82E+04	93%	0.0	82.89%	65633393	NC_015447.3
CopiaCa_17	refeq-genome	8	Nicotiana attenuata strain UT chromosome 7, NIATTR2	<i>Nicotiana attenuata</i>	3607	5,56E+03	88%	0.0	82.79%	114765206	NC_031995.1
CopiaCa_17	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00207	<i>Solanum tuberosum</i>	3591	3,81E+03	80%	0.0	83.69%	1129882	NW_006239133.1
CopiaCa_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	<i>Solanum lycopersicum</i>	3579	3,75E+04	92%	0.0	82.59%	66723567	NC_015442.3
CopiaCa_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	<i>Solanum lycopersicum</i>	3559	4,02E+04	93%	0.0	82.53%	68126176	NC_015449.3
CopiaCa_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	<i>Solanum lycopersicum</i>	3531	4,99E+04	93%	0.0	82.46%	65987440	NC_015445.3
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold53	<i>Capsicum annuum</i>	3530	3,53E+03	82%	0.0	82.09%	1681775	NW_015961414.1
CopiaCa_17	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00085	<i>Solanum tuberosum</i>	3528	3,70E+03	83%	0.0	82.51%	1839851	NW_006239014.1
CopiaCa_17	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00020	<i>Solanum tuberosum</i>	3506	3,70E+03	83%	0.0	82.48%	3188724	NW_006238946.1
CopiaCa_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	<i>Solanum lycopersicum</i>	3483	5,19E+04	93%	0.0	82.65%	72290146	NC_015440.3
CopiaCa_17	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold78222	<i>Nicotiana sylvestris</i>	3448	3,45E+03	86%	0.0	80.87%	35264	NW_009568523.1
CopiaCa_17	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold72590	<i>Nicotiana tabacum</i>	3417	3,55E+03	83%	0.0	81.99%	17365	NW_015932837.1
CopiaCa_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	<i>Solanum lycopersicum</i>	3417	2,46E+04	90%	0.0	81.92%	56597135	NC_015448.3
CopiaCa_17	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold79001	<i>Nicotiana sylvestris</i>	3384	3,53E+03	83%	0.0	81.84%	8522	NW_009569389.1
CopiaCa_17	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1685	<i>Nicotiana tabacum</i>	3362	3,53E+03	82%	0.0	81.84%	110873	NW_015834268.1
CopiaCa_17	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold18579	<i>Nicotiana sylvestris</i>	3362	3,69E+03	83%	0.0	81.84%	126168	NW_009434124.1
CopiaCa_17	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00039	<i>Solanum tuberosum</i>	3301	5,77E+03	84%	0.0	81.57%	2527457	NW_006238968.1
CopiaCa_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	<i>Solanum lycopersicum</i>	3273	2,29E+04	92%	0.0	81.39%	55977580	NC_015439.3
CopiaCa_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	<i>Solanum lycopersicum</i>	3083	2,95E+04	93%	0.0	80.47%	68175699	NC_015444.3
CopiaCa_17	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00227	<i>Solanum tuberosum</i>	3062	3,17E+03	82%	0.0	80.47%	1069887	NW_006239150.1
CopiaCa_17	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00247	<i>Solanum tuberosum</i>	2953	3,65E+03	84%	0.0	81.76%	988620	NW_006239171.1
CopiaCa_17	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00297	<i>Solanum tuberosum</i>	2926	3,55E+03	84%	0.0	81.99%	814182	NW_006239220.1
CopiaCa_17	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00728	<i>Solanum tuberosum</i>	2920	3,95E+03	80%	0.0	83.55%	156345	NW_006239651.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	2811	4,55E+03	89%	0.0	85.10%	3493078	NW_015960517.1
CopiaCa_17	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00011	<i>Solanum tuberosum</i>	2658	2,66E+03	85%	0.0	77.89%	3646191	NW_006238938.1
CopiaCa_17	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold9927	<i>Nicotiana sylvestris</i>	2361	3,80E+03	83%	0.0	82.54%	247963	NW_009591908.1
CopiaCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1341	<i>Capsicum annuum</i>	2226	4,58E+03	85%	0.0	81.90%	1906221	NW_015960665.1
CopiaCa_17	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf01148	<i>Solanum tuberosum</i>	1531	2,72E+03	83%	0.0	78.94%	21312	NW_006240061.1
CopiaCa_18	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9217	1,83E+04	100%	0.0	100.00%	301019445	NC_029977.1
CopiaCa_18	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5936	1,12E+04	100%	0.0	93.08%	215701946	NC_029980.1
CopiaCa_18	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3917	4,22E+03	95%	0.0	83.50%	229934170	NC_029988.1
CopiaCa_18	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2348	6,03E+03	93%	0.0	84.01%	222112641	NC_029983.1
CopiaCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8440	1,68E+08	100%	0.0	100.00%	238794889	
CopiaCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5064	2,39E+08	100%	0.0	89.63%	301019445	
CopiaCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3788	1,33E+08	100%	0.0	84.06%	205736368	
CopiaCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3336	9,18E+04	98%	0.0	92.10%	163962470	
CopiaCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3262	1,76E+08	100%	0.0	91.55%	229934170	
CopiaCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2985	1,93E+08	100%	0.0	89.42%	217274494	
CopiaCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2939	1,04E+08	98%	0.0	87.71%	222112641	
CopiaCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2682	1,27E+08	100%	0.0	87.44%	219521584	
CopiaCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2588	1,51E+08	100%	0.0	88.72%	153299543	
CopiaCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2579	1,69E+08	100%	0.0	86.70%	261510930	
CopiaCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2274	1,70E+08	100%	0.0	84.11%	220335243	

CopiaCa_19	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	2056	1,40E+08	99%	0.0	86.17%	215701946	
CopiaCa_20	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	3336	9,18E+04	98%	0.0	92.10%	163962470	NC_029978.1
CopiaCa_20	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	2939	1,04E+08	98%	0.0	87.71%	222112641	NC_029983.1
CopiaCa_20	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	2056	1,40E+08	99%	0.0	86.17%	215701946	NC_029980.1
CopiaCa_20	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	8440	1,68E+08	100%	0.0	100.00%	238794889	NC_029985.1
CopiaCa_20	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5064	2,39E+08	100%	0.0	89.63%	301019445	NC_029977.1
CopiaCa_20	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	3788	1,33E+08	100%	0.0	84.06%	205736368	NC_029986.1
CopiaCa_20	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	3262	1,76E+08	100%	0.0	91.55%	229934170	NC_029988.1
CopiaCa_20	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	2985	1,93E+08	100%	0.0	89.42%	217274494	NC_029981.1
CopiaCa_20	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	2682	1,27E+08	100%	0.0	87.44%	219521584	NC_029982.1
CopiaCa_20	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	2588	1,51E+08	100%	0.0	88.72%	153299543	NC_029984.1
CopiaCa_20	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	2579	1,69E+08	100%	0.0	86.70%	261510930	NC_029979.1
CopiaCa_20	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	2274	1,70E+08	100%	0.0	84.11%	220335243	NC_029987.1
CopiaCa_21	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	9214	1,01E+04	100%	0.0	100.00%	220335243	NC_029987.1
CopiaCa_21	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4966	8,68E+03	100%	0.0	93.65%	261510930	NC_029979.1
CopiaCa_21	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4170	1,24E+04	95%	0.0	94.51%	215701946	NC_029980.1
CopiaCa_21	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4126	4,20E+03	92%	0.0	83.26%	222112641	NC_029983.1
CopiaCa_21	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	3779	7,63E+03	97%	0.0	91.53%	301019445	NC_029977.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	9503	3,24E+04	100%	0.0	100.00%	153299543	NC_029984.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7901	7,95E+04	100%	0.0	94.41%	301019445	NC_029977.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7696	5,37E+04	100%	0.0	93.69%	261510930	NC_029979.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold450	<i>Capisicum annuum</i>	7635	8,62E+03	100%	0.0	93.50%	175795	NW_015961071.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7474	6,41E+04	100%	0.0	93.02%	219521584	NC_029982.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7315	6,68E+04	100%	0.0	92.41%	205736368	NC_029986.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7306	5,01E+04	100%	0.0	92.38%	229934170	NC_029988.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7293	3,94E+04	100%	0.0	93.02%	238794889	NC_029985.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	6920	6,99E+04	100%	0.0	91.08%	220335243	NC_029987.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	6818	3,21E+04	100%	0.0	90.50%	222112641	NC_029983.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	6560	4,33E+04	100%	0.0	92.89%	215701946	NC_029980.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	6078	3,17E+04	100%	0.0	88.10%	163962470	NC_029978.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold366	<i>Capisicum annuum</i>	6011	6,38E+03	99%	0.0	87.85%	343865	NW_015960794.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4556	4,59E+04	100%	0.0	93.48%	217274494	NC_029981.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1894	<i>Capisicum annuum</i>	3868	7,11E+03	92%	0.0	90.27%	2334823	NW_015961490.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2804	<i>Capisicum annuum</i>	2970	3,60E+03	86%	0.0	80.87%	96601	NW_015961341.1
CopiaCa_22	refeq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold15099	<i>Capisicum annuum</i>	2182	6,96E+03	80%	0.0	91.51%	15847	NW_015960462.1
CopiaCa_23	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	9919	4,68E+04	100%	0.0	99.98%	301019445	NC_029977.1
CopiaCa_23	refeq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4695	<i>Capisicum annuum</i>	7803	9,84E+03	99%	0.0	92.96%	183029	NW_015961566.1
CopiaCa_23	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7625	7,63E+04	100%	0.0	92.51%	261510930	NC_029979.1
CopiaCa_23	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7480	2,69E+04	99%	0.0	91.98%	238794889	NC_029985.1
CopiaCa_23	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	7134	4,40E+04	99%	0.0	93.02%	229934170	NC_029988.1
CopiaCa_23	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	6944	7,62E+04	99%	0.0	92.34%	220335243	NC_029987.1
CopiaCa_23	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	6023	2,06E+04	99%	0.0	90.18%	153299543	NC_029984.1
CopiaCa_23	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5842	6,17E+04	100%	0.0	89.66%	217274494	NC_029981.1
CopiaCa_23	refeq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3745	<i>Capisicum annuum</i>	5653	6,63E+03	82%	0.0	93.70%	97749	NW_015961051.1
CopiaCa_23	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	5345	3,00E+04	100%	0.0	90.52%	163962470	NC_029978.1
CopiaCa_23	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 S16 R44 unplaced genomic scaffold, SolTub_3.0 scf0043	<i>Solanum tuberosum</i>	4813	5,12E+03	89%	0.0	86.19%	2504196	NW_006238969.1
CopiaCa_23	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capisicum annuum</i>	4802	1,58E+04	92%	0.0	86.82%	219521584	NC_029982.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3312	<i>Nicotiana tomentosiformis</i>	4767	4,77E+03	83%	0.0	85.83%	423229	NW_008913731.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold26271	<i>Nicotiana tabacum</i>	4761	9,43E+03	84%	0.0	85.76%	56916	NW_015881256.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13639	<i>Nicotiana tomentosiformis</i>	4758	4,76E+03	82%	0.0	86.11%	337317	NW_008868924.1
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	4700	4,70E+03	82%	0.0	85.92%	286979	NW_017671503.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold50014	<i>Nicotiana tabacum</i>	4697	4,70E+03	84%	0.0	85.52%	226704	NW_015912758.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold15198	<i>Nicotiana tomentosiformis</i>	4632	4,63E+03	83%	0.0	85.30%	103115	NW_008886246.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold77778	<i>Nicotiana tabacum</i>	4623	4,62E+03	84%	0.0	85.22%	46642	NW_015937274.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold8869	<i>Nicotiana tomentosiformis</i>	4595	4,60E+03	84%	0.0	85.16%	68433	NW_008975473.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold113902	<i>Nicotiana tabacum</i>	4590	4,59E+03	83%	0.0	85.10%	78534	NW_015797853.1

CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9861	<i>Nicotiana tabacum</i>	4567	4,57E+03	84%	0.0	85.02%	71947	NW_015954368.1
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT chromosome 2, NIATT2r	<i>Nicotiana attenuata</i>	4554	8,72E+03	82%	0.0	85.34%	96359640	NC_031990.1
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT chromosome 10, NIATT2r	<i>Nicotiana attenuata</i>	4551	8,87E+03	88%	0.0	85.38%	51082206	NC_031998.1
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2r	<i>Nicotiana attenuata</i>	4529	4,53E+03	82%	0.0	85.22%	1005419	NW_017670323.1
CopiaCa_23	refeq-genome	8	Capicum annum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capicum annum</i>	4497	1,84E+04	99%	0.0	84.87%	205736368	NC_029986.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17472	<i>Nicotiana tabacum</i>	4492	4,49E+03	80%	0.0	85.41%	344968	NW_015837843.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold45236	<i>Nicotiana tabacum</i>	4482	4,48E+03	82%	0.0	85.02%	55088	NW_015908360.1
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2r	<i>Nicotiana attenuata</i>	4475	4,48E+03	82%	0.0	85.11%	118424	NW_017673826.1
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT chromosome 7, NIATT2r	<i>Nicotiana attenuata</i>	4451	1,09E+04	82%	0.0	84.94%	114765206	NC_031995.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13769	<i>Nicotiana tabacum</i>	4447	4,45E+03	84%	0.0	84.53%	191847	NW_015814814.1
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2r	<i>Nicotiana attenuata</i>	4440	4,44E+03	82%	0.0	85.02%	255029	NW_017671743.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold14058	<i>Nicotiana tabacum</i>	4429	4,43E+03	84%	0.0	84.46%	106415	NW_015816763.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold3341	<i>Nicotiana tabacum</i>	4414	4,41E+03	82%	0.0	84.82%	132437	NW_015891269.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold41129	<i>Nicotiana tabacum</i>	4414	4,41E+03	82%	0.0	84.74%	150958	NW_015904582.1
CopiaCa_23	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold4960	<i>Nicotiana sylvestris</i>	4397	5,34E+03	82%	0.0	84.56%	38260	NW_009536720.1
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2r	<i>Nicotiana attenuata</i>	4394	4,39E+03	83%	0.0	84.55%	107891	NW_017674123.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold39061	<i>Nicotiana tabacum</i>	4394	4,39E+03	80%	0.0	85.00%	72504	NW_015902667.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold44184	<i>Nicotiana tabacum</i>	4394	8,96E+03	90%	0.0	84.66%	205495	NW_015907404.1
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT chromosome 1, NIATT2r	<i>Nicotiana attenuata</i>	4390	8,54E+03	86%	0.0	84.72%	110538649	NC_031989.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold117898	<i>Nicotiana tabacum</i>	4390	4,79E+03	82%	0.0	84.55%	44475	NW_015800784.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1098	<i>Nicotiana tabacum</i>	4386	4,71E+03	91%	0.0	84.32%	341244	NW_015794785.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold35690	<i>Nicotiana tabacum</i>	4385	4,39E+03	84%	0.0	84.28%	49854	NW_015899509.1
CopiaCa_23	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold941	<i>Nicotiana sylvestris</i>	4385	4,39E+03	82%	0.0	84.57%	55655	NW_009586164.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2975	<i>Nicotiana tomentosiformis</i>	4385	4,39E+03	81%	0.0	84.93%	82171	NW_008909985.1
CopiaCa_23	refeq-genome	8	Capicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold899	<i>Capicum annum</i>	4379	7,04E+03	82%	0.0	93.14%	265334	NW_015961225.1
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2r	<i>Nicotiana attenuata</i>	4377	4,38E+03	83%	0.0	84.48%	38657	NW_017676818.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24788	<i>Nicotiana tabacum</i>	4377	4,38E+03	82%	0.0	84.61%	125848	NW_015874442.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold62478	<i>Nicotiana tabacum</i>	4377	4,38E+03	84%	0.0	84.26%	134539	NW_015924003.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold19939	<i>Nicotiana tabacum</i>	4373	4,37E+03	82%	0.0	84.52%	21566	NW_015851309.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold77431	<i>Nicotiana tabacum</i>	4370	4,85E+03	81%	0.0	85.13%	65915	NW_015936970.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23399	<i>Nicotiana tabacum</i>	4362	4,36E+03	84%	0.0	84.24%	59311	NW_015868045.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold42688	<i>Nicotiana tabacum</i>	4359	4,36E+03	84%	0.0	84.20%	235262	NW_015906017.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold22705	<i>Nicotiana tomentosiformis</i>	4346	4,35E+03	84%	0.0	84.13%	149011	NW_008902159.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold14138	<i>Nicotiana tabacum</i>	4342	4,34E+03	84%	0.0	84.19%	182785	NW_015817312.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold22128	<i>Nicotiana tabacum</i>	4335	4,34E+03	84%	0.0	84.16%	109579	NW_015862097.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23943	<i>Nicotiana tabacum</i>	4335	4,34E+03	83%	0.0	84.29%	41566	NW_015870614.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold29615	<i>Nicotiana tabacum</i>	4327	4,33E+03	82%	0.0	84.33%	90106	NW_015887683.1
CopiaCa_23	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold98422	<i>Nicotiana sylvestris</i>	4324	4,32E+03	82%	0.0	84.32%	53306	NW_009590967.1
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2r	<i>Nicotiana attenuata</i>	4322	4,32E+03	82%	0.0	84.53%	523967	NW_017670641.1
CopiaCa_23	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold7260	<i>Nicotiana sylvestris</i>	4318	4,32E+03	82%	0.0	84.48%	131682	NW_009562276.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold14312	<i>Nicotiana tabacum</i>	4311	4,31E+03	82%	0.0	84.40%	226614	NW_015818477.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold6845	<i>Nicotiana tomentosiformis</i>	4311	4,31E+03	82%	0.0	84.33%	178439	NW_008952985.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1472	<i>Nicotiana tabacum</i>	4309	4,31E+03	82%	0.0	84.42%	38084	NW_015821125.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold99194	<i>Nicotiana tabacum</i>	4309	4,31E+03	84%	0.0	83.98%	44841	NW_015954827.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold20029	<i>Nicotiana tabacum</i>	4301	4,30E+03	83%	0.0	83.98%	89399	NW_015851764.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold35267	<i>Nicotiana tabacum</i>	4289	4,29E+03	82%	0.0	84.21%	82909	NW_015899107.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold99162	<i>Nicotiana tabacum</i>	4289	4,29E+03	82%	0.0	84.26%	49315	NW_015954810.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold18130	<i>Nicotiana tabacum</i>	4287	4,29E+03	82%	0.0	84.22%	36644	NW_015841532.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold15517	<i>Nicotiana tabacum</i>	4283	4,56E+03	89%	0.0	83.97%	67428	NW_015826193.1
CopiaCa_23	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold15697	<i>Nicotiana sylvestris</i>	4283	4,28E+03	82%	0.0	84.22%	71797	NW_009402102.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold83337	<i>Nicotiana tabacum</i>	4281	4,28E+03	82%	0.0	84.22%	53677	NW_015941973.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold7636	<i>Nicotiana tomentosiformis</i>	4276	4,28E+03	84%	0.0	83.89%	14045	NW_008961774.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold14047	<i>Nicotiana tomentosiformis</i>	4272	5,15E+03	82%	0.0	84.56%	44952	NW_008873458.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold12817	<i>Nicotiana tomentosiformis</i>	4266	4,27E+03	84%	0.0	83.82%	436665	NW_008859791.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold39950	<i>Nicotiana tabacum</i>	4259	4,26E+03	84%	0.0	83.94%	72826	NW_015903486.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold105390	<i>Nicotiana tabacum</i>	4253	4,25E+03	84%	0.0	83.79%	65063	NW_015791428.1

CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold3816	<i>Nicotiana tabacum</i>	4241	4,24E+03	83%	0.0	83.83%	32842	NW_015901833.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold46638	<i>Nicotiana tabacum</i>	4239	4,24E+03	82%	0.0	84.05%	101065	NW_015909668.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold7940	<i>Nicotiana tomentosiformis</i>	4239	4,24E+03	82%	0.0	84.06%	69171	NW_008965152.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold15094	<i>Nicotiana tabacum</i>	4229	8,32E+03	82%	0.0	84.00%	128357	NW_015823466.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9230	<i>Nicotiana tabacum</i>	4228	4,23E+03	84%	0.0	83.70%	93700	NW_015949377.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold43213	<i>Nicotiana tabacum</i>	4226	4,23E+03	83%	0.0	83.73%	76524	NW_015906508.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13184	<i>Nicotiana tabacum</i>	4220	4,22E+03	84%	0.0	83.69%	123229	NW_015810784.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold16831	<i>Nicotiana tabacum</i>	4220	4,22E+03	81%	0.0	84.23%	169150	NW_015834148.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1033	<i>Nicotiana tomentosiformis</i>	4220	4,52E+03	89%	0.0	83.72%	196095	NW_008832160.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1267	<i>Nicotiana tabacum</i>	4209	4,21E+03	82%	0.0	83.93%	113957	NW_015807161.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold136176	<i>Nicotiana tabacum</i>	4207	6,53E+03	84%	0.0	83.60%	58202	NW_015813754.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold87539	<i>Nicotiana tabacum</i>	4207	4,21E+03	84%	0.0	83.63%	46761	NW_015945453.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold25576	<i>Nicotiana tabacum</i>	4204	4,20E+03	84%	0.0	83.62%	128471	NW_015877976.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold41351	<i>Nicotiana tabacum</i>	4204	4,47E+03	88%	0.0	83.96%	54750	NW_015904787.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold49214	<i>Nicotiana tabacum</i>	4202	4,20E+03	84%	0.0	83.63%	44441	NW_015912030.1
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	4200	4,20E+03	82%	0.0	83.95%	426980	NW_017670865.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold25158	<i>Nicotiana tabacum</i>	4200	4,20E+03	84%	0.0	83.67%	79416	NW_015876075.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold28051	<i>Nicotiana tomentosiformis</i>	4200	4,20E+03	82%	0.0	83.90%	38680	NW_008908099.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold51493	<i>Nicotiana tabacum</i>	4198	4,20E+03	82%	0.0	83.90%	41816	NW_015914099.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9932	<i>Nicotiana tabacum</i>	4198	7,20E+03	88%	0.0	83.90%	184704	NW_015954932.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold52245	<i>Nicotiana tabacum</i>	4196	4,20E+03	84%	0.0	83.64%	139341	NW_015914800.1
CopiaCa_23	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsy Nsy_scaffold1827	<i>Nicotiana sylvestris</i>	4196	4,20E+03	82%	0.0	83.89%	352154	NW_009430691.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold17434	<i>Nicotiana tomentosiformis</i>	4196	6,59E+03	82%	0.0	83.87%	115607	NW_008896302.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold25025	<i>Nicotiana tomentosiformis</i>	4194	4,19E+03	82%	0.0	83.85%	24943	NW_008904737.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold104383	<i>Nicotiana tabacum</i>	4193	4,95E+03	91%	0.0	83.65%	30580	NW_015790635.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold39257	<i>Nicotiana tabacum</i>	4191	4,19E+03	82%	0.0	83.87%	44533	NW_015902849.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold24828	<i>Nicotiana tomentosiformis</i>	4187	4,19E+03	84%	0.0	83.55%	20520	NW_008904517.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5049	<i>Nicotiana tomentosiformis</i>	4187	4,19E+03	82%	0.0	83.85%	38265	NW_008933030.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold48787	<i>Nicotiana tomentosiformis</i>	4183	4,18E+03	84%	0.0	83.55%	31223	NW_008931137.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold14146	<i>Nicotiana tomentosiformis</i>	4180	4,18E+03	83%	0.0	83.59%	18937	NW_008874558.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4000	<i>Nicotiana tomentosiformis</i>	4172	4,17E+03	84%	0.0	83.45%	627986	NW_008921376.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold120921	<i>Nicotiana tabacum</i>	4170	4,17E+03	82%	0.0	83.80%	18226	NW_015802986.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10039	<i>Nicotiana tabacum</i>	4169	4,17E+03	83%	0.0	83.55%	176966	NW_015787542.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10952	<i>Nicotiana tabacum</i>	4169	4,17E+03	81%	0.0	84.02%	26389	NW_015794579.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold29220	<i>Nicotiana tomentosiformis</i>	4169	5,04E+03	88%	0.0	83.79%	290280	NW_008909398.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold104887	<i>Nicotiana tabacum</i>	4165	4,44E+03	90%	0.0	83.43%	69546	NW_015791025.1
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	4163	4,16E+03	82%	0.0	83.78%	137678	NW_017673397.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold2441	<i>Nicotiana tabacum</i>	4163	4,16E+03	81%	0.0	83.99%	310619	NW_015872755.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3340	<i>Nicotiana tomentosiformis</i>	4157	4,43E+03	88%	0.0	83.79%	152482	NW_008914042.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9568	<i>Nicotiana tomentosiformis</i>	4154	4,15E+03	81%	0.0	83.90%	92013	NW_008983240.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold15228	<i>Nicotiana tabacum</i>	4152	4,15E+03	80%	0.0	84.10%	50820	NW_015824356.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3008	<i>Nicotiana tomentosiformis</i>	4152	4,15E+03	83%	0.0	83.49%	42402	NW_008910353.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold20606	<i>Nicotiana tabacum</i>	4145	5,01E+03	88%	0.0	83.70%	187389	NW_015854711.1
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	4139	4,14E+03	82%	0.0	83.79%	222226	NW_017672067.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold60074	<i>Nicotiana tomentosiformis</i>	4139	4,14E+03	83%	0.0	83.51%	37093	NW_008943680.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7380	<i>Nicotiana tabacum</i>	4132	4,13E+03	83%	0.0	83.38%	120098	NW_015933858.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1654	<i>Nicotiana tomentosiformis</i>	4128	8,29E+03	91%	0.0	83.32%	225051	NW_008895308.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold54820	<i>Nicotiana tabacum</i>	4124	4,32E+03	83%	0.0	84.15%	29679	NW_015917158.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5768	<i>Nicotiana tomentosiformis</i>	4120	4,12E+03	83%	0.0	83.42%	38625	NW_008941018.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold21614	<i>Nicotiana tabacum</i>	4109	4,11E+03	82%	0.0	83.65%	79935	NW_015859602.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold11357	<i>Nicotiana tomentosiformis</i>	4096	4,38E+03	91%	0.0	83.20%	178497	NW_008843569.1
CopiaCa_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4095	2,55E+04	94%	0.0	87.64%	222112641	NC_029983.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17434	<i>Nicotiana tabacum</i>	4091	4,87E+03	82%	0.0	83.72%	146677	NW_015837630.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold27514	<i>Nicotiana tomentosiformis</i>	4091	4,09E+03	83%	0.0	83.33%	47801	NW_008907502.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9827	<i>Nicotiana tomentosiformis</i>	4089	4,85E+03	83%	0.0	83.41%	134862	NW_008986118.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17200	<i>Nicotiana tabacum</i>	4085	4,63E+03	90%	0.0	83.16%	114133	NW_015836286.1

CopiaCa_23	refeq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold258
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1701
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16983
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5452
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold15091
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold96935
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5418
CopiaCa_23	refeq-genome	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold7910
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold38073
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold38411
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold58971
CopiaCa_23	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold49998
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold33146
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold6456
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3398
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold97043
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT chromosome 3, NIATT2
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold17294
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7528
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold24683
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold27552
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold42559
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold748
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold69670
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23203
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold34354
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold61242
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold32902
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT chromosome 11, NIATT2
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4553
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold39882
CopiaCa_23	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold14153
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold28256
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13096
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold53431
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2844
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5546
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13254
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT chromosome 9, NIATT2
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold57661
CopiaCa_23	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold21601
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold8448
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold27863
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold22370
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold53792
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold14924
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold74528
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1715
CopiaCa_23	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold27485
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT chromosome 6, NIATT2
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16562
CopiaCa_23	refeq-genome	8	Capisicum annum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0

<i>Capisicum annum</i>	4084	7,10E+03	92%	0.0	89.71%	3098259	NW_015961180.1
<i>Nicotiana tabacum</i>	4078	4,08E+03	84%	0.0	83.16%	78572	NW_015835234.1
<i>Nicotiana tomentosiformis</i>	4074	4,07E+03	82%	0.0	83.40%	50650	NW_008895800.1
<i>Nicotiana tabacum</i>	4069	4,53E+03	88%	0.0	83.34%	104894	NW_015916888.1
<i>Nicotiana tomentosiformis</i>	4067	4,96E+03	88%	0.0	83.39%	64628	NW_008885058.1
<i>Nicotiana tabacum</i>	4065	4,07E+03	82%	0.0	83.33%	36587	NW_015953035.1
<i>Nicotiana tabacum</i>	4056	6,33E+03	82%	0.0	83.36%	31259	NW_015916578.1
<i>Capisicum annum</i>	4056	9,52E+03	99%	0.0	93.03%	96418	NW_015961469.1
<i>Nicotiana tomentosiformis</i>	4034	4,03E+03	80%	0.0	83.61%	116484	NW_008919234.1
<i>Nicotiana tabacum</i>	4030	4,03E+03	81%	0.0	83.46%	42899	NW_015902064.1
<i>Nicotiana tabacum</i>	4019	4,53E+03	88%	0.0	84.25%	35548	NW_015920877.1
<i>Nicotiana sylvestris</i>	4019	4,53E+03	87%	0.0	84.31%	63176	NW_009537161.1
<i>Nicotiana tomentosiformis</i>	4019	4,02E+03	82%	0.0	83.28%	57481	NW_008913760.1
<i>Nicotiana tomentosiformis</i>	4012	4,82E+03	84%	0.0	86.91%	96054	NW_008948663.1
<i>Nicotiana tomentosiformis</i>	4006	4,01E+03	83%	0.0	82.88%	129453	NW_008914685.1
<i>Nicotiana tabacum</i>	4004	4,00E+03	84%	0.0	82.84%	12295	NW_015953131.1
<i>Nicotiana attenuata</i>	3997	1,24E+04	82%	0.0	85.11%	72893213	NC_031991.1
<i>Nicotiana attenuata</i>	3997	4,46E+03	89%	0.0	83.18%	207768	NW_017672228.1
<i>Nicotiana tomentosiformis</i>	3997	4,00E+03	84%	0.0	82.84%	185449	NW_008896146.1
<i>Nicotiana tabacum</i>	3989	4,88E+03	91%	0.0	82.86%	191274	NW_015935130.1
<i>Nicotiana tomentosiformis</i>	3989	3,99E+03	84%	0.0	82.86%	153191	NW_008904356.1
<i>Nicotiana tomentosiformis</i>	3980	3,98E+03	82%	0.0	83.09%	59904	NW_008907544.1
<i>Nicotiana tabacum</i>	3978	3,98E+03	84%	0.0	82.79%	77038	NW_015905902.1
<i>Nicotiana tabacum</i>	3975	7,50E+03	89%	0.0	82.94%	62261	NW_015934725.1
<i>Nicotiana tabacum</i>	3973	4,25E+03	84%	0.0	82.80%	122493	NW_015930303.1
<i>Nicotiana tabacum</i>	3956	3,96E+03	82%	0.0	82.99%	113944	NW_015867112.1
<i>Nicotiana tabacum</i>	3956	3,96E+03	82%	0.0	83.02%	157734	NW_015892155.1
<i>Nicotiana tomentosiformis</i>	3951	3,95E+03	84%	0.0	82.64%	9627	NW_008944978.1
<i>Nicotiana tabacum</i>	3936	3,94E+03	83%	0.0	82.65%	60418	NW_015890778.1
<i>Nicotiana attenuata</i>	3932	3,93E+03	81%	0.0	82.97%	171278	NW_017672755.1
<i>Nicotiana attenuata</i>	3930	3,93E+03	82%	0.0	82.78%	61252062	NC_031999.1
<i>Nicotiana attenuata</i>	3925	4,50E+03	86%	0.0	85.35%	192159	NW_017672438.1
<i>Nicotiana tabacum</i>	3925	3,93E+03	82%	0.0	82.87%	141234	NW_015908631.1
<i>Nicotiana tabacum</i>	3919	4,48E+03	84%	0.0	84.52%	119430	NW_015903426.1
<i>Nicotiana sylvestris</i>	3906	3,91E+03	83%	0.0	82.44%	54331	NW_009384948.1
<i>Nicotiana tomentosiformis</i>	3906	4,68E+03	81%	0.0	86.09%	101219	NW_008908326.1
<i>Nicotiana tomentosiformis</i>	3899	3,90E+03	82%	0.0	82.80%	84601	NW_008862891.1
<i>Nicotiana tabacum</i>	3890	3,89E+03	82%	0.0	82.76%	85581	NW_015915895.1
<i>Nicotiana tomentosiformis</i>	3869	3,87E+03	81%	0.0	82.87%	243778	NW_008908530.1
<i>Nicotiana tomentosiformis</i>	3868	4,77E+03	84%	0.0	84.12%	99254	NW_008938552.1
<i>Nicotiana attenuata</i>	3842	3,84E+03	82%	0.0	82.41%	924530	NW_017670347.1
<i>Nicotiana tomentosiformis</i>	3808	3,81E+03	82%	0.0	82.48%	110959	NW_008864647.1
<i>Nicotiana attenuata</i>	3801	4,92E+03	88%	0.0	82.87%	50975570	NC_031997.1
<i>Nicotiana tabacum</i>	3792	4,33E+03	86%	0.0	82.68%	73334	NW_015919729.1
<i>Nicotiana sylvestris</i>	3775	4,31E+03	86%	0.0	82.61%	327418	NW_009467704.1
<i>Nicotiana tomentosiformis</i>	3759	3,76E+03	83%	0.0	82.18%	39382	NW_008970796.1
<i>Nicotiana tabacum</i>	3747	3,75E+03	82%	0.0	82.23%	163106	NW_015886036.1
<i>Nicotiana tomentosiformis</i>	3740	7,02E+03	82%	0.0	83.60%	67692	NW_008901787.1
<i>Nicotiana tomentosiformis</i>	3731	3,73E+03	84%	0.0	81.92%	15935	NW_008936699.1
<i>Nicotiana tabacum</i>	3718	4,83E+03	83%	0.0	82.03%	40622	NW_015822408.1
<i>Nicotiana tabacum</i>	3683	4,59E+03	82%	0.0	84.56%	61302	NW_015934492.1
<i>Nicotiana tomentosiformis</i>	3657	4,25E+03	82%	0.0	84.53%	310934	NW_008959586.1
<i>Nicotiana sylvestris</i>	3633	3,63E+03	81%	0.0	81.82%	59671	NW_009512148.1
<i>Nicotiana attenuata</i>	3629	1,16E+04	84%	0.0	82.28%	97571839	NC_031994.1
<i>Nicotiana tomentosiformis</i>	3544	4,34E+03	91%	0.0	83.63%	32233	NW_008895333.1
<i>Capisicum annum</i>	3520	2,20E+04	98%	0.0	88.41%	215701946	NC_029980.1

CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold32132	<i>Nicotiana tabacum</i>	3478	4,21E+03	82%	0.0	83.81%	83779	NW_015890061.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2924	<i>Nicotiana tomentosiformis</i>	3456	4,20E+03	82%	0.0	83.70%	36988	NW_008909419.1
CopiaCa_23	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00031	<i>Solanum tuberosum</i>	3404	5,96E+03	88%	0.0	86.57%	2735125	NW_006238957.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold60349	<i>Nicotiana tabacum</i>	3374	4,14E+03	82%	0.0	84.68%	19147	NW_015922099.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9845	<i>Nicotiana tomentosiformis</i>	3363	5,49E+03	82%	0.0	85.60%	42876	NW_008986318.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold29262	<i>Nicotiana tabacum</i>	3299	4,81E+03	88%	0.0	84.06%	111307	NW_015887349.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold27597	<i>Nicotiana tabacum</i>	3205	4,90E+03	91%	0.0	83.72%	190851	NW_015885772.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold15602	<i>Nicotiana tomentosiformis</i>	3188	3,98E+03	81%	0.0	83.04%	181066	NW_008890736.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold22468	<i>Nicotiana tomentosiformis</i>	3101	4,35E+03	84%	0.0	82.88%	40133	NW_008901895.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold6148	<i>Nicotiana tabacum</i>	3086	4,14E+03	81%	0.0	84.65%	137436	NW_015923102.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold27360	<i>Nicotiana tomentosiformis</i>	3070	4,08E+03	82%	0.0	84.64%	22523	NW_008907331.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1685	<i>Nicotiana tomentosiformis</i>	3068	4,64E+03	84%	0.0	83.74%	262938	NW_008895652.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2406	<i>Nicotiana tomentosiformis</i>	3027	4,04E+03	82%	0.0	83.87%	194317	NW_008903664.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold15086	<i>Nicotiana tabacum</i>	3009	4,87E+03	85%	0.0	85.14%	37718	NW_015823417.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold11681	<i>Nicotiana tabacum</i>	3005	4,01E+03	82%	0.0	83.87%	132606	NW_015800001.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold37558	<i>Nicotiana tomentosiformis</i>	2998	4,30E+03	87%	0.0	84.43%	28999	NW_008918661.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold39138	<i>Nicotiana tomentosiformis</i>	2957	6,79E+03	90%	0.0	83.94%	28341	NW_008920417.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold50667	<i>Nicotiana tabacum</i>	2953	3,70E+03	81%	0.0	83.16%	119393	NW_015913353.1
CopiaCa_23	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsy1 Nsy1_scaffold45693	<i>Nicotiana sylvestris</i>	2953	3,70E+03	81%	0.0	83.16%	76055	NW_009532379.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold19321	<i>Nicotiana tomentosiformis</i>	2953	4,47E+03	82%	0.0	84.43%	34884	NW_008898399.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8727	<i>Nicotiana tabacum</i>	2946	6,95E+03	91%	0.0	84.87%	118427	NW_015945235.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold49708	<i>Nicotiana tabacum</i>	2937	4,48E+03	84%	0.0	83.23%	339750	NW_015912474.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold59279	<i>Nicotiana tabacum</i>	2926	4,33E+03	81%	0.0	83.91%	141922	NW_015921142.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold28866	<i>Nicotiana tabacum</i>	2859	4,18E+03	83%	0.0	82.86%	81388	NW_015886973.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10234	<i>Nicotiana tabacum</i>	2819	4,39E+03	82%	0.0	85.75%	338974	NW_015789061.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold20360	<i>Nicotiana tomentosiformis</i>	2796	4,61E+03	91%	0.0	85.24%	78646	NW_008899554.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold31018	<i>Nicotiana tabacum</i>	2789	4,34E+03	83%	0.0	85.16%	77295	NW_015889007.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16409	<i>Nicotiana tomentosiformis</i>	2772	4,35E+03	83%	0.0	83.63%	79137	NW_008895163.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17610	<i>Nicotiana tabacum</i>	2728	4,32E+03	83%	0.0	85.49%	47695	NW_015838618.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13941	<i>Nicotiana tabacum</i>	2712	4,42E+03	80%	0.0	82.65%	189720	NW_015815959.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold12852	<i>Nicotiana tomentosiformis</i>	2704	4,35E+03	82%	0.0	83.28%	112275	NW_008860180.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold39014	<i>Nicotiana tabacum</i>	2693	8,43E+03	82%	0.0	85.47%	85645	NW_015902622.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3500	<i>Nicotiana tomentosiformis</i>	2686	4,37E+03	91%	0.0	82.55%	183243	NW_008915820.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5153	<i>Nicotiana tomentosiformis</i>	2686	4,18E+03	84%	0.0	82.37%	113207	NW_008934186.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold34342	<i>Nicotiana tabacum</i>	2669	4,13E+03	84%	0.0	82.28%	77082	NW_015892143.1
CopiaCa_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capsicum annuum</i>	2647	1,22E+04	97%	0.0	93.06%	2397553	NW_015961318.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1859	<i>Nicotiana tomentosiformis</i>	2595	3,73E+03	80%	0.0	82.02%	82017	NW_008897585.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold10055	<i>Nicotiana tomentosiformis</i>	2582	4,41E+03	84%	0.0	82.79%	34584	NW_008829105.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold32186	<i>Nicotiana tomentosiformis</i>	2569	5,53E+03	82%	0.0	86.71%	146298	NW_008912693.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13768	<i>Nicotiana tomentosiformis</i>	2529	4,52E+03	84%	0.0	85.05%	105966	NW_008870357.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold7702	<i>Nicotiana tomentosiformis</i>	2492	4,07E+03	83%	0.0	83.91%	115835	NW_008962508.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold6697	<i>Nicotiana tomentosiformis</i>	2484	5,63E+03	88%	0.0	82.34%	47924	NW_008951340.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold114148	<i>Nicotiana tabacum</i>	2473	4,43E+03	84%	0.0	83.10%	16839	NW_015798048.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold70901	<i>Nicotiana tabacum</i>	2455	4,33E+03	84%	0.0	84.70%	52426	NW_015931375.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold22940	<i>Nicotiana tabacum</i>	2436	4,20E+03	80%	0.0	86.27%	76236	NW_015865915.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold12089	<i>Nicotiana tabacum</i>	2420	4,53E+03	81%	0.0	84.95%	86612	NW_015802956.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold7133	<i>Nicotiana tomentosiformis</i>	2403	4,31E+03	82%	0.0	83.09%	60173	NW_008956186.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8306	<i>Nicotiana tabacum</i>	2379	4,28E+03	82%	0.0	82.87%	85779	NW_015941741.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5433	<i>Nicotiana tomentosiformis</i>	2372	4,89E+03	90%	0.0	84.07%	137307	NW_00893297.1
CopiaCa_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5566	<i>Capsicum annuum</i>	2362	1,03E+04	90%	0.0	92.45%	186051	NW_015960765.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold37901	<i>Nicotiana tabacum</i>	2350	4,47E+03	84%	0.0	85.48%	12230	NW_015901576.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold88767	<i>Nicotiana tabacum</i>	2331	4,31E+03	88%	0.0	81.94%	34630	NW_015946459.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold34388	<i>Nicotiana tabacum</i>	2316	4,17E+03	82%	0.0	82.91%	33290	NW_015892185.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold14895	<i>Nicotiana tomentosiformis</i>	2300	4,16E+03	82%	0.0	82.80%	125663	NW_008882879.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold21580	<i>Nicotiana tomentosiformis</i>	2296	7,69E+03	91%	0.0	84.97%	124788	NW_008900909.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold443	<i>Nicotiana tomentosiformis</i>	2289	4,69E+03	88%	0.0	84.97%	91647	NW_008926152.1

CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7074	<i>Nicotiana tabacum</i>	2285	4,41E+03	90%	0.0	85.16%	145082	NW_015931223.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold40570	<i>Nicotiana tabacum</i>	2209	4,28E+03	81%	0.0	85.38%	70498	NW_015904048.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold26686	<i>Nicotiana tomentosiformis</i>	2196	4,21E+03	83%	0.0	82.41%	30193	NW_008906581.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold66881	<i>Nicotiana tabacum</i>	2137	4,28E+03	80%	0.0	86.18%	86455	NW_015927884.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24001	<i>Nicotiana tabacum</i>	2126	4,21E+03	80%	0.0	84.61%	228954	NW_015870881.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold137110	<i>Nicotiana tomentosiformis</i>	2115	4,41E+03	80%	0.0	82.56%	198271	NW_008946929.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold63	<i>Nicotiana tabacum</i>	2067	4,06E+03	81%	0.0	84.47%	207694	NW_015885126.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold26913	<i>Nicotiana tomentosiformis</i>	2054	4,95E+03	88%	0.0	81.89%	37248	NW_008839037.1
CopiaCa_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	<i>Nicotiana attenuata</i>	2039	3,68E+03	82%	0.0	83.91%	150974	NW_017673108.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold15777	<i>Nicotiana tomentosiformis</i>	1986	6,33E+03	83%	0.0	87.10%	17460	NW_008892679.1
CopiaCa_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold12883	<i>Nicotiana tomentosiformis</i>	1967	4,09E+03	80%	0.0	82.65%	123148	NW_008860524.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold19855	<i>Nicotiana tabacum</i>	1921	4,60E+03	89%	0.0	86.16%	95589	NW_015850863.1
CopiaCa_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold11196	<i>Nicotiana tabacum</i>	1888	4,04E+03	80%	0.0	82.12%	99269	NW_015796410.1
CopiaCa_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8486	2,20E+04	100%	0.0	100.00%	261510930	NC_029979.1
CopiaCa_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6682	2,20E+04	100%	0.0	92.99%	220335243	NC_029987.1
CopiaCa_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold619	<i>Capsicum annuum</i>	6433	6,83E+03	100%	0.0	91.92%	2512642	NW_015960498.1
CopiaCa_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6414	4,03E+04	100%	0.0	92.41%	301019445	NC_029977.1
CopiaCa_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6375	2,58E+04	100%	0.0	91.70%	219521584	NC_029982.1
CopiaCa_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6335	1,06E+04	100%	0.0	91.60%	229934170	NC_029988.1
CopiaCa_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5729	9,73E+03	100%	0.0	91.34%	222112641	NC_029983.1
CopiaCa_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	5578	6,17E+03	100%	0.0	88.75%	3493078	NW_015960517.1
CopiaCa_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5570	7,49E+03	93%	0.0	94.65%	153299543	NC_029984.1
CopiaCa_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5533	3,03E+04	100%	0.0	90.87%	238794889	NC_029985.1
CopiaCa_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3611	<i>Capsicum annuum</i>	5393	5,39E+03	99%	0.0	87.89%	177705	NW_015960430.1
CopiaCa_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00197	<i>Solanum tuberosum</i>	4806	4,81E+03	96%	0.0	86.38%	1164124	NW_006239125.1
CopiaCa_24	refeq-genome	8	Solanum pennellii chromosome 9, SPENN9V200	<i>Solanum pennellii</i>	4519	7,63E+03	92%	0.0	86.38%	84057508	NC_028645.1
CopiaCa_24	refeq-genome	8	Solanum pennellii chromosome 7, SPENN9V200	<i>Solanum pennellii</i>	4514	1,57E+04	90%	0.0	86.27%	79292169	NC_028643.1
CopiaCa_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00044	<i>tuberosum Solanum</i>	4510	4,51E+03	97%	0.0	85.09%	2434495	NW_006238970.1
CopiaCa_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00198	<i>tuberosum Capsicum</i>	4479	4,48E+03	98%	0.0	84.66%	1172864	NW_006239121.1
CopiaCa_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4197	<i>annuum Nicotiana</i>	4455	6,54E+03	100%	0.0	90.68%	182937	NW_015961354.1
CopiaCa_24	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold11322	<i>tomentosiformis Nicotiana</i>	4414	4,41E+03	92%	0.0	85.55%	12745	NW_008843181.1
CopiaCa_24	refeq-genome	8	Nicotiana attenuata strain UT chromosome 12, NIATTr2	<i>attenuata Solanum</i>	4392	4,39E+03	89%	0.0	86.12%	53491057	NC_032000.1
CopiaCa_24	refeq-genome	8	Solanum pennellii chromosome 4, SPENN9V200	<i>pennellii Nicotiana</i>	4377	1,62E+04	92%	0.0	85.64%	77197300	NC_028640.1
CopiaCa_24	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold127524	<i>tabacum Solanum</i>	4359	4,36E+03	92%	0.0	85.32%	25554	NW_015807743.1
CopiaCa_24	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	<i>lycopersicum Solanum</i>	4327	7,16E+03	90%	0.0	85.54%	72906345	NC_015446.3
CopiaCa_24	refeq-genome	8	Solanum pennellii chromosome 6, SPENN9V200	<i>pennellii Nicotiana</i>	4313	4,31E+03	95%	0.0	84.53%	60730942	NC_028642.1
CopiaCa_24	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	<i>attenuata Nicotiana</i>	4217	4,22E+03	92%	0.0	84.70%	883723	NW_017670363.1
CopiaCa_24	refeq-genome	8	Nicotiana attenuata strain UT chromosome 11, NIATTr2	<i>attenuata Solanum</i>	4196	4,20E+03	96%	0.0	83.95%	61250262	NC_031999.1
CopiaCa_24	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	<i>lycopersicum Solanum</i>	4169	1,10E+04	92%	0.0	84.92%	66557038	NC_015441.3
CopiaCa_24	refeq-genome	8	Solanum pennellii chromosome 11, SPENN9V200	<i>pennellii</i>	4048	1,74E+04	90%	0.0	84.49%	66223686	NC_028647.1
CopiaCa_24	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	<i>Solanum lycopersicum</i>	3938	3,94E+03	90%	0.0	83.90%	49794276	NC_015443.3
CopiaCa_24	refeq-genome	8	Solanum pennellii chromosome 1, SPENN9V200	<i>Solanum pennellii</i>	3934	1,63E+04	92%	0.0	83.82%	109333515	NC_028637.1
CopiaCa_24	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold105839	<i>Nicotiana sylvestris</i>	3917	4,38E+03	96%	0.0	85.64%	37463	NW_009345291.1
CopiaCa_24	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold50197	<i>Nicotiana tabacum</i>	3916	4,37E+03	93%	0.0	86.21%	39211	NW_01591296.1
CopiaCa_24	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	<i>Nicotiana attenuata</i>	3784	3,78E+03	86%	0.0	83.91%	586478	NW_017670551.1
CopiaCa_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3482	1,45E+04	100%	0.0	89.50%	205736368	NC_029986.1
CopiaCa_24	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	<i>Solanum lycopersicum</i>	3419	6,44E+03	90%	0.0	83.43%	98455869	NC_015438.3
CopiaCa_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3269	1,02E+04	100%	0.0	91.21%	217274494	NC_029981.1
CopiaCa_24	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	<i>Solanum lycopersicum</i>	3251	4,02E+03	86%	0.0	85.10%	65987440	NC_015445.3
CopiaCa_24	refeq-genome	8	Solanum pennellii chromosome 2, SPENN9V200	<i>Solanum pennellii</i>	3101	9,84E+03	96%	0.0	85.68%	59803892	NC_028638.1
CopiaCa_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00460	<i>Solanum tuberosum</i>	2977	4,57E+03	93%	0.0	86.81%	452291	NW_006239377.1
CopiaCa_24	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	<i>Solanum lycopersicum</i>	2959	4,67E+03	89%	0.0	84.80%	55977580	NC_015439.3
CopiaCa_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00216	<i>Solanum tuberosum</i>	2876	4,67E+03	92%	0.0	86.13%	1093816	NW_006239143.1
CopiaCa_24	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	<i>Solanum lycopersicum</i>	2824	3,54E+03	88%	0.0	81.99%	56597135	NC_015448.3
CopiaCa_24	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	<i>Solanum lycopersicum</i>	2736	8,25E+03	90%	0.0	83.95%	68175699	NC_015444.3

CopiaCa_24	refeq-genome	8	Solanum pennellii chromosome 3, SPENN200	<i>Solanum pennellii</i>	2678	1,14E+04	96%	0.0	81.82%	75414019	NC_028639.1
CopiaCa_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00066	<i>Solanum tuberosum</i>	2507	3,85E+03	88%	0.0	83.29%	2159599	NW_006238989.1
CopiaCa_24	refeq-genome	8	Solanum pennellii chromosome 10, SPENN200	<i>Solanum pennellii</i>	2423	1,12E+04	90%	0.0	83.67%	82529941	NC_028646.1
CopiaCa_24	refeq-genome	8	Solanum pennellii chromosome 8, SPENN200	<i>Solanum pennellii</i>	2320	5,89E+03	88%	0.0	85.57%	70546659	NC_028644.1
CopiaCa_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00404	<i>Solanum tuberosum</i>	2285	4,29E+03	82%	0.0	87.86%	496490	NW_006239353.1
CopiaCa_24	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	2222	3,58E+03	83%	0.0	85.44%	246988	NW_017671827.1
CopiaCa_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00182	<i>Solanum tuberosum</i>	2183	3,73E+03	85%	0.0	83.50%	1231519	NW_006239108.1
CopiaCa_24	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1027	<i>Nicotiana tabacum</i>	2098	3,83E+03	87%	0.0	82.48%	230627	NW_015789328.1
CopiaCa_24	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold23420	<i>Nicotiana tomentosiformis</i>	2087	3,73E+03	87%	0.0	82.40%	75929	NW_008902954.1
CopiaCa_24	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold25864	<i>Nicotiana sylvestris</i>	2008	3,36E+03	86%	0.0	80.86%	180178	NW_009510347.1
CopiaCa_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00031	<i>Solanum tuberosum</i>	1816	4,69E+03	85%	0.0	89.58%	2735125	NW_006238957.1
CopiaCa_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00022	<i>Solanum tuberosum</i>	1389	4,12E+03	81%	0.0	87.61%	3032639	NW_006238948.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8597	5,59E+04	100%	0.0	100.00%	153299543	NC_029984.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5794	3,59E+08	100%	0.0	89.32%	219521584	NC_029982.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5705	2,10E+08	100%	0.0	88.87%	220335243	NC_029987.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5642	3,58E+08	100%	0.0	88.59%	301019445	NC_029977.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5626	2,93E+08	100%	0.0	88.66%	229934170	NC_029988.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold11487	<i>Capsicum annuum</i>	5618	5,94E+03	100%	0.0	88.49%	98637	NW_015960853.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1874	<i>Capsicum annuum</i>	5618	6,37E+03	100%	0.0	88.65%	719064	NW_015961147.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold53	<i>Capsicum annuum</i>	5613	5,61E+03	99%	0.0	88.55%	1681775	NW_015961414.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5605	1,42E+08	100%	0.0	88.50%	238794889	NC_029985.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5579	1,83E+08	100%	0.0	88.50%	222112641	NC_029983.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1399	<i>Capsicum annuum</i>	5553	1,59E+04	100%	0.0	88.26%	1742971	NW_015961630.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3606	<i>Capsicum annuum</i>	5520	6,26E+03	99%	0.0	89.44%	166922	NW_015961502.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5404	1,77E+08	100%	0.0	87.82%	205736368	NC_029986.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5387	3,04E+08	100%	0.0	88.96%	261510930	NC_029979.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5301	1,58E+08	100%	0.0	87.46%	215701946	NC_029980.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1038	<i>Capsicum annuum</i>	5301	1,34E+04	98%	0.0	88.54%	2164954	NW_015960831.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4660	<i>Capsicum annuum</i>	5241	5,48E+03	99%	0.0	87.27%	152652	NW_015961314.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5153	1,71E+08	100%	0.0	88.06%	217274494	NC_029981.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5149	1,61E+08	100%	0.0	86.87%	163962470	NC_029978.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capsicum annuum</i>	5049	6,51E+03	94%	0.0	87.50%	2397553	NW_015961318.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1037	<i>Capsicum annuum</i>	5012	1,13E+04	99%	0.0	87.45%	2186250	NW_015960681.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1730	<i>Capsicum annuum</i>	4992	4,99E+03	99%	0.0	86.34%	1080651	NW_015961122.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2375	<i>Capsicum annuum</i>	4929	1,73E+04	95%	0.0	86.81%	1171512	NW_015961637.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3800	<i>Capsicum annuum</i>	4861	4,86E+03	94%	0.0	86.75%	452370	NW_015961629.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3354	<i>Capsicum annuum</i>	4737	4,79E+03	94%	0.0	86.30%	55019	NW_015960344.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold949	<i>Capsicum annuum</i>	4719	2,46E+04	100%	0.0	86.26%	3128562	NW_015961487.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold13303	<i>Capsicum annuum</i>	4479	4,48E+03	94%	0.0	85.17%	61598	NW_015961592.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1633	<i>Capsicum annuum</i>	4475	7,83E+03	95%	0.0	85.24%	1620172	NW_015961041.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1929	<i>Capsicum annuum</i>	4455	4,46E+03	94%	0.0	85.10%	996492	NW_015961795.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4129	<i>Capsicum annuum</i>	4453	1,79E+04	99%	0.0	87.15%	328737	NW_015961069.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold723	<i>Capsicum annuum</i>	4433	5,35E+03	92%	0.0	87.31%	559742	NW_015961191.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1245	<i>Capsicum annuum</i>	4311	5,37E+03	93%	0.0	86.26%	307737	NW_015960675.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1973	<i>Capsicum annuum</i>	4274	4,27E+03	94%	0.0	84.40%	1714326	NW_015961422.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold7165	<i>Capsicum annuum</i>	4263	8,30E+03	83%	0.0	86.59%	114031	NW_015960684.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	4220	4,22E+03	94%	0.0	84.21%	7059680	NW_015960855.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold450	<i>Capsicum annuum</i>	4207	5,61E+03	86%	0.0	88.08%	175795	NW_015961071.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1174	<i>Capsicum annuum</i>	4115	5,50E+03	99%	0.0	88.36%	656414	NW_015961459.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold16	<i>Capsicum annuum</i>	3914	1,99E+04	99%	0.0	84.60%	2023789	NW_015960591.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold519	<i>Capsicum annuum</i>	3897	1,99E+04	94%	0.0	85.38%	1101621	NW_015961578.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9	<i>Capsicum annuum</i>	3871	6,46E+03	90%	0.0	84.60%	6567951	NW_015961815.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold450	<i>Capsicum annuum</i>	3866	9,47E+03	99%	0.0	85.50%	928404	NW_015960451.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1339	<i>Capsicum annuum</i>	3795	4,80E+03	99%	0.0	85.64%	749978	NW_015961154.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold306	<i>Capsicum annuum</i>	3613	4,49E+03	92%	0.0	84.15%	1897442	NW_015960687.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1741	<i>Capsicum annuum</i>	3581	1,55E+04	99%	0.0	87.09%	912839	NW_015961757.1

CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold7487	<i>Capsicum annuum</i>	3345	5,14E+03	98%	0.0	85.45%	147954	NW_015960866.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold659	<i>Capsicum annuum</i>	3293	8,90E+03	99%	0.0	88.31%	613124	NW_015961271.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1124	<i>Capsicum annuum</i>	3168	1,06E+04	93%	0.0	84.57%	2110277	NW_015961370.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3653	<i>Capsicum annuum</i>	2955	5,66E+03	94%	0.0	86.89%	172086	NW_015961357.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3822	<i>Capsicum annuum</i>	2785	4,67E+03	87%	0.0	87.29%	410103	NW_015960760.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold814	<i>Capsicum annuum</i>	2484	4,63E+03	88%	0.0	88.08%	1128876	NW_015960740.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold7595	<i>Capsicum annuum</i>	2451	5,11E+03	93%	0.0	88.46%	38816	NW_015960439.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold181	<i>Capsicum annuum</i>	2303	5,22E+03	92%	0.0	87.62%	2703691	NW_015961388.1
CopiaCa_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2447	<i>Capsicum annuum</i>	2008	4,53E+03	89%	0.0	86.47%	421792	NW_015960841.1
CopiaCa_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7598	9,59E+08	100%	0.0	100.00%	217274494	NC_029981.1
CopiaCa_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5251	7,64E+08	100%	0.0	89.76%	205736368	NC_029986.1
CopiaCa_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5168	6,33E+08	100%	0.0	89.34%	301019445	NC_029977.1
CopiaCa_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5092	5,22E+08	100%	0.0	89.15%	153299543	NC_029984.1
CopiaCa_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5077	1,25E+09	100%	0.0	88.96%	219521584	NC_029982.1
CopiaCa_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4793	3,81E+08	100%	0.0	87.83%	222112641	NC_029983.1
CopiaCa_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4715	8,78E+08	100%	0.0	87.49%	215701946	NC_029980.1
CopiaCa_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4569	1,23E+09	100%	0.0	89.04%	261510930	NC_029979.1
CopiaCa_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4100	5,13E+08	100%	0.0	86.42%	220335243	NC_029987.1
CopiaCa_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8588	1,89E+04	100%	0.0	100.00%	220335243	NC_029987.1
CopiaCa_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5295	2,37E+04	100%	0.0	89.99%	215701946	NC_029980.1
CopiaCa_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4953	1,76E+04	100%	0.0	88.82%	261510930	NC_029979.1
CopiaCa_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4638	1,37E+04	100%	0.0	87.41%	219521584	NC_029982.1
CopiaCa_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1514	<i>Capsicum annuum</i>	4590	5,88E+03	98%	0.0	87.14%	427058	NW_015960414.1
CopiaCa_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1174	<i>Capsicum annuum</i>	4578	4,94E+03	95%	0.0	87.14%	656414	NW_015961459.1
CopiaCa_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4571	8,42E+03	95%	0.0	88.34%	301019445	NC_029977.1
CopiaCa_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3729	5,11E+03	84%	0.0	84.09%	238794889	NC_029985.1
CopiaCa_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3397	2,18E+04	100%	0.0	84.58%	222112641	NC_029983.1
CopiaCa_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2970	5,50E+03	80%	0.0	83.97%	229934170	NC_029988.1
CopiaCa_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2150	1,55E+04	98%	0.0	87.60%	205736368	NC_029986.1
CopiaCa_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2043	1,29E+04	98%	0.0	85.36%	217274494	NC_029981.1
CopiaCa_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1124	<i>Capsicum annuum</i>	1459	3,63E+03	82%	0.0	84.30%	2110277	NW_015961370.1
CopiaCa_28	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	<i>Nicotiana attenuata</i>	3362	3,36E+03	81%	0.0	82.80%	467702	NW_017670748.1
CopiaCa_28	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold20124	<i>Nicotiana sylvestris</i>	3345	3,35E+03	81%	0.0	82.71%	137542	NW_009451293.1
CopiaCa_28	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold10240	<i>Nicotiana sylvestris</i>	1652	3,16E+03	81%	0.0	82.11%	329378	NW_009341471.1
CopiaCa_28	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold81250	<i>Nicotiana tabacum</i>	3339	3,34E+03	81%	0.0	82.69%	82904	NW_015940238.1
CopiaCa_28	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold92772	<i>Nicotiana tabacum</i>	1668	3,17E+03	81%	0.0	82.29%	131555	NW_015949767.1
CopiaCa_28	refeq-genome	8	Solanum pennellii chromosome 1, SPENNIV200	<i>Solanum pennellii</i>	2809	5,11E+03	81%	0.0	80.28%	109333515	NC_028637.1
CopiaCa_28	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf01081	<i>Solanum tuberosum</i>	4678	4,90E+03	87%	0.0	88.89%	24442	NW_006240019.1
CopiaCa_28	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00009	<i>Solanum tuberosum</i>	3273	3,27E+03	81%	0.0	82.43%	3955149	NW_006238935.1
CopiaCa_28	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00564	<i>Solanum tuberosum</i>	2939	1,05E+04	81%	0.0	80.85%	285676	NW_006239498.1
CopiaCa_28	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00052	<i>Solanum tuberosum</i>	2933	3,35E+03	81%	0.0	82.83%	2278663	NW_006238977.1
CopiaCa_28	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00063	<i>Solanum tuberosum</i>	2867	3,27E+03	80%	0.0	82.53%	2166112	NW_006238988.1
CopiaCa_28	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00121	<i>Solanum tuberosum</i>	2608	2,61E+03	81%	0.0	79.43%	1607711	NW_006239046.1
CopiaCa_28	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00223	<i>Solanum tuberosum</i>	1877	2,82E+03	81%	0.0	81.81%	1084529	NW_006239146.1
CopiaCa_28	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00109	<i>Solanum tuberosum</i>	1757	4,01E+03	87%	0.0	85.99%	1626695	NW_006239042.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11060	1,56E+08	100%	0.0	100.00%	215701946	NC_029980.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6063	1,92E+08	99%	0.0	85.71%	229934170	NC_029988.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6006	7,27E+04	98%	0.0	88.92%	153299543	NC_029984.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6004	1,91E+08	99%	0.0	87.78%	301019445	NC_029977.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5999	2,07E+08	98%	0.0	88.69%	205736368	NC_029986.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5991	2,59E+08	98%	0.0	85.27%	163962470	NC_029978.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5984	1,75E+08	97%	0.0	88.49%	261510930	NC_029979.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5947	1,71E+08	99%	0.0	87.86%	217274494	NC_029981.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5849	1,87E+08	96%	0.0	87.75%	219521584	NC_029982.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5803	1,45E+08	99%	0.0	84.63%	238794889	NC_029985.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5766	1,34E+08	99%	0.0	87.76%	220335243	NC_029987.1

CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5648	1,45E+08	96%	0.0	87.82%	222112641	NC_029983.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9053	<i>Capsicum annuum</i>	5548	5,99E+03	84%	0.0	86.57%	132257	NW_015960629.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold814	<i>Capsicum annuum</i>	5548	6,29E+03	94%	0.0	86.61%	1128876	NW_015960740.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold16	<i>Capsicum annuum</i>	5518	5,52E+03	86%	0.0	85.90%	2023789	NW_015960959.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2265	<i>Capsicum annuum</i>	5467	7,87E+03	95%	0.0	89.04%	1257389	NW_015961797.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold619	<i>Capsicum annuum</i>	5347	5,47E+03	84%	0.0	86.86%	2512642	NW_015960498.1
CopiaCa_30	refeq-genome	8	Solanum pennellii chromosome 6, SPENN200	<i>Solanum pennellii</i>	5289	7,69E+04	90%	0.0	86.33%	60730942	NC_028642.1
CopiaCa_30	refeq-genome	8	Solanum pennellii chromosome 2, SPENN200	<i>Solanum pennellii</i>	5280	9,44E+04	89%	0.0	86.01%	59803892	NC_028638.1
CopiaCa_30	refeq-genome	8	Solanum pennellii chromosome 4, SPENN200	<i>Solanum pennellii</i>	5278	1,02E+08	89%	0.0	86.32%	77197300	NC_028640.1
CopiaCa_30	refeq-genome	8	Solanum pennellii chromosome 5, SPENN200	<i>Solanum pennellii</i>	5278	1,09E+08	86%	0.0	86.32%	77991103	NC_028641.1
CopiaCa_30	refeq-genome	8	Solanum pennellii chromosome 8, SPENN200	<i>Solanum pennellii</i>	5278	9,23E+04	88%	0.0	86.28%	70546659	NC_028644.1
CopiaCa_30	refeq-genome	8	Solanum pennellii chromosome 10, SPENN200	<i>Solanum pennellii</i>	5273	1,07E+08	87%	0.0	86.30%	82529941	NC_028646.1
CopiaCa_30	refeq-genome	8	Solanum pennellii chromosome 3, SPENN200	<i>Solanum pennellii</i>	5267	5,48E+04	88%	0.0	86.24%	75414019	NC_028639.1
CopiaCa_30	refeq-genome	8	Solanum pennellii chromosome 9, SPENN200	<i>Solanum pennellii</i>	5267	1,22E+08	88%	0.0	86.25%	84057508	NC_028645.1
CopiaCa_30	refeq-genome	8	Solanum pennellii chromosome 1, SPENN200	<i>Solanum pennellii</i>	5262	1,87E+08	89%	0.0	86.26%	109333515	NC_028637.1
CopiaCa_30	refeq-genome	8	Solanum pennellii chromosome 7, SPENN200	<i>Solanum pennellii</i>	5262	1,08E+08	88%	0.0	86.41%	79291269	NC_028643.1
CopiaCa_30	refeq-genome	8	Solanum pennellii chromosome 11, SPENN200	<i>Solanum pennellii</i>	5254	1,01E+08	88%	0.0	86.33%	66223686	NC_028647.1
CopiaCa_30	refeq-genome	8	Solanum pennellii chromosome 12, SPENN200	<i>Solanum pennellii</i>	5243	1,65E+08	89%	0.0	86.17%	83305730	NC_028648.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold680	<i>Capsicum annuum</i>	5219	5,58E+03	99%	0.0	84.74%	788567	NW_015961649.1
CopiaCa_30	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	<i>Solanum lycopersicum</i>	5110	8,96E+04	87%	0.0	85.93%	56597135	NC_015448.3
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9	<i>Capsicum annuum</i>	5088	1,09E+04	84%	0.0	85.52%	6567951	NW_015961815.1
CopiaCa_30	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	<i>Solanum lycopersicum</i>	5044	9,85E+04	89%	0.0	85.61%	65633393	NC_015447.3
CopiaCa_30	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	<i>Solanum lycopersicum</i>	5020	1,30E+08	89%	0.0	85.40%	66723567	NC_015442.3
CopiaCa_30	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	<i>Solanum lycopersicum</i>	5020	8,95E+04	89%	0.0	85.54%	72906345	NC_015446.3
CopiaCa_30	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	<i>Solanum lycopersicum</i>	4975	1,95E+08	89%	0.0	85.39%	98455869	NC_015438.3
CopiaCa_30	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	<i>Solanum lycopersicum</i>	4964	8,95E+04	89%	0.0	84.89%	68175699	NC_015444.3
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2003	<i>Capsicum annuum</i>	4924	6,93E+03	95%	0.0	86.68%	314382	NW_015961213.1
CopiaCa_30	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	<i>Solanum lycopersicum</i>	4922	8,31E+04	88%	0.0	85.05%	55977580	NC_015439.3
CopiaCa_30	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	<i>Solanum lycopersicum</i>	4883	1,14E+08	88%	0.0	85.00%	68126176	NC_015449.3
CopiaCa_30	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	<i>Solanum lycopersicum</i>	4879	4,52E+04	88%	0.0	84.63%	65987440	NC_015445.3
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00142	<i>Solanum tuberosum</i>	4817	5,28E+03	87%	0.0	84.60%	1452299	NW_006239067.1
CopiaCa_30	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	<i>Solanum lycopersicum</i>	4798	6,40E+04	87%	0.0	84.59%	72290146	NC_015440.3
CopiaCa_30	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	<i>Solanum lycopersicum</i>	4767	6,70E+04	89%	0.0	84.64%	66557038	NC_015441.3
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	4732	5,77E+03	92%	0.0	84.09%	3311724	NW_015960635.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00743	<i>Solanum tuberosum</i>	4662	4,90E+03	87%	0.0	83.98%	154618	NW_006239654.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold8606	<i>Capsicum annuum</i>	4630	4,79E+03	81%	0.0	84.56%	138485	NW_015960788.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold490	<i>Capsicum annuum</i>	4626	2,94E+04	93%	0.0	86.60%	695234	NW_015960447.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00315	<i>Solanum tuberosum</i>	4514	5,77E+03	88%	0.0	86.12%	757876	NW_006239239.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00187	<i>Solanum tuberosum</i>	4421	4,75E+03	84%	0.0	84.57%	1214092	NW_006239111.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00452	<i>Solanum tuberosum</i>	4407	4,85E+03	88%	0.0	82.83%	457427	NW_006239375.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00218	<i>Solanum tuberosum</i>	4357	9,64E+03	90%	0.0	83.22%	1093294	NW_006239144.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold231	<i>Capsicum annuum</i>	4289	8,65E+03	92%	0.0	90.19%	1482909	NC_015961568.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold979	<i>Capsicum annuum</i>	4268	6,27E+03	90%	0.0	86.85%	1172749	NW_015961157.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00038	<i>Solanum tuberosum</i>	4248	5,08E+03	81%	0.0	86.47%	2533956	NW_006238966.1
CopiaCa_30	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	<i>Solanum lycopersicum</i>	4237	4,57E+04	87%	0.0	85.17%	49794276	NC_015443.3
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00740	<i>Solanum tuberosum</i>	4213	1,68E+04	87%	0.0	82.40%	151380	NW_006239662.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00258	<i>Solanum tuberosum</i>	4211	4,21E+03	81%	0.0	82.45%	953633	NW_006239179.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00177	<i>Solanum tuberosum</i>	4120	5,05E+03	80%	0.0	85.92%	1275411	NW_006239101.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00001	<i>Solanum tuberosum</i>	4019	1,43E+04	82%	0.0	85.52%	7090834	NW_006238927.1
CopiaCa_30	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl_Nsyl_scaffold44737	<i>Nicotiana sylvestris</i>	4010	4,01E+03	80%	0.0	81.99%	48392	NW_009531317.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00126	<i>Solanum tuberosum</i>	4006	1,02E+04	85%	0.0	85.46%	1558646	NW_006239051.1
CopiaCa_30	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90_Ntab-TN90_scaffold90716	<i>Nicotiana tabacum</i>	4002	4,00E+03	80%	0.0	81.96%	31948	NW_015948067.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00008	<i>Solanum tuberosum</i>	3999	9,44E+03	88%	0.0	85.32%	4063183	NW_006238934.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00055	<i>Solanum tuberosum</i>	3997	5,69E+03	85%	0.0	85.39%	2254777	NW_006238981.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00002	<i>Solanum tuberosum</i>	3995	8,54E+03	86%	0.0	85.18%	6551949	NW_006238928.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capsicum annuum</i>	3936	6,56E+03	95%	0.0	89.18%	2397553	NW_015961318.1

CopiaCa_30	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold96870	<i>Nicotiana sylvestris</i>	3838	3,84E+03	80%	0.0	81.35%	79525	NW_009589242.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00346	<i>Solanum tuberosum</i>	3829	4,20E+03	80%	0.0	82.25%	666711	NW_006239265.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00387	<i>Solanum tuberosum</i>	3794	4,34E+03	81%	0.0	81.55%	560296	NW_006239312.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold742	<i>Capsicum annuum</i>	3557	5,59E+03	80%	0.0	86.01%	2382255	NW_015961471.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold53	<i>Capsicum annuum</i>	3520	6,56E+03	90%	0.0	90.38%	1681775	NW_015961414.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00199	<i>Solanum tuberosum</i>	3288	9,01E+03	86%	0.0	85.34%	1110059	NW_006239136.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00037	<i>Solanum tuberosum</i>	3227	7,63E+03	82%	0.0	80.29%	2613480	NW_006238961.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3800	<i>Capsicum annuum</i>	3173	6,71E+03	92%	0.0	86.83%	452370	NW_015961629.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2635	<i>Capsicum annuum</i>	3120	5,70E+03	86%	0.0	87.42%	699991	NW_015960547.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00133	<i>Solanum tuberosum</i>	3027	1,66E+04	88%	0.0	86.29%	1531558	NW_006239056.1
CopiaCa_30	refeq-genome	8	Nicotiana attenuata strain UT chromosome 1, NIATT2	<i>Nicotiana attenuata</i>	2955	6,11E+03	83%	0.0	82.48%	110538649	NC_031989.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold578	<i>Capsicum annuum</i>	2926	5,32E+03	91%	0.0	86.62%	1213777	NW_015961091.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00244	<i>Solanum tuberosum</i>	2918	5,25E+03	87%	0.0	83.55%	945173	NW_006239181.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00006	<i>Solanum tuberosum</i>	2904	9,63E+03	86%	0.0	81.99%	4150974	NW_006238932.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00031	<i>Solanum tuberosum</i>	2789	1,06E+04	85%	0.0	81.20%	2735125	NW_006238957.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00097	<i>Solanum tuberosum</i>	2695	7,63E+03	82%	0.0	83.87%	1770913	NW_006239022.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00287	<i>Solanum tuberosum</i>	2643	4,86E+03	84%	0.0	83.05%	844541	NW_006239211.1
CopiaCa_30	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC00000001	<i>Solanum lycopersicum</i>	2638	7,47E+03	81%	0.0	83.98%	2014460	NW_020442482.1
CopiaCa_30	refeq-genome	8	Nicotiana attenuata strain UT chromosome 8, NIATT2	<i>Nicotiana attenuata</i>	2615	1,27E+04	81%	0.0	81.92%	47755514	NC_031996.1
CopiaCa_30	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	2586	4,19E+03	80%	0.0	82.18%	87492	NW_017674709.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00168	<i>Solanum tuberosum</i>	2547	8,63E+03	80%	0.0	84.87%	1293034	NW_006239096.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3079	<i>Capsicum annuum</i>	2495	6,15E+03	86%	0.0	92.07%	299956	NW_015961555.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00924	<i>Solanum tuberosum</i>	2436	6,06E+03	81%	0.0	84.48%	54308	NW_006239841.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00130	<i>Solanum tuberosum</i>	2407	5,17E+03	86%	0.0	85.26%	1515740	NW_006239060.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00454	<i>Solanum tuberosum</i>	2272	5,62E+03	87%	0.0	85.89%	461660	NW_006239370.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00018	<i>Solanum tuberosum</i>	2187	1,29E+04	88%	0.0	86.67%	3280163	NW_006238944.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00061	<i>Solanum tuberosum</i>	2180	4,34E+03	84%	0.0	83.72%	2128288	NW_006238993.1
CopiaCa_30	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	2161	3,95E+03	80%	0.0	82.95%	207768	NW_017672228.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00411	<i>Solanum tuberosum</i>	1986	6,99E+03	82%	0.0	85.79%	508257	NW_006239344.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00045	<i>Solanum tuberosum</i>	1954	6,23E+03	80%	0.0	84.49%	2407939	NW_006238972.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00269	<i>Solanum tuberosum</i>	1823	6,93E+03	86%	0.0	84.79%	905297	NW_006239191.1
CopiaCa_30	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3682	<i>Capsicum annuum</i>	1718	5,56E+03	85%	0.0	83.31%	568842	NW_015961135.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00320	<i>Solanum tuberosum</i>	1552	4,32E+03	80%	0.0	84.27%	738063	NW_006239242.1
CopiaCa_30	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00003	<i>Solanum tuberosum</i>	1450	5,29E+03	83%	0.0	86.55%	4590222	NW_006238929.1
CopiaCa_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8288	2,40E+04	100%	0.0	100.00%	153299543	NC_029984.1
CopiaCa_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6763	1,76E+04	100%	0.0	93.86%	301019445	NC_029977.1
CopiaCa_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6678	2,78E+04	100%	0.0	93.53%	163962470	NC_029978.1
CopiaCa_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6615	2,79E+04	100%	0.0	93.36%	229934170	NC_029988.1
CopiaCa_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6034	1,60E+04	100%	0.0	91.00%	261510930	NC_029979.1
CopiaCa_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5751	4,30E+04	100%	0.0	90.59%	219521584	NC_029982.1
CopiaCa_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5666	1,21E+04	98%	0.0	93.16%	205736368	NC_029986.1
CopiaCa_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5024	2,43E+04	100%	0.0	90.72%	238794889	NC_029985.1
CopiaCa_33	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00071	<i>Solanum tuberosum</i>	4545	4,55E+03	99%	0.0	85.06%	2054262	NW_006238997.1
CopiaCa_33	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00397	<i>Solanum tuberosum</i>	4098	4,10E+03	91%	0.0	84.78%	523264	NW_006239334.1
CopiaCa_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4041	7,82E+03	88%	0.0	85.05%	220335243	NC_029987.1
CopiaCa_33	refeq-genome	8	Solanum pennellii chromosome 2, SPENN200	<i>Solanum pennellii</i>	3886	4,17E+03	88%	0.0	84.57%	59803892	NC_028638.1
CopiaCa_33	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00081	<i>Solanum tuberosum</i>	3856	3,86E+03	93%	0.0	83.53%	1932198	NW_006239008.1
CopiaCa_33	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00079	<i>Solanum tuberosum</i>	3779	4,52E+03	96%	0.0	85.61%	1965697	NW_006239004.1
CopiaCa_33	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00013	<i>Solanum tuberosum</i>	3698	4,19E+03	90%	0.0	84.84%	3422333	NW_006238940.1
CopiaCa_33	refeq-genome	8	Solanum pennellii chromosome 11, SPENN200	<i>Solanum pennellii</i>	3452	4,44E+03	80%	0.0	86.10%	66223686	NC_028647.1
CopiaCa_33	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	<i>Solanum lycopersicum</i>	3448	4,19E+03	88%	0.0	84.14%	55977580	NC_015439.3
CopiaCa_33	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00163	<i>Solanum tuberosum</i>	3393	3,39E+03	85%	0.0	82.76%	1340955	NW_006239087.1
CopiaCa_33	refeq-genome	8	Solanum pennellii chromosome 10, SPENN200	<i>Solanum pennellii</i>	3160	5,04E+03	92%	0.0	80.92%	82529941	NC_028646.1
CopiaCa_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2752	1,73E+04	100%	0.0	81.77%	217274494	NC_029981.1
CopiaCa_33	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00211	<i>Solanum tuberosum</i>	2717	4,20E+03	96%	0.0	85.46%	1109151	NW_006239137.1
CopiaCa_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold660	<i>Capsicum annuum</i>	2364	3,76E+03	86%	0.0	82.24%	663463	NW_015961299.1

CopiaCa_33	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00178	<i>Solanum tuberosum</i>	2322	4,57E+03	90%	0.0	85.17%	1266720	NW_006239102.1
CopiaCa_33	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf0025	<i>Solanum tuberosum</i>	1683	4,18E+03	89%	0.0	88.33%	2980670	NW_006238951.1
CopiaCa_33	refeq-genome	8	Solanum pennellii chromosome 1, SPENN200	<i>Solanum pennellii</i>	1659	1,89E+04	90%	0.0	81.46%	109333515	NC_028637.1
CopiaCa_33	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf0028	<i>Solanum tuberosum</i>	1489	2,67E+03	80%	0.0	78.42%	2805095	NW_006238954.1
CopiaCa_35	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9311	2,69E+04	100%	0.0	100.00%	205736368	NC_029986.1
CopiaCa_35	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4780	3,11E+04	100%	0.0	92.95%	219521584	NC_029982.1
CopiaCa_35	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4488	8,97E+03	99%	0.0	91.40%	217274494	NC_029981.1
CopiaCa_35	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4102	1,44E+04	100%	0.0	91.82%	301019445	NC_029977.1
CopiaCa_35	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4039	3,09E+04	100%	0.0	89.41%	261510930	NC_029979.1
CopiaCa_35	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3421	1,23E+04	93%	0.0	91.60%	238794889	NC_029985.1
CopiaCa_35	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3256	1,67E+04	100%	0.0	88.10%	222112641	NC_029983.1
CopiaCa_35	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2970	8,07E+03	95%	0.0	91.10%	229934170	NC_029988.1
CopiaCa_35	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1058	<i>Capsicum annuum</i>	2795	5,43E+03	86%	0.0	88.61%	165946	NW_015961110.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold19599	<i>Nicotiana sylvestris</i>	2734	4,38E+03	97%	0.0	82.09%	81334	NW_009445457.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold43824	<i>Nicotiana tabacum</i>	2728	6,17E+03	96%	0.0	82.35%	79168	NW_015907074.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold11737	<i>Nicotiana sylvestris</i>	2726	4,24E+03	92%	0.0	82.05%	151702	NW_009358103.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold28896	<i>Nicotiana sylvestris</i>	2726	4,31E+03	98%	0.0	82.07%	61887	NW_009513715.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold53213	<i>Nicotiana tabacum</i>	2723	4,23E+03	93%	0.0	82.00%	91520	NW_015915692.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold40233	<i>Nicotiana sylvestris</i>	2723	4,35E+03	98%	0.0	82.03%	58827	NW_009526314.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold99637	<i>Nicotiana tabacum</i>	2721	5,70E+03	98%	0.0	82.01%	43769	NW_015955177.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold53637	<i>Nicotiana sylvestris</i>	2721	4,34E+03	98%	0.0	82.00%	84359	NW_009541206.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5769	<i>Nicotiana tabacum</i>	2717	4,36E+03	98%	0.0	81.99%	79149	NW_015919754.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold22919	<i>Nicotiana sylvestris</i>	2717	5,35E+03	97%	0.0	81.99%	169066	NW_009482347.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold21003	<i>Nicotiana sylvestris</i>	2715	5,21E+03	98%	0.0	81.98%	123044	NW_009461060.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold18728	<i>Nicotiana tabacum</i>	2713	3,75E+03	84%	0.0	81.89%	107316	NW_015844842.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold63522	<i>Nicotiana tabacum</i>	2712	4,30E+03	97%	0.0	81.96%	57186	NW_015924936.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold75014	<i>Nicotiana tabacum</i>	2712	4,34E+03	98%	0.0	81.95%	46485	NW_015934912.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold151	<i>Nicotiana sylvestris</i>	2712	4,34E+03	98%	0.0	81.96%	58005	NW_009395469.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold16273	<i>Nicotiana sylvestris</i>	2712	4,30E+03	97%	0.0	81.96%	73974	NW_009048053.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold3203	<i>Nicotiana sylvestris</i>	2712	4,34E+03	98%	0.0	81.96%	207024	NW_009517199.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold42321	<i>Nicotiana sylvestris</i>	2712	6,99E+03	98%	0.0	81.96%	87778	NW_009528634.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold6057	<i>Nicotiana sylvestris</i>	2712	4,34E+03	98%	0.0	81.96%	77567	NW_009548909.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold73883	<i>Nicotiana sylvestris</i>	2712	4,35E+03	97%	0.0	81.96%	48559	NW_009563701.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold14077	<i>Nicotiana sylvestris</i>	2710	4,34E+03	98%	0.0	81.95%	37651	NW_009384103.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold35181	<i>Nicotiana sylvestris</i>	2710	4,35E+03	98%	0.0	81.96%	223463	NW_009520700.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold43454	<i>Nicotiana sylvestris</i>	2710	6,37E+03	98%	0.0	81.96%	60659	NW_009529892.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold145244	<i>Nicotiana tabacum</i>	2706	4,34E+03	98%	0.0	81.94%	39048	NW_015819852.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17072	<i>Nicotiana tabacum</i>	2706	4,34E+03	98%	0.0	81.93%	46069	NW_015835586.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold64357	<i>Nicotiana tabacum</i>	2704	4,11E+03	90%	0.0	81.91%	52930	NW_015925684.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold71353	<i>Nicotiana tabacum</i>	2704	4,34E+03	98%	0.0	81.93%	249447	NW_015931757.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold78083	<i>Nicotiana tabacum</i>	2704	7,85E+03	98%	0.0	81.93%	84057	NW_015937537.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold4106	<i>Nicotiana sylvestris</i>	2704	4,34E+03	98%	0.0	81.91%	300521	NW_009527232.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold43344	<i>Nicotiana sylvestris</i>	2704	4,33E+03	99%	0.0	81.93%	69731	NW_009529770.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold15997	<i>Nicotiana tabacum</i>	2702	4,19E+03	94%	0.0	81.90%	28426	NW_015829182.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold75228	<i>Nicotiana tabacum</i>	2702	4,26E+03	97%	0.0	82.03%	119599	NW_015935085.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7779	<i>Nicotiana tabacum</i>	2700	4,34E+03	98%	0.0	81.90%	122697	NW_015937379.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8323	<i>Nicotiana tabacum</i>	2700	4,34E+03	97%	0.0	82.08%	63792	NW_015941879.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold10493	<i>Nicotiana sylvestris</i>	2700	4,24E+03	94%	0.0	81.90%	407140	NW_009344281.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1297	<i>Nicotiana sylvestris</i>	2700	4,33E+03	98%	0.0	81.90%	163881	NW_009371802.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold276	<i>Nicotiana sylvestris</i>	2700	4,33E+03	98%	0.0	81.90%	204994	NW_009512275.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold35527	<i>Nicotiana sylvestris</i>	2700	4,33E+03	98%	0.0	81.90%	106024	NW_009521084.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold37089	<i>Nicotiana sylvestris</i>	2700	8,65E+03	98%	0.0	81.90%	61481	NW_009522819.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold37970	<i>Nicotiana sylvestris</i>	2700	4,34E+03	98%	0.0	81.91%	109875	NW_009523798.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold4890	<i>Nicotiana sylvestris</i>	2700	3,61E+03	83%	0.0	81.90%	10393	NW_009535942.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold28353	<i>Nicotiana tabacum</i>	2699	8,67E+03	98%	0.0	81.89%	89541	NW_015886501.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold11757	<i>Nicotiana sylvestris</i>	2699	4,33E+03	98%	0.0	81.90%	536095	NW_009358325.1

CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold82825	<i>Nicotiana tabacum</i>	2697	4,33E+03	97%	0.0	82.01%	84468	NW_015941556.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold56335	<i>Nicotiana sylvestris</i>	2697	4,33E+03	97%	0.0	82.00%	51274	NW_009544204.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17462	<i>Nicotiana tabacum</i>	2695	4,32E+03	96%	0.0	82.16%	105279	NW_015837790.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold41786	<i>Nicotiana tabacum</i>	2695	4,33E+03	96%	0.0	82.20%	156880	NW_015905192.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold6187	<i>Nicotiana tabacum</i>	2695	1,16E+04	98%	0.0	81.87%	121797	NW_015923456.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold72763	<i>Nicotiana tabacum</i>	2695	4,32E+03	98%	0.0	81.87%	94243	NW_015932988.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold11163	<i>Nicotiana sylvestris</i>	2695	8,65E+03	98%	0.0	81.87%	70535	NW_009351726.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold15615	<i>Nicotiana sylvestris</i>	2695	8,66E+03	97%	0.0	81.86%	58351	NW_009401192.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold37263	<i>Nicotiana sylvestris</i>	2695	9,92E+03	98%	0.0	81.87%	91883	NW_009523013.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold40245	<i>Nicotiana sylvestris</i>	2695	4,33E+03	99%	0.0	81.87%	46148	NW_009526327.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold6906	<i>Nicotiana sylvestris</i>	2695	7,83E+03	98%	0.0	81.87%	200015	NW_009558342.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold69336	<i>Nicotiana sylvestris</i>	2695	4,70E+03	99%	0.0	81.87%	37475	NW_009558649.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold7973	<i>Nicotiana sylvestris</i>	2695	4,33E+03	97%	0.0	81.88%	171798	NW_009570197.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold107274	<i>Nicotiana tabacum</i>	2693	4,33E+03	97%	0.0	82.02%	17883	NW_015792848.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold126364	<i>Nicotiana tabacum</i>	2693	4,33E+03	98%	0.0	81.87%	59964	NW_015806933.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold18120	<i>Nicotiana tabacum</i>	2693	4,33E+03	97%	0.0	81.86%	74335	NW_015841478.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24420	<i>Nicotiana tabacum</i>	2693	4,34E+03	97%	0.0	81.87%	37691	NW_015872804.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold15099	<i>Nicotiana sylvestris</i>	2691	4,31E+03	97%	0.0	81.97%	144529	NW_009395458.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold38148	<i>Nicotiana sylvestris</i>	2691	4,30E+03	96%	0.0	82.16%	133774	NW_009523996.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold40110	<i>Nicotiana sylvestris</i>	2691	4,29E+03	97%	0.0	81.88%	27008	NW_009526178.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold131047	<i>Nicotiana tabacum</i>	2689	4,31E+03	97%	0.0	81.84%	19879	NW_015810225.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold136175	<i>Nicotiana tabacum</i>	2689	4,32E+03	97%	0.0	81.84%	44171	NW_015813753.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold14535	<i>Nicotiana tabacum</i>	2689	4,32E+03	97%	0.0	81.86%	43771	NW_015819923.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold27552	<i>Nicotiana tabacum</i>	2689	4,33E+03	98%	0.0	81.84%	82479	NW_015885731.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold32847	<i>Nicotiana tabacum</i>	2689	4,32E+03	98%	0.0	81.84%	100428	NW_015890727.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold43111	<i>Nicotiana tabacum</i>	2689	4,34E+03	98%	0.0	81.85%	106924	NW_015906410.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold59918	<i>Nicotiana tabacum</i>	2689	4,33E+03	97%	0.0	81.99%	72977	NW_015921710.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold64433	<i>Nicotiana tabacum</i>	2689	4,33E+03	98%	0.0	81.84%	166873	NW_015925749.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold6485	<i>Nicotiana tabacum</i>	2689	4,33E+03	98%	0.0	81.82%	65313	NW_015926110.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8982	<i>Nicotiana tabacum</i>	2689	4,32E+03	97%	0.0	81.84%	199813	NW_015947320.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold26465	<i>Nicotiana sylvestris</i>	2689	8,67E+03	97%	0.0	81.84%	206821	NW_009511015.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold37329	<i>Nicotiana sylvestris</i>	2689	4,32E+03	98%	0.0	81.84%	49510	NW_009523086.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold3878	<i>Nicotiana sylvestris</i>	2689	4,41E+03	97%	0.0	81.84%	148721	NW_009524697.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold8580	<i>Nicotiana sylvestris</i>	2689	4,33E+03	99%	0.0	81.84%	113779	NW_009576942.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold9351	<i>Nicotiana sylvestris</i>	2689	4,32E+03	97%	0.0	81.84%	268250	NW_009585509.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10332	<i>Nicotiana tabacum</i>	2687	4,33E+03	97%	0.0	81.98%	209402	NW_015789816.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold16640	<i>Nicotiana tabacum</i>	2687	4,30E+03	97%	0.0	81.99%	105449	NW_015833017.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold10932	<i>Nicotiana sylvestris</i>	2687	4,33E+03	97%	0.0	81.85%	75826	NW_009349159.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold7532	<i>Nicotiana sylvestris</i>	2687	4,31E+03	98%	0.0	81.83%	107574	NW_009565298.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold16242	<i>Nicotiana tabacum</i>	2686	4,32E+03	96%	0.0	82.19%	129443	NW_015830679.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4590	<i>Nicotiana tabacum</i>	2686	4,32E+03	97%	0.0	81.94%	37824	NW_015908982.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold13069	<i>Nicotiana sylvestris</i>	2686	4,32E+03	97%	0.0	81.94%	168331	NW_009372903.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1970	<i>Nicotiana sylvestris</i>	2686	4,32E+03	97%	0.0	81.94%	173175	NW_009446580.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold29616	<i>Nicotiana sylvestris</i>	2686	4,32E+03	97%	0.0	81.94%	33055	NW_009514516.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold53512	<i>Nicotiana sylvestris</i>	2686	4,31E+03	98%	0.0	81.82%	31353	NW_009541068.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold101329	<i>Nicotiana tabacum</i>	2684	4,30E+03	97%	0.0	81.81%	74040	NW_015788280.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold156423	<i>Nicotiana tabacum</i>	2684	4,32E+03	97%	0.0	81.81%	34134	NW_015826970.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold25321	<i>Nicotiana tabacum</i>	2684	4,32E+03	97%	0.0	81.81%	132464	NW_015876807.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold29060	<i>Nicotiana tabacum</i>	2684	4,30E+03	98%	0.0	81.81%	52040	NW_015887154.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold33066	<i>Nicotiana tabacum</i>	2684	4,31E+03	97%	0.0	81.94%	87385	NW_015890936.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4685	<i>Nicotiana tabacum</i>	2684	4,31E+03	97%	0.0	81.82%	200851	NW_015909858.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold73784	<i>Nicotiana tabacum</i>	2684	4,32E+03	97%	0.0	81.82%	69102	NW_015933844.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold77066	<i>Nicotiana tabacum</i>	2684	4,31E+03	99%	0.0	81.81%	66194	NW_015936658.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8771	<i>Nicotiana tabacum</i>	2684	4,32E+03	97%	0.0	81.81%	61761	NW_015945587.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold10118	<i>Nicotiana sylvestris</i>	2684	4,32E+03	97%	0.0	81.81%	78143	NW_009340115.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold12429	<i>Nicotiana sylvestris</i>	2684	4,32E+03	97%	0.0	81.81%	15007	NW_009365792.1

CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold17472	<i>Nicotiana sylvestris</i>	2684	4,32E+03	98%	0.0	81.85%	90010	NW_009421825.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold24346	<i>Nicotiana sylvestris</i>	2684	4,32E+03	97%	0.0	81.81%	52571	NW_009498203.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold42360	<i>Nicotiana sylvestris</i>	2684	8,62E+03	97%	0.0	81.81%	49038	NW_009528677.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold70476	<i>Nicotiana sylvestris</i>	2684	7,54E+03	98%	0.0	81.81%	119459	NW_009559916.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold73465	<i>Nicotiana sylvestris</i>	2684	8,62E+03	98%	0.0	81.82%	98251	NW_009563237.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold20485	<i>Nicotiana tabacum</i>	2682	4,33E+03	96%	0.0	82.08%	159734	NW_015854099.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold31263	<i>Nicotiana tabacum</i>	2682	4,29E+03	96%	0.0	82.09%	47106	NW_015889240.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold57563	<i>Nicotiana tabacum</i>	2682	4,31E+03	97%	0.0	81.99%	64684	NW_015919640.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold11114	<i>Nicotiana sylvestris</i>	2682	4,31E+03	97%	0.0	81.81%	57395	NW_009351182.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold14701	<i>Nicotiana sylvestris</i>	2682	4,30E+03	97%	0.0	81.80%	86106	NW_009391037.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13430	<i>Nicotiana tabacum</i>	2680	4,31E+03	96%	0.0	82.12%	68702	NW_015812463.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold27829	<i>Nicotiana tabacum</i>	2678	4,30E+03	97%	0.0	81.79%	55049	NW_015886005.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold57033	<i>Nicotiana sylvestris</i>	2678	6,30E+03	97%	0.0	81.79%	34725	NW_009544980.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold126703	<i>Nicotiana tabacum</i>	2676	4,31E+03	98%	0.0	81.79%	31219	NW_015807164.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold54562	<i>Nicotiana tabacum</i>	2675	4,12E+03	90%	0.0	81.88%	77179	NW_015916924.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24465	<i>Nicotiana tabacum</i>	2673	4,31E+03	98%	0.0	81.76%	216158	NW_015872997.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold45667	<i>Nicotiana tabacum</i>	2673	5,75E+03	97%	0.0	81.75%	108823	NW_015908758.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold85457	<i>Nicotiana tabacum</i>	2673	4,29E+03	97%	0.0	81.77%	60309	NW_015943734.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold772	<i>Nicotiana sylvestris</i>	2673	4,31E+03	96%	0.0	81.87%	70679	NW_009567386.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold5327	<i>Nicotiana sylvestris</i>	2671	4,70E+03	97%	0.0	81.75%	63779	NW_009543084.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold44740	<i>Nicotiana sylvestris</i>	2667	5,47E+03	98%	0.0	81.79%	85499	NW_009531321.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold86986	<i>Nicotiana sylvestris</i>	2654	5,63E+03	98%	0.0	81.67%	39570	NW_009578259.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold76778	<i>Nicotiana tabacum</i>	2652	4,29E+03	96%	0.0	81.93%	37160	NW_015936416.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold38145	<i>Nicotiana tabacum</i>	2651	4,19E+03	97%	0.0	81.89%	29522	NW_015901818.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold11621	<i>Nicotiana sylvestris</i>	2645	4,28E+03	97%	0.0	81.72%	209220	NW_009356815.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold4264	<i>Nicotiana sylvestris</i>	2645	4,70E+03	96%	0.0	82.08%	185422	NW_009528987.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold47780	<i>Nicotiana sylvestris</i>	2641	5,39E+03	97%	0.0	81.75%	51280	NW_009534698.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold43939	<i>Nicotiana sylvestris</i>	2627	4,04E+03	91%	0.0	81.81%	109796	NW_009530430.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold48501	<i>Nicotiana tabacum</i>	2621	4,23E+03	91%	0.0	82.95%	72251	NW_015911376.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold116498	<i>Nicotiana tabacum</i>	2615	4,29E+03	91%	0.0	82.92%	85930	NW_015799768.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold11117	<i>Nicotiana tabacum</i>	2612	4,24E+03	92%	0.0	82.67%	168162	NW_015795818.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8459	<i>Nicotiana tabacum</i>	2603	4,22E+03	97%	0.0	81.56%	55876	NW_015943004.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold45200	<i>Nicotiana tabacum</i>	2601	4,22E+03	97%	0.0	81.53%	112687	NW_015908326.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold75712	<i>Nicotiana tabacum</i>	2599	5,24E+03	91%	0.0	82.97%	66547	NW_015935507.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold16434	<i>Nicotiana sylvestris</i>	2588	5,11E+03	97%	0.0	81.54%	29763	NW_009410292.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold57650	<i>Nicotiana sylvestris</i>	2584	4,06E+03	92%	0.0	81.41%	17623	NW_009545665.1
CopiaCa_35	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	2577	4,21E+03	97%	0.0	81.42%	262618	NW_017671692.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13726	<i>Nicotiana tabacum</i>	2573	4,02E+03	97%	0.0	81.26%	75802	NW_015814528.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold25287	<i>Nicotiana sylvestris</i>	2558	4,18E+03	97%	0.0	81.24%	9436	NW_009508658.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold10970	<i>Nicotiana sylvestris</i>	2551	6,69E+03	85%	0.0	81.60%	109043	NW_009349581.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold27274	<i>Nicotiana sylvestris</i>	2551	4,08E+03	96%	0.0	81.45%	62986	NW_009511914.1
CopiaCa_35	refeq-genome	8	Nicotiana attenuata strain UT chromosome 7, NIATT2	<i>Nicotiana attenuata</i>	2549	1,15E+04	91%	0.0	81.37%	114765206	NC_031995.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10718	<i>Nicotiana tabacum</i>	2532	5,08E+03	98%	0.0	81.02%	65994	NW_015792768.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8288	<i>Nicotiana tabacum</i>	2527	4,09E+03	91%	0.0	82.39%	88475	NW_015941604.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1060	<i>Nicotiana sylvestris</i>	2527	4,10E+03	96%	0.0	81.46%	33055	NW_009345470.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold21037	<i>Nicotiana sylvestris</i>	2523	3,80E+03	89%	0.0	81.11%	66317	NW_009461437.1
CopiaCa_35	refeq-genome	8	Nicotiana attenuata strain UT chromosome 9, NIATT2	<i>Nicotiana attenuata</i>	2514	7,99E+03	98%	0.0	80.94%	50975570	NC_031997.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold3881	<i>Nicotiana sylvestris</i>	2483	6,07E+03	81%	0.0	83.51%	56468	NW_009524731.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold56271	<i>Nicotiana sylvestris</i>	2473	4,11E+03	85%	0.0	83.60%	128651	NW_009544133.1
CopiaCa_35	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	2457	4,07E+03	97%	0.0	80.71%	147415	NW_017673182.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold119754	<i>Nicotiana sylvestris</i>	2446	4,08E+03	84%	0.0	83.74%	21668	NW_009360752.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold18799	<i>Nicotiana sylvestris</i>	2444	4,41E+03	92%	0.0	82.23%	244701	NW_009436568.1
CopiaCa_35	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	2442	4,04E+03	98%	0.0	80.54%	210470	NW_017672203.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold43406	<i>Nicotiana sylvestris</i>	2425	5,89E+03	97%	0.0	81.55%	75838	NW_009529839.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold41151	<i>Nicotiana sylvestris</i>	2423	4,06E+03	84%	0.0	83.65%	26785	NW_009527334.1
CopiaCa_35	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	2418	3,96E+03	97%	0.0	80.42%	615692	NW_017670515.1

CopiaCa_35	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	2416	6,92E+03	91%	0.0	81.78%	432553	NW_017670854.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold18006	<i>Nicotiana sylvestris</i>	2403	4,03E+03	90%	0.0	82.58%	15207	NW_009427759.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold30390	<i>Nicotiana sylvestris</i>	2401	4,04E+03	92%	0.0	81.51%	24551	NW_009515377.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold16090	<i>Nicotiana sylvestris</i>	2398	4,03E+03	84%	0.0	83.66%	59265	NW_009406470.1
CopiaCa_35	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold42570	<i>Nicotiana tomentosiformis</i>	2386	8,00E+03	88%	0.0	82.63%	72771	NW_008924231.1
CopiaCa_35	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4354	<i>Nicotiana tomentosiformis</i>	2386	5,18E+03	88%	0.0	82.62%	90878	NW_008925308.1
CopiaCa_35	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold892	<i>Nicotiana tomentosiformis</i>	2381	8,01E+03	88%	0.0	82.59%	190525	NW_008976040.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold11260	<i>Nicotiana sylvestris</i>	2377	4,04E+03	84%	0.0	83.53%	187450	NW_009352804.1
CopiaCa_35	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold44434	<i>Nicotiana tomentosiformis</i>	2375	4,01E+03	87%	0.0	82.65%	215043	NW_008926197.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold104025	<i>Nicotiana tabacum</i>	2374	3,22E+03	81%	0.0	80.31%	15581	NW_015790369.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold36793	<i>Nicotiana sylvestris</i>	2366	5,80E+03	86%	0.0	82.19%	37617	NW_009522490.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold97760	<i>Nicotiana tabacum</i>	2362	3,82E+03	96%	0.0	80.35%	38281	NW_015953711.1
CopiaCa_35	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold33285	<i>Nicotiana tomentosiformis</i>	2362	4,01E+03	87%	0.0	82.57%	148793	NW_008913914.1
CopiaCa_35	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	2361	5,57E+03	97%	0.0	80.17%	535319	NW_017670623.1
CopiaCa_35	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold11745	<i>Nicotiana tomentosiformis</i>	2353	4,00E+03	88%	0.0	82.41%	74301	NW_008847880.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold5517	<i>Nicotiana sylvestris</i>	2351	4,51E+03	85%	0.0	82.90%	133282	NW_009542909.1
CopiaCa_35	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16270	<i>Nicotiana tomentosiformis</i>	2350	3,99E+03	88%	0.0	82.27%	128676	NW_008895009.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold39517	<i>Nicotiana tabacum</i>	2344	3,26E+03	82%	0.0	80.18%	68001	NW_015903087.1
CopiaCa_35	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold46707	<i>Nicotiana tomentosiformis</i>	2337	3,95E+03	88%	0.0	82.30%	129297	NW_008928827.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold81330	<i>Nicotiana sylvestris</i>	2326	3,92E+03	85%	0.0	82.65%	32773	NW_009571977.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13677	<i>Nicotiana tabacum</i>	2318	3,55E+03	88%	0.0	80.10%	210890	NW_015814184.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold33614	<i>Nicotiana tabacum</i>	2302	3,97E+03	84%	0.0	82.77%	49368	NW_015891460.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold43324	<i>Nicotiana tabacum</i>	2302	3,92E+03	81%	0.0	83.88%	79402	NW_015906612.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold18160	<i>Nicotiana tabacum</i>	2300	3,59E+03	88%	0.0	79.93%	53219	NW_015841694.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold149028	<i>Nicotiana tabacum</i>	2296	4,29E+03	97%	0.0	81.76%	21928	NW_015822282.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold14314	<i>Nicotiana sylvestris</i>	2274	4,40E+03	95%	0.0	81.80%	142707	NW_009386737.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold34568	<i>Nicotiana tabacum</i>	2268	4,29E+03	97%	0.0	81.67%	41091	NW_015892406.1
CopiaCa_35	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold61868	<i>Nicotiana tomentosiformis</i>	2261	3,58E+03	83%	0.0	81.81%	32916	NW_008945672.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold63832	<i>Nicotiana tabacum</i>	2242	3,53E+03	82%	0.0	81.71%	55722	NW_015925227.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold50661	<i>Nicotiana sylvestris</i>	2231	3,72E+03	89%	0.0	81.19%	47467	NW_009537900.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold39050	<i>Nicotiana sylvestris</i>	2230	3,80E+03	85%	0.0	81.90%	58263	NW_009524999.1
CopiaCa_35	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	2213	3,63E+03	82%	0.0	81.44%	6613	NW_017685767.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold35096	<i>Nicotiana sylvestris</i>	2209	4,36E+03	82%	0.0	81.70%	48636	NW_009520605.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold19268	<i>Nicotiana tabacum</i>	2207	9,56E+03	92%	0.0	83.10%	107039	NW_015847773.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7886	<i>Nicotiana tabacum</i>	2206	3,68E+03	85%	0.0	81.90%	36287	NW_015938204.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24092	<i>Nicotiana tabacum</i>	2196	3,79E+03	88%	0.0	81.37%	71342	NW_015871287.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold28180	<i>Nicotiana tabacum</i>	2185	3,76E+03	87%	0.0	81.34%	31379	NW_015886337.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1169	<i>Nicotiana tabacum</i>	2170	4,18E+03	94%	0.0	81.86%	118507	NW_01580066.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold61005	<i>Nicotiana tabacum</i>	2170	4,04E+03	97%	0.0	80.94%	14812	NW_015922689.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold9638	<i>Nicotiana sylvestris</i>	2169	5,06E+03	84%	0.0	83.31%	235636	NW_009588697.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24228	<i>Nicotiana tabacum</i>	2150	3,89E+03	95%	0.0	81.21%	90245	NW_015871912.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold18240	<i>Nicotiana sylvestris</i>	2146	3,60E+03	87%	0.0	81.09%	50093	NW_009430359.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold39223	<i>Nicotiana sylvestris</i>	2141	3,81E+03	92%	0.0	80.77%	91384	NW_009525191.1
CopiaCa_35	refeq-genome	8	Solanum pennellii chromosome 6, SPENNv200	<i>Solanum pennellii</i>	2139	1,88E+04	91%	0.0	80.62%	60730942	NC_028642.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold40064	<i>Nicotiana tabacum</i>	2134	3,66E+03	88%	0.0	80.97%	69012	NW_015903590.1
CopiaCa_35	refeq-genome	8	Nicotiana attenuata strain UT chromosome 10, NIATTR2	<i>Nicotiana attenuata</i>	2128	6,51E+03	91%	0.0	80.36%	51082206	NC_031988.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold29436	<i>Nicotiana tabacum</i>	2113	3,99E+03	95%	0.0	81.25%	133853	NW_015887512.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold32660	<i>Nicotiana tabacum</i>	2111	3,44E+03	83%	0.0	81.06%	61121	NW_015890550.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold26301	<i>Nicotiana tabacum</i>	2102	3,98E+03	96%	0.0	80.67%	57823	NW_015881398.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold24492	<i>Nicotiana sylvestris</i>	2095	4,04E+03	91%	0.0	80.84%	48786	NW_009499825.1
CopiaCa_35	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold76839	<i>Nicotiana tomentosiformis</i>	2091	5,79E+03	82%	0.0	82.79%	7407	NW_008962306.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold10566	<i>Nicotiana sylvestris</i>	2080	4,13E+03	81%	0.0	80.97%	55123	NW_009345092.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold39418	<i>Nicotiana sylvestris</i>	2063	3,52E+03	89%	0.0	80.30%	161234	NW_009525407.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold60047	<i>Nicotiana sylvestris</i>	2061	3,43E+03	83%	0.0	81.38%	50218	NW_009548329.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold8429	<i>Nicotiana sylvestris</i>	2060	3,46E+03	96%	0.0	78.89%	63105	NW_009575264.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold25495	<i>Nicotiana sylvestris</i>	2056	3,41E+03	86%	0.0	80.48%	14424	NW_009509937.1

CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold30504	<i>Nicotiana tabacum</i>	2049	3,42E+03	86%	0.0	80.45%	72675	NW_015888515.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold79425	<i>Nicotiana tabacum</i>	2047	3,44E+03	88%	0.0	80.52%	82404	NW_015938673.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold12157	<i>Nicotiana sylvestris</i>	2047	3,59E+03	90%	0.0	80.52%	129264	NW_009362770.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold59573	<i>Nicotiana tabacum</i>	2039	3,58E+03	87%	0.0	80.49%	44109	NW_015921401.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold35466	<i>Nicotiana tabacum</i>	2034	3,63E+03	82%	0.0	81.87%	74899	NW_015899295.1
CopiaCa_35	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	2019	3,18E+03	81%	0.0	80.45%	66948	NW_017675454.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold40956	<i>Nicotiana tabacum</i>	2017	3,87E+03	96%	0.0	80.42%	65777	NW_015904408.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold37816	<i>Nicotiana tabacum</i>	2013	3,84E+03	95%	0.0	80.66%	138130	NW_015901499.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold35490	<i>Nicotiana tabacum</i>	2012	3,32E+03	81%	0.0	80.24%	60584	NW_015899320.1
CopiaCa_35	refeq-genome	8	Nicotiana attenuata strain UT chromosome 6, NIATT2	<i>Nicotiana attenuata</i>	2006	8,84E+03	81%	0.0	82.14%	97571839	NC_031994.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold22438	<i>Nicotiana tabacum</i>	2001	3,83E+03	96%	0.0	80.07%	150234	NW_015863524.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold28556	<i>Nicotiana sylvestris</i>	1962	3,60E+03	92%	0.0	79.99%	167357	NW_009513338.1
CopiaCa_35	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	1958	3,28E+03	82%	0.0	80.05%	52228	NW_017676094.1
CopiaCa_35	refeq-genome	8	Solanum pennellii chromosome 4, SPENN200	<i>Solanum pennellii</i>	1951	2,56E+04	82%	0.0	81.75%	77197300	NC_028640.1
CopiaCa_35	refeq-genome	8	Solanum pennellii chromosome 3, SPENN200	<i>Solanum pennellii</i>	1940	3,78E+04	90%	0.0	81.72%	75414019	NC_028639.1
CopiaCa_35	refeq-genome	8	Solanum pennellii chromosome 9, SPENN200	<i>Solanum pennellii</i>	1940	2,29E+04	82%	0.0	81.71%	84057508	NC_028645.1
CopiaCa_35	refeq-genome	8	Solanum pennellii chromosome 5, SPENN200	<i>Solanum pennellii</i>	1934	1,47E+04	82%	0.0	81.62%	77991103	NC_028641.1
CopiaCa_35	refeq-genome	8	Solanum pennellii chromosome 10, SPENN200	<i>Solanum pennellii</i>	1929	2,71E+04	90%	0.0	82.44%	82529941	NC_028646.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold9090	<i>Nicotiana sylvestris</i>	1929	3,80E+03	98%	0.0	79.60%	169429	NW_009582609.1
CopiaCa_35	refeq-genome	8	Solanum pennellii chromosome 1, SPENN200	<i>Solanum pennellii</i>	1923	1,85E+04	91%	0.0	82.40%	109333515	NC_028637.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold22534	<i>Nicotiana sylvestris</i>	1923	4,17E+03	97%	0.0	79.79%	72389	NW_009478070.1
CopiaCa_35	refeq-genome	8	Solanum pennellii chromosome 7, SPENN200	<i>Solanum pennellii</i>	1917	1,99E+04	91%	0.0	82.35%	79292169	NC_028643.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold75417	<i>Nicotiana tabacum</i>	1897	3,74E+03	89%	0.0	81.35%	47045	NW_015935246.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold34623	<i>Nicotiana sylvestris</i>	1895	3,36E+03	82%	0.0	80.86%	61985	NW_009520080.1
CopiaCa_35	refeq-genome	8	Solanum pennellii chromosome 2, SPENN200	<i>Solanum pennellii</i>	1884	1,03E+04	82%	0.0	81.25%	59803892	NC_028638.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold69689	<i>Nicotiana tabacum</i>	1884	3,81E+03	89%	0.0	81.48%	49780	NW_015930317.1
CopiaCa_35	refeq-genome	8	Solanum pennellii chromosome 11, SPENN200	<i>Solanum pennellii</i>	1877	9,92E+03	82%	0.0	81.22%	66223686	NC_028647.1
CopiaCa_35	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	<i>Solanum lycopersicum</i>	1873	8,88E+03	82%	0.0	81.17%	66557038	NC_015441.3
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold18555	<i>Nicotiana sylvestris</i>	1862	3,73E+03	90%	0.0	81.26%	112746	NW_009433858.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold11601	<i>Nicotiana sylvestris</i>	1858	3,55E+03	80%	0.0	82.26%	99084	NW_009356593.1
CopiaCa_35	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3330	<i>Nicotiana tomentosiformis</i>	1857	3,58E+03	82%	0.0	81.55%	338133	NW_008913931.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold34002	<i>Nicotiana tabacum</i>	1851	3,57E+03	82%	0.0	81.51%	108041	NW_015891825.1
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold78293	<i>Nicotiana tabacum</i>	1825	3,27E+03	81%	0.0	80.84%	27611	NW_015937720.1
CopiaCa_35	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	<i>Solanum lycopersicum</i>	1820	1,60E+04	90%	0.0	79.33%	72906345	NC_015446.3
CopiaCa_35	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	<i>Solanum lycopersicum</i>	1816	9,36E+03	91%	0.0	80.74%	68175699	NC_015444.3
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold36157	<i>Nicotiana sylvestris</i>	1796	4,97E+03	89%	0.0	82.70%	38647	NW_009521784.1
CopiaCa_35	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	<i>Solanum lycopersicum</i>	1775	1,26E+04	82%	0.0	81.22%	98455869	NC_015438.3
CopiaCa_35	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold11011	<i>Nicotiana tabacum</i>	1764	3,47E+03	80%	0.0	81.04%	170479	NW_015795033.1
CopiaCa_35	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold87086	<i>Nicotiana sylvestris</i>	1760	4,24E+03	91%	0.0	83.50%	22610	NW_009578371.1
CopiaCa_37	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9487	1,79E+04	100%	0.0	100.00%	219521584	NC_029982.1
CopiaCa_37	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7265	1,58E+04	100%	0.0	92.14%	220335243	NC_029987.1
CopiaCa_37	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7003	1,38E+04	100%	0.0	91.22%	215701946	NC_029980.1
CopiaCa_37	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1245	<i>Capsicum annuum</i>	6656	7,45E+03	100%	0.0	90.21%	307737	NW_015960675.1
CopiaCa_37	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6360	1,85E+04	99%	0.0	90.37%	301019445	NC_029977.1
CopiaCa_37	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5099	6,12E+03	90%	0.0	88.07%	229934170	NC_029988.1
CopiaCa_37	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1246	<i>Capsicum annuum</i>	5051	7,65E+03	98%	0.0	90.70%	3388803	NW_015960834.1
CopiaCa_37	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold756	<i>Capsicum annuum</i>	4909	9,72E+03	99%	0.0	90.94%	884456	NW_015961254.1
CopiaCa_37	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4220	5,69E+03	94%	0.0	87.56%	217274494	NC_029981.1
CopiaCa_37	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3882	8,10E+03	100%	0.0	92.15%	222112641	NC_029983.1
CopiaCa_37	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3092	7,27E+03	88%	0.0	91.70%	205736368	NC_029986.1
CopiaCa_41	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10543	1,27E+04	100%	0.0	100.00%	205736368	NC_029986.1
CopiaCa_41	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold65545	<i>Capsicum annuum</i>	5622	8,94E+03	97%	0.0	93.25%	15526	NW_015961722.1
CopiaCa_41	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2610	6,15E+03	89%	0.0	80.15%	238794889	NC_029985.1
CopiaCa_41	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2252	4,53E+03	85%	0.0	81.82%	153299543	NC_029984.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7697	1,12E+08	100%	0.0	100.00%	229934170	NC_029988.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4307	1,21E+08	100%	0.0	89.51%	238794889	NC_029985.1

CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4257	1,23E+08	100%	0.0	89.12%	215701946	NC_029980.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4156	1,07E+08	100%	0.0	88.84%	222112641	NC_029983.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold376	<i>Capsicum annuum</i>	3989	4,29E+03	85%	0.0	88.27%	776404	NW_015961666.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3611	1,62E+08	100%	0.0	90.32%	261510930	NC_029977.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3598	1,61E+08	100%	0.0	90.20%	301019445	NC_029977.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3592	8,00E+04	100%	0.0	90.12%	153299543	NC_029984.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4912	<i>Capsicum annuum</i>	3592	4,42E+03	83%	0.0	90.17%	178551	NW_015960903.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3552	1,29E+08	100%	0.0	89.90%	205736368	NC_029986.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3524	1,03E+08	100%	0.0	88.88%	220335243	NC_029987.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3489	1,08E+08	100%	0.0	89.65%	163962470	NC_029978.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3489	8,14E+04	95%	0.0	89.50%	217274494	NC_029981.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold891	<i>Capsicum annuum</i>	3480	6,76E+03	83%	0.0	89.44%	1431046	NW_015960469.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2322	<i>Capsicum annuum</i>	3478	1,09E+04	99%	0.0	89.47%	1668020	NW_015961155.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3226	<i>Capsicum annuum</i>	3358	4,57E+03	90%	0.0	88.67%	1101908	NW_015960413.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1038	<i>Capsicum annuum</i>	3328	7,80E+03	83%	0.0	88.63%	2164954	NW_015960831.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold8630	<i>Capsicum annuum</i>	2950	4,48E+03	85%	0.0	88.13%	66653	NW_015961494.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2944	7,76E+04	100%	0.0	88.93%	219521584	NC_029982.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold13708	<i>Capsicum annuum</i>	2446	6,60E+03	88%	0.0	83.96%	186116	NW_015960578.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold189	<i>Capsicum annuum</i>	2436	7,45E+03	90%	0.0	88.70%	3791111	NW_015960691.1
CopiaCa_43	refeq-genome	8	Solanum pennellii chromosome 9, SPENN200	<i>Solanum pennellii</i>	2377	1,70E+04	82%	0.0	83.45%	84057508	NC_028645.1
CopiaCa_43	refeq-genome	8	Solanum pennellii chromosome 7, SPENN200	<i>Solanum pennellii</i>	2355	1,81E+04	80%	0.0	83.31%	79292169	NC_028643.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1390	<i>Capsicum annuum</i>	2346	4,93E+03	81%	0.0	85.94%	1730405	NW_015960783.1
CopiaCa_43	refeq-genome	8	Solanum pennellii chromosome 11, SPENN200	<i>Solanum pennellii</i>	2292	1,32E+04	80%	0.0	82.79%	66223686	NC_028647.1
CopiaCa_43	refeq-genome	8	Solanum pennellii chromosome 12, SPENN200	<i>Solanum pennellii</i>	2278	2,51E+04	82%	0.0	82.84%	83305730	NC_028648.1
CopiaCa_43	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	<i>Solanum lycopersicum</i>	2185	1,33E+04	82%	0.0	82.09%	56597135	NC_015448.3
CopiaCa_43	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	<i>Solanum lycopersicum</i>	1886	9,95E+03	81%	0.0	80.34%	72906345	NC_015446.3
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1434	<i>Capsicum annuum</i>	1857	8,01E+03	91%	0.0	88.78%	1100793	NW_015960373.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold90	<i>Capsicum annuum</i>	1712	4,78E+03	87%	0.0	88.28%	6074945	NW_015960458.1
CopiaCa_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold349	<i>Capsicum annuum</i>	1286	6,68E+03	84%	0.0	84.55%	2130419	NW_015960729.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8658	9,71E+04	100%	0.0	100.00%	261510930	NC_029979.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6141	8,03E+04	100%	0.0	93.04%	238794889	NC_029985.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6085	1,32E+08	100%	0.0	92.73%	301019445	NC_029977.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5980	7,47E+04	100%	0.0	92.31%	215701946	NC_029980.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5936	5,11E+04	100%	0.0	92.10%	153299543	NC_029984.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capsicum annuum</i>	5746	9,06E+03	100%	0.0	91.28%	2397553	NW_015961318.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5729	3,76E+04	100%	0.0	91.19%	205736368	NC_029986.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5659	9,22E+04	100%	0.0	90.92%	219521584	NC_029982.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5480	1,16E+08	100%	0.0	90.17%	229934170	NC_029988.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5426	6,37E+04	100%	0.0	89.94%	222112641	NC_029983.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5398	1,36E+08	100%	0.0	89.82%	220335243	NC_029987.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1765	<i>Capsicum annuum</i>	5332	1,49E+04	95%	0.0	89.52%	289511	NW_015961400.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9	<i>Capsicum annuum</i>	5083	5,08E+03	89%	0.0	88.52%	6567951	NW_015961815.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1246	<i>Capsicum annuum</i>	5012	6,18E+03	95%	0.0	88.26%	3388803	NW_015960834.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold16	<i>Capsicum annuum</i>	4776	9,89E+03	95%	0.0	87.14%	2023789	NW_015960595.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold189	<i>Capsicum annuum</i>	4656	1,17E+04	97%	0.0	86.76%	3791111	NW_015960691.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4621	7,12E+04	100%	0.0	92.32%	163962470	NC_029978.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00554	<i>Solanum tuberosum</i>	4545	4,63E+03	91%	0.0	86.16%	306819	NW_006239480.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4516	8,33E+04	100%	0.0	88.83%	217274494	NC_029981.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00148	<i>Solanum tuberosum</i>	4481	4,73E+03	88%	0.0	87.15%	1437876	NW_006239071.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00533	<i>Solanum tuberosum</i>	4475	8,48E+03	91%	0.0	86.23%	333517	NW_006239459.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00629	<i>Solanum tuberosum</i>	4205	4,32E+03	93%	0.0	84.84%	239099	NW_006239549.1
CopiaCa_45	refeq-genome	8	Solanum pennellii chromosome 11, SPENN200	<i>Solanum pennellii</i>	4187	1,23E+04	92%	0.0	84.75%	66223686	NC_028647.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00017	<i>Solanum tuberosum</i>	4145	4,42E+03	89%	0.0	85.67%	3280810	NW_006238943.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00232	<i>Solanum tuberosum</i>	3991	4,55E+03	89%	0.0	85.76%	1045062	NW_006239158.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00005	<i>Solanum tuberosum</i>	3971	5,70E+03	84%	0.0	84.83%	4371776	NW_006238931.1
CopiaCa_45	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	<i>Solanum lycopersicum</i>	3903	8,57E+03	89%	0.0	85.24%	66557038	NC_015441.3

CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold756	<i>Capsicum annuum</i>	3899	6,65E+03	95%	0.0	90.74%	884456	NW_015961254.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00188	<i>Solanum tuberosum</i>	3816	3,82E+03	84%	0.0	84.11%	1178348	NW_006239119.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00070	<i>Solanum tuberosum</i>	3807	3,81E+03	84%	0.0	84.06%	2061072	NW_006238996.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold519	<i>Capsicum annuum</i>	3794	5,92E+03	80%	0.0	91.61%	1101621	NW_015961578.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00022	<i>Solanum tuberosum</i>	3757	7,39E+03	84%	0.0	84.99%	3032639	NW_006238948.1
CopiaCa_45	refeq-genome	8	Solanum pennellii chromosome 1, SPENN200	<i>Solanum pennellii</i>	3672	2,60E+04	97%	0.0	83.49%	109333515	NC_028637.1
CopiaCa_45	refeq-genome	8	Solanum pennellii chromosome 12, SPENN200	<i>Solanum pennellii</i>	3664	1,73E+04	93%	0.0	82.56%	83305730	NC_028648.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1399	<i>Capsicum annuum</i>	3648	6,16E+03	88%	0.0	89.33%	1742971	NW_015961630.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00662	<i>Solanum tuberosum</i>	3605	3,61E+03	84%	0.0	83.27%	215247	NW_006239580.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00063	<i>Solanum tuberosum</i>	3568	1,07E+04	89%	0.0	84.14%	2166112	NW_006238988.1
CopiaCa_45	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	<i>Solanum lycopersicum</i>	3565	1,48E+04	92%	0.0	82.05%	66723567	NC_015442.3
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00446	<i>Solanum tuberosum</i>	3550	5,26E+03	89%	0.0	84.43%	469271	NW_006239368.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00007	<i>Solanum tuberosum</i>	3535	5,65E+03	89%	0.0	82.11%	4094051	NW_006238933.1
CopiaCa_45	refeq-genome	8	Nicotiana attenuata strain UT chromosome 2, NIATT2	<i>Nicotiana attenuata</i>	3530	5,49E+03	84%	0.0	86.75%	96359640	NC_031990.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00175	<i>Solanum tuberosum</i>	3520	6,49E+03	87%	0.0	82.36%	1280083	NW_006239098.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00185	<i>Solanum tuberosum</i>	3507	4,07E+03	86%	0.0	84.65%	1171467	NW_006239122.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00071	<i>Solanum tuberosum</i>	3472	1,38E+04	84%	0.0	82.62%	2054262	NW_006238997.1
CopiaCa_45	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	3446	4,20E+03	84%	0.0	86.29%	113208	NW_017673983.1
CopiaCa_45	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	3421	4,22E+03	84%	0.0	86.19%	11831	NW_017681131.1
CopiaCa_45	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	<i>Solanum lycopersicum</i>	3413	1,06E+04	91%	0.0	82.74%	98455869	NC_015438.3
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00162	<i>Solanum tuberosum</i>	3410	3,41E+03	84%	0.0	82.23%	1340945	NW_006239088.1
CopiaCa_45	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold77694	<i>Nicotiana tabacum</i>	3304	4,01E+03	84%	0.0	85.49%	62952	NW_015937204.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf0024	<i>Solanum tuberosum</i>	3201	7,93E+03	87%	0.0	84.86%	2984907	NW_006238950.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00067	<i>Solanum tuberosum</i>	3199	9,30E+03	89%	0.0	86.89%	2156588	NW_006238991.1
CopiaCa_45	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	<i>Solanum lycopersicum</i>	3199	4,30E+03	85%	0.0	81.42%	68175699	NC_015444.3
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	3153	4,12E+03	86%	0.0	85.43%	3493078	NW_015960517.1
CopiaCa_45	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold55798	<i>Nicotiana sylvestris</i>	3151	4,61E+03	84%	0.0	85.28%	52338	NW_009543606.1
CopiaCa_45	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold22748	<i>Nicotiana tabacum</i>	3123	4,59E+03	84%	0.0	85.12%	46108	NW_015865027.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00455	<i>Solanum tuberosum</i>	3094	3,91E+03	89%	0.0	83.96%	461042	NW_006239371.1
CopiaCa_45	refeq-genome	8	Solanum pennellii chromosome 7, SPENN200	<i>Solanum pennellii</i>	3025	5,12E+03	94%	0.0	80.57%	79292169	NC_028643.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Heinz 1706 chromosome 9, SL3.0	<i>Capsicum annuum</i>	3020	5,96E+03	92%	0.0	90.86%	913988	NW_015961387.1
CopiaCa_45	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold14202	<i>Nicotiana sylvestris</i>	3020	3,62E+03	84%	0.0	84.05%	171652	NW_009355881.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00297	<i>Solanum tuberosum</i>	3020	8,02E+03	91%	0.0	82.76%	814182	NW_006239220.1
CopiaCa_45	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	<i>Solanum lycopersicum</i>	3016	1,17E+04	90%	0.0	83.96%	72906345	NC_015446.3
CopiaCa_45	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold14202	<i>Nicotiana sylvestris</i>	3014	3,69E+03	83%	0.0	84.05%	214048	NW_009385493.1
CopiaCa_45	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold220809	<i>Nicotiana tabacum</i>	2998	3,58E+03	84%	0.0	83.92%	33449	NW_015861866.1
CopiaCa_45	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold2477	<i>Nicotiana tabacum</i>	2998	3,65E+03	84%	0.0	83.89%	158578	NW_015874368.1
CopiaCa_45	refeq-genome	8	Solanum pennellii chromosome 5, SPENN200	<i>Solanum pennellii</i>	2968	1,86E+04	94%	0.0	80.78%	77991103	NC_028641.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00411	<i>Solanum tuberosum</i>	2961	4,45E+03	84%	0.0	87.21%	508257	NW_006239441.1
CopiaCa_45	refeq-genome	8	Solanum pennellii chromosome 4, SPENN200	<i>Solanum pennellii</i>	2953	1,68E+04	91%	0.0	85.33%	77197300	NC_028640.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00106	<i>Solanum tuberosum</i>	2909	4,57E+03	84%	0.0	86.70%	1697319	NW_006239032.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	2867	3,84E+03	88%	0.0	83.55%	7059680	NW_015960855.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00013	<i>Solanum tuberosum</i>	2826	6,42E+03	80%	0.0	83.39%	3422333	NW_006238940.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00310	<i>Solanum tuberosum</i>	2800	3,78E+03	81%	0.0	85.87%	764739	NW_006239236.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf01132	<i>Solanum tuberosum</i>	2778	4,35E+03	87%	0.0	86.71%	22280	NW_006240048.1
CopiaCa_45	refeq-genome	8	Nicotiana attenuata strain UT chromosome 6, NIATT2	<i>Nicotiana attenuata</i>	2761	4,22E+03	86%	0.0	86.52%	97571839	NC_031994.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	2603	5,71E+03	89%	0.0	90.33%	2470984	NW_015961229.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1174	<i>Capsicum annuum</i>	2597	5,92E+03	93%	0.0	91.42%	656414	NW_015961459.1
CopiaCa_45	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold8508	<i>Nicotiana sylvestris</i>	2562	3,98E+03	82%	0.0	86.66%	44195	NW_009576142.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00189	<i>Solanum tuberosum</i>	2418	4,31E+03	84%	0.0	87.64%	1194366	NW_006239113.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00355	<i>Solanum tuberosum</i>	2351	5,92E+03	83%	0.0	83.36%	640234	NW_006239277.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00570	<i>Solanum tuberosum</i>	2351	5,01E+03	81%	0.0	86.34%	294773	NW_006239488.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00009	<i>Solanum tuberosum</i>	2305	4,37E+03	88%	0.0	81.52%	3955149	NW_006238935.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00207	<i>Solanum tuberosum</i>	2303	4,15E+03	84%	0.0	84.16%	1129882	NW_006239133.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold723	<i>Capsicum annuum</i>	2180	5,47E+03	84%	0.0	80.94%	559742	NW_015961191.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00255	<i>Solanum tuberosum</i>	2154	4,60E+03	83%	0.0	85.88%	929374	NW_006239184.1

CopiaCa_45	refeq-genome	8	Solanum pennellii chromosome 10, SPENN200	<i>Solanum pennellii</i>	2130	1,29E+04	91%	0.0	79.91%	82529941	NC_028646.1
CopiaCa_45	refeq-genome	8	Solanum pennellii chromosome 8, SPENN200	<i>Solanum pennellii</i>	2089	4,19E+03	90%	0.0	83.68%	70546659	NC_028644.1
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00113	<i>Solanum tuberosum</i>	2087	3,37E+03	81%	0.0	83.05%	1648986	NW_006239039.1
CopiaCa_45	refeq-genome	8	Solanum pennellii chromosome 9, SPENN200	<i>Solanum pennellii</i>	2060	1,01E+04	86%	0.0	83.54%	84057508	NC_028645.1
CopiaCa_45	refeq-genome	8	Solanum pennellii chromosome 2, SPENN200	<i>Solanum pennellii</i>	2056	1,02E+04	89%	0.0	80.96%	59803892	NC_028638.1
CopiaCa_45	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	<i>Solanum lycopersicum</i>	1953	9,77E+03	93%	0.0	81.02%	68126176	NC_015449.3
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2256	<i>Capsicum annuum</i>	1917	9,36E+03	83%	0.0	87.77%	41427	NW_015960920.1
CopiaCa_45	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	<i>Solanum lycopersicum</i>	1897	5,40E+03	89%	0.0	82.29%	65987440	NC_015445.3
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00279	<i>Solanum tuberosum</i>	1845	3,98E+03	83%	0.0	84.30%	864930	NW_006239204.1
CopiaCa_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold349	<i>Capsicum annuum</i>	1724	4,05E+03	84%	0.0	82.17%	2130419	NW_015960729.1
CopiaCa_45	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	<i>Solanum lycopersicum</i>	1692	8,21E+03	94%	0.0	83.30%	55977580	NC_015439.3
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00097	<i>Solanum tuberosum</i>	1690	4,70E+03	93%	0.0	87.59%	1770913	NW_006239022.1
CopiaCa_45	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	<i>Solanum lycopersicum</i>	1646	9,86E+03	84%	0.0	81.63%	65633393	NC_015447.3
CopiaCa_45	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00141	<i>Solanum tuberosum</i>	1482	4,34E+03	85%	0.0	83.81%	1429992	NW_006239075.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9029	3,02E+04	100%	0.0	100.00%	229934170	NC_029988.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3578	1,82E+04	100%	0.0	92.37%	205736368	NC_029986.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3323	3,95E+04	100%	0.0	90.58%	215701946	NC_029980.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3205	3,18E+04	100%	0.0	89.68%	261510930	NC_029979.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3155	1,56E+04	99%	0.0	89.70%	163962470	NC_029978.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3075	2,33E+04	100%	0.0	89.98%	222112641	NC_029983.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3049	1,92E+04	94%	0.0	88.94%	219521584	NC_029982.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3025	2,85E+04	94%	0.0	88.78%	301019445	NC_029977.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3005	4,14E+04	99%	0.0	88.33%	238794889	NC_029985.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3466	<i>Capsicum annuum</i>	2948	5,40E+03	92%	0.0	88.01%	143631	NW_015960416.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2944	5,53E+03	94%	0.0	88.17%	153299543	NC_029984.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold979	<i>Capsicum annuum</i>	2898	5,81E+03	99%	0.0	87.45%	1172749	NW_015961157.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold448	<i>Capsicum annuum</i>	2874	5,72E+03	90%	0.0	90.62%	1323857	NW_015960958.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4977	<i>Capsicum annuum</i>	2874	5,12E+03	92%	0.0	87.71%	543020	NW_015961055.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2835	2,07E+04	99%	0.0	87.89%	220335243	NC_029987.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2782	1,02E+04	94%	0.0	87.00%	217274494	NC_029981.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1730	<i>Capsicum annuum</i>	2761	5,10E+03	94%	0.0	87.02%	1080651	NW_015961122.1
CopiaCa_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3313	<i>Capsicum annuum</i>	1740	5,12E+03	83%	0.0	89.22%	770237	NW_015961029.1
monotypic-Ch01d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8602	5,10E+04	100%	0.0	100.00%	301019445	NC_029977.1
monotypic-Ch01d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6119	2,68E+04	99%	0.0	92.24%	215701946	NC_029980.1
monotypic-Ch01d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6017	2,53E+04	99%	0.0	93.29%	153299543	NC_029984.1
monotypic-Ch01d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5842	2,09E+04	99%	0.0	91.00%	217274494	NC_029981.1
monotypic-Ch01d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5692	2,38E+04	98%	0.0	90.59%	219521584	NC_029982.1
monotypic-Ch01d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5435	2,48E+04	98%	0.0	89.64%	238794889	NC_029985.1
monotypic-Ch01d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5086	2,85E+04	96%	0.0	89.04%	163962470	NC_029978.1
monotypic-Ch01d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1741	<i>Capsicum annuum</i>	5077	5,36E+03	95%	0.0	87.99%	1897442	NW_015960687.1
monotypic-Ch01d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5018	1,86E+04	92%	0.0	90.01%	229934170	NC_029988.1
monotypic-Ch01d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2998	<i>Capsicum annuum</i>	4427	6,15E+03	88%	0.0	93.40%	26483	NW_015961441.1
monotypic-Ch01d_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00665	<i>Solanum tuberosum</i>	4313	4,31E+03	90%	0.0	85.18%	215440	NW_006239579.1
monotypic-Ch01d_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00168	<i>Solanum tuberosum</i>	4143	4,74E+03	89%	0.0	84.73%	1293034	NW_006239096.1
monotypic-Ch01d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3965	3,45E+04	99%	0.0	85.09%	205736368	NC_029986.1
monotypic-Ch01d_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00003	<i>Solanum tuberosum</i>	3888	3,89E+03	90%	0.0	83.43%	459022	NW_006238929.1
monotypic-Ch01d_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00375	<i>Solanum tuberosum</i>	3773	3,77E+03	81%	0.0	84.83%	589396	NW_006239299.1
monotypic-Ch01d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3714	2,63E+04	99%	0.0	89.53%	220335243	NC_029987.1
monotypic-Ch01d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3703	2,38E+04	99%	0.0	89.53%	261510930	NC_029979.1
monotypic-Ch01d_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00236	<i>Solanum tuberosum</i>	3417	4,19E+03	89%	0.0	84.25%	1037213	NW_006239160.1
monotypic-Ch01d_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00176	<i>Solanum tuberosum</i>	3295	3,30E+03	87%	0.0	81.48%	1277157	NW_006239100.1
monotypic-Ch01d_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00407	<i>Solanum tuberosum</i>	3125	6,22E+03	89%	0.0	80.64%	532697	NW_006239324.1
monotypic-Ch01d_24	refeq-genome	8	Solanum pennellii chromosome 5, SPENN200	<i>Solanum pennellii</i>	2988	5,02E+03	82%	0.0	80.89%	77991103	NC_028641.1
monotypic-Ch01d_24	refeq-genome	8	Solanum pennellii chromosome 1, SPENN200	<i>Solanum pennellii</i>	2865	1,00E+04	89%	0.0	79.70%	109333515	NC_028637.1
monotypic-Ch01d_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00250	<i>Solanum tuberosum</i>	2861	8,59E+03	89%	0.0	85.71%	939400	NW_006239183.1
monotypic-Ch01d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold258	<i>Capsicum annuum</i>	2481	7,14E+03	85%	0.0	93.17%	3098259	NC_015961180.1

monotypic-Ch01d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2447	7,21E+03	93%	0.0	83.91%	222112641	NC_029983.1
monotypic-Ch01d_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00018	<i>Solanum tuberosum</i>	2427	4,14E+03	80%	0.0	87.24%	3280163	NW_006238944.1
monotypic-Ch01d_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00400	<i>Solanum tuberosum</i>	2370	4,17E+03	90%	0.0	84.90%	525930	NW_006239330.1
monotypic-Ch01d_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00123	<i>Solanum tuberosum</i>	2230	5,67E+03	81%	0.0	84.75%	1602102	NW_006239048.1
monotypic-Ch01d_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00029	<i>Solanum tuberosum</i>	2209	3,37E+03	87%	0.0	81.06%	2785282	NW_006238951.1
monotypic-Ch01d_24	refeq-genome	8	Solanum pennellii chromosome 11, SPENN200	<i>Solanum pennellii</i>	2206	3,14E+03	88%	0.0	79.99%	66223686	NC_028647.1
monotypic-Ch01d_24	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00071	<i>Solanum tuberosum</i>	2161	7,90E+03	89%	0.0	80.40%	2054262	NW_006238997.1
monotypic-Ch01d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold619	<i>Capsicum annuum</i>	1592	4,68E+03	89%	0.0	88.52%	2512642	NW_015960498.1
monotypic-Ch01f_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8841	8,93E+04	100%	0.0	100.00%	301019445	NC_029977.1
monotypic-Ch01f_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5147	6,29E+04	100%	0.0	93.50%	238794889	NC_029985.1
monotypic-Ch01f_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4669	6,27E+04	99%	0.0	91.52%	217274494	NC_029981.1
monotypic-Ch01f_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4669	4,39E+04	100%	0.0	91.05%	153299543	NC_029984.1
monotypic-Ch01f_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4639	6,09E+04	99%	0.0	91.12%	220335243	NC_029987.1
monotypic-Ch01f_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1271	<i>Capsicum annuum</i>	4370	5,99E+03	93%	0.0	92.63%	192768	NW_015961561.1
monotypic-Ch01f_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4287	6,47E+04	100%	0.0	89.30%	261510930	NC_029979.1
monotypic-Ch01f_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4130	5,82E+04	99%	0.0	88.40%	215701946	NC_029980.1
monotypic-Ch01f_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2003	<i>Capsicum annuum</i>	4078	6,66E+03	94%	0.0	92.42%	314382	NW_015961213.1
monotypic-Ch01f_34	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00528	<i>Solanum tuberosum</i>	3620	4,85E+03	95%	0.0	86.05%	350176	NW_006239446.1
monotypic-Ch01f_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3437	5,69E+04	99%	0.0	93.53%	219521584	NC_029982.1
monotypic-Ch01f_34	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00343	<i>Solanum tuberosum</i>	3378	4,52E+03	95%	0.0	84.74%	664124	NW_006239267.1
monotypic-Ch01f_34	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00079	<i>Solanum tuberosum</i>	3319	4,54E+03	99%	0.0	84.47%	1965697	NW_006239004.1
monotypic-Ch01f_34	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00332	<i>Solanum tuberosum</i>	3249	4,43E+03	92%	0.0	84.21%	692937	NW_006239255.1
monotypic-Ch01f_34	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00044	<i>Solanum tuberosum</i>	3101	4,11E+03	95%	0.0	83.42%	2434495	NW_006238970.1
monotypic-Ch01f_34	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7362	<i>Nicotiana tabacum</i>	2737	3,61E+03	83%	0.0	82.74%	72050	NW_015933702.1
monotypic-Ch01f_34	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31337	<i>Nicotiana tomentosiformis</i>	2732	3,61E+03	83%	0.0	82.71%	95121	NW_008911750.1
monotypic-Ch01f_34	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23933	<i>Nicotiana tabacum</i>	2623	3,50E+03	83%	0.0	82.08%	213559	NW_015870571.1
monotypic-Ch01f_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold181	<i>Capsicum annuum</i>	2495	4,62E+03	86%	0.0	83.20%	2703691	NW_015961388.1
monotypic-Ch01f_34	refeq-genome	8	Solanum pennellii chromosome 12, SPENN200	<i>Solanum pennellii</i>	2473	7,24E+03	98%	0.0	80.23%	83305730	NC_028648.1
monotypic-Ch01f_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold919	<i>Capsicum annuum</i>	2353	6,03E+03	93%	0.0	94.13%	689688	NW_015961602.1
monotypic-Ch01f_34	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00516	<i>Solanum tuberosum</i>	2313	3,54E+03	80%	0.0	81.76%	363052	NW_006239432.1
monotypic-Ch01f_34	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	<i>Solanum lycopersicum</i>	2102	1,05E+04	99%	0.0	78.36%	68126176	NC_015449.3
monotypic-Ch01f_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2004	4,99E+04	99%	0.0	89.58%	222112641	NC_029983.1
monotypic-Ch01f_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	1687	3,01E+04	84%	0.0	86.77%	163962470	NC_029978.1
monotypic-Ch01f_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold555	<i>Capsicum annuum</i>	1461	3,80E+03	88%	0.0	81.53%	1396435	NW_015960461.1
monotypic-Ch01f_35	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10033	1,14E+04	100%	0.0	100.00%	301019445	NC_029977.1
monotypic-Ch01f_35	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8161	9,20E+03	100%	0.0	93.86%	229934170	NC_029988.1
monotypic-Ch01f_35	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7978	9,63E+03	100%	0.0	93.19%	205736368	NC_029986.1
monotypic-Ch01f_35	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7553	1,32E+04	100%	0.0	93.87%	261510930	NC_029979.1
monotypic-Ch01f_35	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6874	8,46E+03	99%	0.0	89.63%	219521584	NC_029982.1
monotypic-Ch01f_35	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5609	9,05E+03	100%	0.0	89.82%	215701946	NC_029980.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8961	1,62E+08	100%	0.0	100.00%	163962470	NC_029978.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8944	2,31E+08	100%	0.0	99.94%	261510930	NC_029979.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8944	1,52E+08	100%	0.0	99.94%	219521584	NC_029982.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8909	3,04E+08	100%	0.0	99.81%	301019445	NC_029977.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8907	1,70E+08	100%	0.0	99.81%	229934170	NC_029988.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8901	8,83E+04	100%	0.0	99.79%	153299543	NC_029984.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8896	1,90E+08	100%	0.0	99.81%	215701946	NC_029980.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8887	1,55E+08	100%	0.0	99.73%	238794889	NC_029985.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4537	<i>Capsicum annuum</i>	8879	9,41E+03	100%	0.0	99.71%	176915	NW_015961775.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold891	<i>Capsicum annuum</i>	8874	1,57E+04	100%	0.0	99.69%	1431046	NW_015960469.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8872	1,63E+08	100%	0.0	99.69%	222112641	NC_029983.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8848	1,36E+08	100%	0.0	99.57%	217274494	NC_029981.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8796	1,25E+08	100%	0.0	99.40%	220335243	NC_029987.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8778	1,81E+08	100%	0.0	99.30%	205736368	NC_029986.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold367	<i>Capsicum annuum</i>	8386	1,02E+04	97%	0.0	99.98%	625659	NW_015960847.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold7798	<i>Capsicum annuum</i>	8379	9,49E+03	100%	0.0	99.61%	106903	NW_015961464.1

monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2635	<i>Capsicum annuum</i>	8362	1,78E+04	97%	0.0	99.87%	699991	NW_015960547.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1191	<i>Capsicum annuum</i>	8347	9,45E+03	98%	0.0	99.82%	145056	NW_015961591.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold448	<i>Capsicum annuum</i>	8342	9,31E+03	98%	0.0	99.80%	1323857	NW_015960958.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold90	<i>Capsicum annuum</i>	8335	9,99E+03	98%	0.0	99.78%	6074945	NW_015960458.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1246	<i>Capsicum annuum</i>	8333	2,73E+04	98%	0.0	99.76%	3388803	NW_015960834.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2615	<i>Capsicum annuum</i>	8314	9,27E+03	98%	0.0	99.71%	131126	NW_015961557.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold181	<i>Capsicum annuum</i>	8279	9,20E+03	98%	0.0	99.78%	2703691	NW_015961388.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold301	<i>Capsicum annuum</i>	8277	9,28E+03	98%	0.0	99.73%	1226190	NW_015960697.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold519	<i>Capsicum annuum</i>	8264	9,22E+03	97%	0.0	99.80%	1101621	NW_015961578.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1037	<i>Capsicum annuum</i>	8263	9,15E+03	98%	0.0	99.71%	2186250	NW_015960681.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2298	<i>Capsicum annuum</i>	8242	9,23E+03	97%	0.0	99.71%	1494726	NW_015960454.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capsicum annuum</i>	8209	9,13E+03	97%	0.0	99.80%	2397553	NW_015961318.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	8189	9,13E+03	97%	0.0	99.87%	2470984	NW_015961229.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2719	<i>Capsicum annuum</i>	8176	9,09E+03	99%	0.0	99.60%	277434	NW_015960900.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1874	<i>Capsicum annuum</i>	8139	9,02E+03	96%	0.0	99.62%	719064	NW_015961147.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1360	<i>Capsicum annuum</i>	7958	8,85E+03	94%	0.0	99.66%	360446	NW_015960954.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1399	<i>Capsicum annuum</i>	7893	9,46E+03	98%	0.0	99.81%	1742971	NW_015961630.1
monotypic-Ch02d_66	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2138	<i>Capsicum annuum</i>	6890	7,16E+03	80%	0.0	99.89%	681955	NW_015960370.1
monotypic-Ch02d_70	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8700	1,67E+04	100%	0.0	100.00%	163962470	NC_029978.1
monotypic-Ch02d_70	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7269	7,74E+03	100%	0.0	94.52%	238794889	NC_029985.1
monotypic-Ch02d_70	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7097	7,64E+03	100%	0.0	93.89%	215701946	NC_029980.1
monotypic-Ch02d_70	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7075	1,77E+04	100%	0.0	93.84%	229934170	NC_029988.1
monotypic-Ch02d_70	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6868	1,12E+04	100%	0.0	92.97%	220335243	NC_029987.1
monotypic-Ch02d_70	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6811	1,50E+04	100%	0.0	92.75%	219521584	NC_029982.1
monotypic-Ch02d_70	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold258	<i>Capsicum annuum</i>	6373	6,79E+03	100%	0.0	91.12%	3098259	NW_015961180.1
monotypic-Ch02d_70	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5502	5,86E+03	97%	0.0	89.21%	217274494	NC_029981.1
monotypic-Ch02d_70	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2348	<i>Capsicum annuum</i>	2636	6,25E+03	100%	0.0	89.85%	373378	NW_015960596.1
monotypic-Ch02d_70	refeq-genome	8	Nicotiana attenuata strain Ut chromosome 12, NIATTr2	<i>Nicotiana attenuata</i>	2237	3,70E+03	90%	0.0	82.12%	53491057	NC_032000.1
monotypic-Ch02d_70	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9192	<i>Nicotiana tabacum</i>	2145	3,40E+03	91%	0.0	81.52%	31423	NW_015949058.1
monotypic-Ch02d_70	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold7467	<i>Nicotiana sylvestris</i>	2130	3,42E+03	91%	0.0	81.40%	59109	NW_009564575.1
monotypic-Ch02d_70	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4210	<i>Nicotiana tabacum</i>	2102	3,57E+03	88%	0.0	81.62%	190891	NW_015905470.1
monotypic-Ch02d_70	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2856	<i>Nicotiana tomentosiformis</i>	2102	3,58E+03	88%	0.0	81.62%	378280	NW_008908663.1
monotypic-Ch02d_70	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold51410	<i>Nicotiana tabacum</i>	1971	3,23E+03	90%	0.0	80.67%	117067	NW_015914026.1
monotypic-Ch02d_70	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold389	<i>Nicotiana tomentosiformis</i>	1949	3,24E+03	90%	0.0	80.51%	82980	NW_008920151.1
monotypic-Ch02d_70	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold12917	<i>Nicotiana tabacum</i>	1936	3,27E+03	91%	0.0	80.26%	107750	NW_015808904.1
monotypic-Ch02d_70	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold15092	<i>Nicotiana tabacum</i>	1116	3,05E+03	90%	0.0	79.64%	108898	NW_015823451.1
monotypic-Ch02d_71	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10009	1,10E+04	100%	0.0	100.00%	163962470	NC_029978.1
monotypic-Ch02d_71	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4863	4,94E+03	89%	0.0	84.97%	215701946	NC_029980.1
monotypic-Ch02d_71	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4820	5,97E+03	84%	0.0	85.77%	220335243	NC_029987.1
monotypic-Ch02d_71	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4547	5,03E+03	100%	0.0	83.98%	238794889	NC_029985.1
monotypic-Ch02d_71	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3831	6,52E+03	99%	0.0	85.24%	229934170	NC_029988.1
monotypic-Ch02d_71	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3807	8,36E+03	100%	0.0	89.70%	205736368	NC_029986.1
monotypic-Ch02d_71	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3476	4,80E+03	90%	0.0	80.31%	222112641	NC_029983.1
monotypic-Ch02d_71	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	1855	3,81E+03	83%	0.0	83.64%	219521584	NC_029982.1
monotypic-Ch02e_08	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9060	1,12E+04	100%	0.0	99.96%	163962470	NC_029978.1
monotypic-Ch02e_08	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4397	8,14E+03	95%	0.0	93.76%	220335243	NC_029987.1
monotypic-Ch02e_08	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4165	1,20E+04	95%	0.0	92.34%	301019445	NC_029977.1
monotypic-Ch02e_08	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4036	7,27E+03	94%	0.0	91.82%	238794889	NC_029985.1
monotypic-Ch02e_08	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2671	6,68E+03	95%	0.0	90.95%	215701946	NC_029980.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10275	3,84E+08	100%	0.0	100.00%	163962470	NC_029978.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5445	3,79E+08	100%	0.0	92.20%	229934170	NC_029988.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5433	3,13E+08	100%	0.0	91.46%	222112641	NC_029983.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5369	1,92E+08	93%	0.0	91.24%	153299543	NC_029984.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5341	3,24E+08	94%	0.0	92.27%	215701946	NC_029980.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5321	3,46E+08	94%	0.0	91.96%	220335243	NC_029987.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5306	3,61E+08	94%	0.0	90.95%	238794889	NC_029985.1

monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold8268	<i>Capsicum annuum</i>	5301	6,55E+03	90%	0.0	91.15%	176351	NW_015960588.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5280	2,72E+08	93%	0.0	90.93%	217274494	NC_029981.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capsicum annuum</i>	5275	1,41E+04	92%	0.0	90.83%	2397553	NW_015961318.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5264	3,54E+08	94%	0.0	91.07%	205736368	NC_029982.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5245	5,48E+08	100%	0.0	90.77%	261510930	NC_029979.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1341	<i>Capsicum annuum</i>	5238	8,13E+03	88%	0.0	90.85%	1906221	NW_015960665.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5210	4,95E+08	94%	0.0	90.70%	301019445	NC_029977.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5195	3,28E+08	100%	0.0	90.45%	219521584	NC_029982.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2719	<i>Capsicum annuum</i>	5190	8,91E+03	90%	0.0	90.47%	277434	NW_015960900.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold277	<i>Capsicum annuum</i>	5101	7,47E+03	89%	0.0	90.44%	448122	NW_015961390.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold175	<i>Capsicum annuum</i>	5040	6,80E+03	87%	0.0	90.66%	655560	NW_015960845.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	4843	1,51E+04	90%	0.0	89.63%	3493078	NW_015960517.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold937	<i>Capsicum annuum</i>	4835	7,50E+03	89%	0.0	89.15%	1465884	NW_015961782.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold814	<i>Capsicum annuum</i>	4824	1,18E+04	88%	0.0	89.38%	1128876	NW_015960740.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold306	<i>Capsicum annuum</i>	4796	6,03E+03	87%	0.0	89.60%	749978	NW_015961154.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5785	<i>Capsicum annuum</i>	4771	6,86E+03	90%	0.0	90.84%	79892	NW_015961741.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3026	<i>Capsicum annuum</i>	4761	8,76E+03	87%	0.0	89.46%	1386929	NW_015961102.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2523	<i>Capsicum annuum</i>	4747	6,72E+03	85%	0.0	89.65%	735434	NW_015960464.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold231	<i>Capsicum annuum</i>	4652	6,02E+03	86%	0.0	88.29%	1482909	NW_015961568.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold170	<i>Capsicum annuum</i>	4578	5,87E+03	88%	0.0	87.97%	2217302	NW_015960390.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1670	<i>Capsicum annuum</i>	4401	6,15E+03	89%	0.0	87.10%	1202455	NW_015961465.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1888	<i>Capsicum annuum</i>	3976	7,41E+03	86%	0.0	90.98%	361877	NW_015960979.1
monotypic-Ch02e_17	refeq-genome	8	Solanum pennellii chromosome 5, SPENN200	<i>Solanum pennellii</i>	3904	4,79E+04	82%	0.0	86.00%	77991103	NC_028641.1
monotypic-Ch02e_17	refeq-genome	8	Solanum pennellii chromosome 12, SPENN200	<i>Solanum pennellii</i>	3901	3,62E+04	82%	0.0	86.02%	83305730	NC_028648.1
monotypic-Ch02e_17	refeq-genome	8	Solanum pennellii chromosome 1, SPENN200	<i>Solanum pennellii</i>	3884	5,33E+04	83%	0.0	85.94%	109333515	NC_028637.1
monotypic-Ch02e_17	refeq-genome	8	Solanum pennellii chromosome 3, SPENN200	<i>Solanum pennellii</i>	3884	5,24E+04	82%	0.0	85.94%	75414019	NC_028639.1
monotypic-Ch02e_17	refeq-genome	8	Solanum pennellii chromosome 9, SPENN200	<i>Solanum pennellii</i>	3884	4,43E+04	82%	0.0	85.94%	84057508	NC_028645.1
monotypic-Ch02e_17	refeq-genome	8	Solanum pennellii chromosome 11, SPENN200	<i>Solanum pennellii</i>	3873	4,30E+04	82%	0.0	85.88%	66223686	NC_028647.1
monotypic-Ch02e_17	refeq-genome	8	Solanum pennellii chromosome 4, SPENN200	<i>Solanum pennellii</i>	3856	3,65E+04	82%	0.0	85.80%	77197300	NC_028640.1
monotypic-Ch02e_17	refeq-genome	8	Solanum pennellii chromosome 8, SPENN200	<i>Solanum pennellii</i>	3856	6,42E+04	82%	0.0	85.80%	70546659	NC_028644.1
monotypic-Ch02e_17	refeq-genome	8	Solanum pennellii chromosome 6, SPENN200	<i>Solanum pennellii</i>	3851	4,70E+04	82%	0.0	85.77%	60730942	NC_028642.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold8073	<i>Capsicum annuum</i>	3851	5,53E+03	84%	0.0	87.45%	91573	NW_015961746.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1165	<i>Capsicum annuum</i>	3847	6,44E+03	89%	0.0	89.81%	67805	NW_015960689.1
monotypic-Ch02e_17	refeq-genome	8	Solanum pennellii chromosome 2, SPENN200	<i>Solanum pennellii</i>	3808	3,64E+04	81%	0.0	85.56%	59803892	NC_028638.1
monotypic-Ch02e_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	<i>Solanum lycopersicum</i>	3709	3,85E+04	82%	0.0	85.12%	68175699	NC_015444.3
monotypic-Ch02e_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	<i>Solanum lycopersicum</i>	3707	4,29E+04	82%	0.0	85.08%	72906345	NC_015446.3
monotypic-Ch02e_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	<i>Solanum lycopersicum</i>	3683	5,14E+04	82%	0.0	84.96%	56597135	NC_015448.3
monotypic-Ch02e_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	<i>Solanum lycopersicum</i>	3668	3,06E+04	80%	0.0	84.84%	49794276	NC_015443.3
monotypic-Ch02e_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	<i>Solanum lycopersicum</i>	3629	3,19E+04	82%	0.0	84.71%	66723567	NC_015442.3
monotypic-Ch02e_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	<i>Solanum lycopersicum</i>	3581	3,71E+04	82%	0.0	84.47%	72290146	NC_015440.3
monotypic-Ch02e_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	<i>Solanum lycopersicum</i>	3574	3,37E+04	82%	0.0	84.48%	66557038	NC_015441.3
monotypic-Ch02e_17	refeq-genome	8	Solanum pennellii chromosome 10, SPENN200	<i>Solanum pennellii</i>	3531	3,41E+04	82%	0.0	85.26%	82529941	NC_028646.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold258	<i>Capsicum annuum</i>	3531	9,82E+03	85%	0.0	91.83%	3098259	NW_015961180.1
monotypic-Ch02e_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	<i>Solanum lycopersicum</i>	3507	3,95E+04	81%	0.0	84.12%	55977580	NC_015439.3
monotypic-Ch02e_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	<i>Solanum lycopersicum</i>	3465	4,36E+04	82%	0.0	83.96%	65633393	NC_015447.3
monotypic-Ch02e_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	<i>Solanum lycopersicum</i>	3463	2,89E+04	81%	0.0	83.83%	98455869	NC_015438.3
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2102	<i>Capsicum annuum</i>	3428	6,67E+03	89%	0.0	90.50%	627640	NW_015961440.1
monotypic-Ch02e_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	<i>Solanum lycopersicum</i>	3421	3,88E+04	82%	0.0	83.88%	65987440	NC_015445.3
monotypic-Ch02e_17	refeq-genome	8	Solanum pennellii chromosome 7, SPENN200	<i>Solanum pennellii</i>	3384	3,77E+04	83%	0.0	85.88%	79292169	NC_028643.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1336	<i>Capsicum annuum</i>	3363	6,68E+03	89%	0.0	90.03%	463180	NW_015961656.1
monotypic-Ch02e_17	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	<i>Solanum lycopersicum</i>	3354	2,30E+04	81%	0.0	83.35%	68126176	NC_015449.3
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2138	<i>Capsicum annuum</i>	3153	7,35E+03	87%	0.0	90.64%	681955	NW_015960370.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3348	<i>Capsicum annuum</i>	3011	6,39E+03	90%	0.0	90.11%	31107	NW_015961241.1
monotypic-Ch02e_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2400	<i>Capsicum annuum</i>	2953	1,34E+04	89%	0.0	90.76%	1353284	NW_015960968.1
monotypic-Ch02e_68	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8637	4,64E+08	100%	0.0	100.00%	163962470	NC_029978.1
monotypic-Ch02e_68	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5264	2,06E+08	100%	0.0	92.71%	222112641	NC_029983.1

monotypic-Ch02e_68	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3760	7,15E+08	100%	0.0	86.44%	229934170	NC_029988.1
monotypic-Ch02e_68	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3615	4,34E+08	99%	0.0	85.52%	215701946	NC_029980.1
monotypic-Ch02e_68	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3542	2,99E+08	84%	0.0	86.24%	301019445	NC_029977.1
monotypic-Ch02e_68	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3459	6,18E+08	97%	0.0	84.77%	261510930	NC_029979.1
monotypic-Ch02e_68	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2763	2,59E+08	89%	0.0	86.48%	153299543	NC_029984.1
monotypic-Ch02e_68	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2741	6,73E+08	100%	0.0	99.80%	238794889	NC_029985.1
monotypic-Ch02e_68	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2736	3,68E+08	96%	0.0	99.73%	205736368	NC_029986.1
monotypic-Ch02e_68	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2730	2,53E+08	99%	0.0	99.67%	220335243	NC_029987.1
monotypic-Ch03a_32	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10013	1,46E+04	100%	0.0	100.00%	261510930	NC_029979.1
monotypic-Ch03a_32	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6828	9,20E+03	100%	0.0	94.94%	220335243	NC_029987.1
monotypic-Ch03a_32	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6802	9,48E+03	100%	0.0	94.82%	301019445	NC_029977.1
monotypic-Ch03a_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9463	1,50E+04	100%	0.0	100.00%	261510930	NC_029979.1
monotypic-Ch03a_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5531	5,78E+03	99%	0.0	86.98%	222112641	NC_029983.1
monotypic-Ch03a_77	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9659	1,05E+04	100%	0.0	100.00%	261510930	NC_029979.1
monotypic-Ch03a_77	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2968	5,39E+03	100%	0.0	82.95%	219521584	NC_029982.1
monotypic-Ch03b_38	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10100	1,13E+04	100%	0.0	99.98%	261510930	NC_029979.1
monotypic-Ch03b_38	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5110	1,34E+04	100%	0.0	90.10%	205736368	NC_029986.1
monotypic-Ch03b_38	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2933	5,90E+03	90%	0.0	87.41%	238794889	NC_029985.1
monotypic-Ch03g_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9468	1,05E+04	100%	0.0	100.00%	261510930	NC_029979.1
monotypic-Ch03g_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7986	8,93E+03	100%	0.0	94.73%	163962470	NC_029978.1
monotypic-Ch03g_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3877	8,30E+03	96%	0.0	92.78%	217274494	NC_029981.1
monotypic-Ch03g_76	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9095	1,87E+04	100%	0.0	100.00%	261510930	NC_029979.1
monotypic-Ch03g_76	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7463	2,95E+04	99%	0.0	94.38%	220335243	NC_029987.1
monotypic-Ch03g_76	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7289	2,62E+04	99%	0.0	93.96%	301019445	NC_029977.1
monotypic-Ch03g_76	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7260	2,08E+04	99%	0.0	93.32%	215701946	NC_029980.1
monotypic-Ch03g_76	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7057	3,47E+04	99%	0.0	92.58%	238794889	NC_029985.1
monotypic-Ch03g_76	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7042	3,23E+04	99%	0.0	93.22%	205736368	NC_029986.1
monotypic-Ch03g_76	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold433	<i>Capsicum annuum</i>	5060	6,68E+03	97%	0.0	90.94%	1277097	NW_015961081.1
monotypic-Ch03g_76	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4885	1,16E+04	99%	0.0	87.36%	217274494	NC_029981.1
monotypic-Ch03g_76	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4185	2,72E+04	99%	0.0	91.51%	222112641	NC_029983.1
monotypic-Ch03g_76	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTTr2	<i>Nicotiana attenuata</i>	4054	4,05E+03	85%	0.0	84.17%	966234	NW_017670333.1
monotypic-Ch03g_76	refeq-genome	8	Nicotiana attenuata strain UT chromosome 11, NIATTTr2	<i>Nicotiana attenuata</i>	4004	4,00E+03	85%	0.0	83.97%	61252062	NC_031999.1
monotypic-Ch03g_76	refeq-genome	8	Solanum pennellii chromosome 10, SPENNv200	<i>Solanum pennellii</i>	3803	7,99E+03	85%	0.0	83.07%	82529941	NC_028646.1
monotypic-Ch03g_76	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00278	<i>Solanum tuberosum</i>	3670	5,00E+03	89%	0.0	81.94%	824609	NW_006239215.1
monotypic-Ch03g_76	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00200	<i>Solanum tuberosum</i>	2671	4,27E+03	89%	0.0	85.25%	1164273	NW_006239124.1
monotypic-Ch03g_76	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold50102	<i>Nicotiana tabacum</i>	1949	3,41E+03	82%	0.0	82.92%	74331	NW_015912835.1
monotypic-Ch04a_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9598	1,05E+04	100%	0.0	100.00%	215701946	NC_029980.1
monotypic-Ch04a_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6082	8,61E+03	82%	0.0	95.19%	229934170	NC_029988.1
monotypic-Ch04a_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold148	<i>Capsicum annuum</i>	4745	8,65E+03	98%	0.0	95.38%	1323090	NW_015961290.1
monotypic-Ch04a_25	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3376	7,28E+03	99%	0.0	90.42%	261510930	NC_029979.1
monotypic-Ch04d_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8074	1,30E+04	100%	0.0	100.00%	215701946	NC_029980.1
monotypic-Ch04d_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3162	5,47E+03	99%	0.0	87.38%	220335243	NC_029987.1
monotypic-Ch04d_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2423	1,16E+04	98%	0.0	89.81%	261510930	NC_029979.1
monotypic-Ch04d_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold6237	<i>Capsicum annuum</i>	2368	5,49E+03	98%	0.0	88.23%	101780	NW_015960658.1
monotypic-Ch04d_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2350	5,07E+03	98%	0.0	89.12%	205736368	NC_029986.1
monotypic-Ch04d_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold102420	<i>Nicotiana tabacum</i>	2167	3,77E+03	94%	0.0	82.88%	39892	NW_015789122.1
monotypic-Ch04d_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold14165	<i>Nicotiana tomentosiformis</i>	2145	3,75E+03	94%	0.0	82.72%	110726	NW_008874769.1
monotypic-Ch04d_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	1934	9,78E+03	99%	0.0	87.90%	153299543	NC_029984.1
monotypic-Ch04d_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold86662	<i>Nicotiana tomentosiformis</i>	1807	4,25E+03	94%	0.0	85.96%	8422	NW_008973221.1
monotypic-Ch04d_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold20485	<i>Nicotiana tabacum</i>	1779	4,20E+03	94%	0.0	85.66%	159734	NW_015854099.1
monotypic-Ch04d_23	refeq-genome	8	Nicotiana attenuata strain UT chromosome 1, NIATTTr2	<i>Nicotiana attenuata</i>	1773	4,10E+03	92%	0.0	85.61%	110538649	NC_031989.1
monotypic-Ch04d_23	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold131556	<i>Nicotiana tabacum</i>	1725	3,76E+03	85%	0.0	85.09%	53056	NW_015810592.1
monotypic-Ch04d_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	1687	5,02E+03	85%	0.0	87.05%	238794889	NC_029985.1
monotypic-Ch04d_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTTr2	<i>Nicotiana attenuata</i>	1642	3,96E+03	94%	0.0	84.37%	176171	NW_017672684.1
monotypic-Ch04d_23	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5305	<i>Nicotiana tomentosiformis</i>	1570	3,60E+03	86%	0.0	83.49%	128670	NW_008935875.1
monotypic-Ch04d_23	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTTr2	<i>Nicotiana attenuata</i>	1568	3,93E+03	90%	0.0	83.99%	1295229	NW_017670282.1

monotypic-Ch05a_52	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8541	5,09E+04	100%	0.0	100.00%	217274494	NC_029981.1
monotypic-Ch05a_52	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1653	<i>Capsicum annuum</i>	3864	6,23E+03	82%	0.0	94.26%	913988	NW_015961387.1
monotypic-Ch05a_52	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3526	4,48E+04	94%	0.0	93.15%	205736368	NC_029986.1
monotypic-Ch05a_52	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3653	<i>Capsicum annuum</i>	3493	5,43E+03	82%	0.0	91.71%	172086	NW_015961357.1
monotypic-Ch05a_52	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3448	4,06E+04	94%	0.0	91.43%	219521584	NC_029982.1
monotypic-Ch05a_52	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3417	4,28E+04	93%	0.0	92.15%	222112641	NC_029983.1
monotypic-Ch05a_52	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3347	6,69E+04	94%	0.0	90.65%	301019445	NC_029977.1
monotypic-Ch05a_52	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3308	2,78E+04	96%	0.0	91.42%	229934170	NC_029988.1
monotypic-Ch05a_52	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3212	3,81E+04	96%	0.0	91.33%	261510930	NC_029979.1
monotypic-Ch05a_52	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1399	<i>Capsicum annuum</i>	3201	8,84E+03	89%	0.0	90.75%	1742971	NW_015961630.1
monotypic-Ch05a_52	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2375	<i>Capsicum annuum</i>	3199	5,24E+03	80%	0.0	90.49%	1171512	NW_015961637.1
monotypic-Ch05a_52	refeq-genome	8	Solanum pennellii chromosome 9, SPENN200	<i>Solanum pennellii</i>	3151	8,42E+03	84%	0.0	89.18%	84057508	NC_028645.1
monotypic-Ch05a_52	refeq-genome	8	Solanum pennellii chromosome 4, SPENN200	<i>Solanum pennellii</i>	3094	2,65E+04	85%	0.0	88.79%	77197300	NC_028640.1
monotypic-Ch05a_52	refeq-genome	8	Solanum pennellii chromosome 5, SPENN200	<i>Solanum pennellii</i>	3049	8,34E+03	84%	0.0	88.47%	77991103	NC_028641.1
monotypic-Ch05a_52	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	<i>Solanum lycopersicum</i>	3040	9,98E+03	84%	0.0	88.40%	66723567	NC_015442.3
monotypic-Ch05a_52	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	<i>Solanum lycopersicum</i>	3018	2,93E+04	84%	0.0	88.26%	66555708	NC_015441.3
monotypic-Ch05a_52	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00039	<i>Solanum tuberosum</i>	2994	4,85E+03	81%	0.0	89.24%	2527457	NW_006238968.1
monotypic-Ch05a_52	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00008	<i>Solanum tuberosum</i>	2985	4,77E+03	80%	0.0	89.18%	4063183	NW_006238934.1
monotypic-Ch05a_52	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	<i>Solanum lycopersicum</i>	2981	1,48E+04	84%	0.0	88.00%	98455869	NC_015438.3
monotypic-Ch05a_52	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	<i>Solanum lycopersicum</i>	2924	5,19E+03	84%	0.0	87.63%	68126176	NC_015449.3
monotypic-Ch05a_52	refeq-genome	8	Solanum pennellii chromosome 12, SPENN200	<i>Solanum pennellii</i>	2904	1,37E+04	84%	0.0	87.57%	83305730	NC_028648.1
monotypic-Ch05a_52	refeq-genome	8	Nicotiana attenuata strain UT chromosome 9, NIATT2	<i>Nicotiana attenuata</i>	2800	4,76E+03	81%	0.0	87.20%	50975570	NC_031997.1
monotypic-Ch05a_52	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	2771	4,48E+03	81%	0.0	86.99%	217169	NW_017672119.1
monotypic-Ch05a_52	refeq-genome	8	Nicotiana attenuata strain UT chromosome 5, NIATT2	<i>Nicotiana attenuata</i>	2747	4,50E+03	81%	0.0	86.83%	57012777	NC_031993.1
monotypic-Ch05a_52	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold9133	<i>Nicotiana sylvestris</i>	2741	4,44E+03	81%	0.0	86.80%	69551	NW_009583087.1
monotypic-Ch05a_52	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold229741	<i>Nicotiana tabacum</i>	2678	4,36E+03	81%	0.0	86.33%	44276	NW_015866064.1
monotypic-Ch05a_52	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold30058	<i>Nicotiana sylvestris</i>	2582	4,20E+03	81%	0.0	85.75%	59914	NW_009515008.1
monotypic-Ch05a_52	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold40764	<i>Nicotiana tabacum</i>	2571	4,19E+03	81%	0.0	85.67%	9400	NW_015904227.1
monotypic-Ch05a_52	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00123	<i>Solanum tuberosum</i>	2564	4,29E+03	80%	0.0	86.06%	1602102	NW_006239048.1
monotypic-Ch05a_52	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1107	<i>Capsicum annuum</i>	1864	5,68E+03	82%	0.0	91.89%	318496	NW_015961239.1
monotypic-Ch05a_59	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8495	4,89E+04	100%	0.0	100.00%	217274494	NC_029981.1
monotypic-Ch05a_59	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5273	4,92E+04	94%	0.0	91.96%	222112641	NC_029983.1
monotypic-Ch05a_59	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5097	5,40E+04	94%	0.0	91.25%	301019445	NC_029977.1
monotypic-Ch05a_59	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5042	4,70E+04	94%	0.0	90.88%	219521584	NC_029982.1
monotypic-Ch05a_59	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5031	5,48E+04	100%	0.0	90.98%	215701946	NC_029980.1
monotypic-Ch05a_59	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4990	1,60E+04	93%	0.0	90.63%	153299543	NC_029984.1
monotypic-Ch05a_59	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1339	<i>Capsicum annuum</i>	4979	9,08E+03	84%	0.0	90.66%	928404	NW_015960451.1
monotypic-Ch05a_59	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3819	3,76E+04	93%	0.0	91.68%	238794889	NC_029985.1
monotypic-Ch05a_59	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2828	4,28E+04	93%	0.0	86.79%	229934170	NC_029988.1
monotypic-Ch05a_59	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2686	3,46E+04	100%	0.0	81.46%	205736368	NC_029986.1
monotypic-Ch05a_59	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2289	3,70E+04	92%	0.0	77.91%	163962470	NC_029978.1
monotypic-Ch05f_50	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8466	2,40E+04	100%	0.0	100.00%	217274494	NC_029981.1
monotypic-Ch05f_50	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4290	2,40E+04	99%	0.0	93.78%	222112641	NC_029983.1
monotypic-Ch05f_50	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4200	8,94E+03	99%	0.0	93.28%	261510930	NC_029979.1
monotypic-Ch05f_50	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3923	1,35E+04	99%	0.0	91.76%	220335243	NC_029987.1
monotypic-Ch05f_50	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3825	1,07E+04	99%	0.0	91.02%	215701946	NC_029980.1
monotypic-Ch05f_50	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold175	<i>Capsicum annuum</i>	3757	6,42E+03	98%	0.0	90.62%	655560	NW_015960451.1
monotypic-Ch05f_50	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3747	1,10E+04	99%	0.0	90.54%	219521584	NC_029982.1
monotypic-Ch05f_50	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3411	1,58E+04	99%	0.0	92.74%	301019445	NC_029977.1
monotypic-Ch05f_50	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2946	5,55E+03	83%	0.0	92.89%	205736368	NC_029986.1
monotypic-Ch05f_50	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1339	<i>Capsicum annuum</i>	2808	7,13E+03	99%	0.0	93.50%	928404	NW_015960451.1
monotypic-Ch05f_50	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2639	8,16E+03	87%	0.0	85.62%	229934170	NC_029988.1
monotypic-Ch06a_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10056	1,11E+04	100%	0.0	100.00%	219521584	NC_029982.1
monotypic-Ch06a_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1795	<i>Capsicum annuum</i>	4126	8,57E+03	99%	0.0	91.99%	162617	NW_015961207.1
monotypic-Ch06a_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3799	7,98E+03	99%	0.0	90.16%	217274494	NC_029981.1
monotypic-Ch06a_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3770	2,47E+04	99%	0.0	89.81%	163962470	NC_029978.1

monotypic-Ch06a_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3513	9,57E+03	98%	0.0	91.19%	261510930	NC_029979.1
monotypic-Ch06a_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3083	8,22E+03	91%	0.0	92.38%	153299543	NC_029984.1
monotypic-Ch06a_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2577	5,43E+03	82%	0.0	87.67%	229934170	NC_029988.1
monotypic-Ch06a_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1422	<i>Capsicum annuum</i>	2174	1,17E+04	94%	0.0	91.28%	802096	NW_015960747.1
monotypic-Ch06a_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2091	8,92E+03	99%	0.0	92.41%	220335243	NC_029987.1
monotypic-Ch06a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9278	2,29E+04	100%	0.0	100.00%	219521584	NC_029982.1
monotypic-Ch06a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7057	3,11E+04	100%	0.0	92.89%	261510930	NC_029979.1
monotypic-Ch06a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold12872	<i>Capsicum annuum</i>	6935	7,59E+03	100%	0.0	91.71%	51304	NW_015961659.1
monotypic-Ch06a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6898	1,46E+04	100%	0.0	91.54%	215701946	NC_029980.1
monotypic-Ch06a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold814	<i>Capsicum annuum</i>	6852	7,42E+03	100%	0.0	91.37%	1128876	NW_015960740.1
monotypic-Ch06a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold592	<i>Capsicum annuum</i>	5999	6,00E+03	89%	0.0	90.86%	1905997	NW_015960988.1
monotypic-Ch06a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5720	1,16E+04	100%	0.0	90.76%	222112641	NC_029983.1
monotypic-Ch06a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4767	2,08E+04	99%	0.0	84.69%	205736368	NC_029986.1
monotypic-Ch06a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4263	1,83E+04	96%	0.0	84.90%	217274494	NC_029981.1
monotypic-Ch06a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3463	1,40E+04	99%	0.0	79.94%	163962470	NC_029978.1
monotypic-Ch06a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3326	1,17E+04	94%	0.0	88.35%	301019445	NC_029977.1
monotypic-Ch06a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2891	6,48E+03	100%	0.0	82.57%	153299543	NC_029984.1
monotypic-Ch06a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1929	<i>Capsicum annuum</i>	2615	3,41E+03	90%	0.0	83.75%	996492	NW_015961795.1
monotypic-Ch06a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1874	<i>Capsicum annuum</i>	2551	2,55E+03	93%	0.0	76.79%	719064	NW_015961147.1
monotypic-Ch06a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2145	1,17E+04	96%	0.0	84.97%	238794889	NC_029985.1
monotypic-Ch06a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	1893	4,08E+03	89%	0.0	85.43%	220335243	NC_029987.1
monotypic-Ch06a_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9383	1,03E+04	100%	0.0	100.00%	219521584	NC_029982.1
monotypic-Ch06a_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3199	9,25E+03	100%	0.0	93.58%	301019445	NC_029977.1
monotypic-Ch06a_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3169	8,59E+03	100%	0.0	93.29%	205736368	NC_029986.1
monotypic-Ch06a_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3120	1,37E+04	100%	0.0	92.93%	222112641	NC_029983.1
monotypic-Ch06a_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3088	1,62E+04	100%	0.0	92.46%	261510930	NC_029979.1
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10510	7,13E+04	100%	0.0	100.00%	219521584	NC_029982.1
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7382	8,30E+04	99%	0.0	90.26%	229934170	NC_029988.1
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7138	1,40E+08	99%	0.0	89.65%	261510930	NC_029979.1
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6691	5,88E+04	99%	0.0	88.48%	222112641	NC_029983.1
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6396	1,13E+08	99%	0.0	89.73%	301019445	NC_029977.1
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	6150	7,62E+03	98%	0.0	89.46%	3493078	NW_015960517.1
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5997	7,27E+04	99%	0.0	89.29%	220335243	NC_029987.1
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5642	5,01E+04	94%	0.0	86.97%	215701946	NC_029980.1
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5566	6,39E+04	97%	0.0	87.33%	205736368	NC_029986.1
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5315	9,11E+04	99%	0.0	87.77%	217274494	NC_029981.1
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5188	7,49E+04	97%	0.0	87.73%	163962470	NC_029978.1
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5018	1,41E+04	99%	0.0	88.05%	153299543	NC_029984.1
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4706	7,91E+04	99%	0.0	88.51%	238794889	NC_029985.1
monotypic-Ch06a_26	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00013	<i>Solanum tuberosum</i>	4634	4,63E+03	85%	0.0	83.94%	3422333	NW_006238940.1
monotypic-Ch06a_26	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00120	<i>Solanum tuberosum</i>	4359	5,98E+03	86%	0.0	82.93%	1603502	NW_006239047.1
monotypic-Ch06a_26	refeq-genome	8	Solanum pennellii chromosome 7, SPENN200	<i>Solanum pennellii</i>	4224	4,58E+04	86%	0.0	82.59%	79292169	NC_028643.1
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold231	<i>Capsicum annuum</i>	4191	7,56E+03	92%	0.0	83.58%	1482909	NW_015961568.1
monotypic-Ch06a_26	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00256	<i>Solanum tuberosum</i>	4102	4,10E+03	85%	0.0	82.05%	954438	NW_006239178.1
monotypic-Ch06a_26	refeq-genome	8	Solanum pennellii chromosome 10, SPENN200	<i>Solanum pennellii</i>	4085	5,11E+04	86%	0.0	82.38%	82529941	NC_028646.1
monotypic-Ch06a_26	refeq-genome	8	Solanum pennellii chromosome 3, SPENN200	<i>Solanum pennellii</i>	4060	4,08E+04	86%	0.0	81.92%	75414019	NC_028639.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	4024	4,02E+03	85%	0.0	81.71%	55101	NW_017675972.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT chromosome 6, NIATT2	<i>Nicotiana attenuata</i>	4012	2,32E+04	86%	0.0	81.66%	97571839	NC_031994.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	4006	4,01E+03	85%	0.0	81.63%	2491754	NW_017670241.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	4002	4,00E+03	85%	0.0	81.61%	167244	NW_017672821.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	3960	3,96E+03	85%	0.0	81.47%	310446	NW_017671335.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT chromosome 8, NIATT2	<i>Nicotiana attenuata</i>	3958	1,08E+04	86%	0.0	81.48%	47755514	NC_031996.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	3954	3,95E+03	85%	0.0	81.52%	217963	NW_017672111.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	3949	6,48E+03	85%	0.0	81.50%	442368	NW_017670825.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	<i>Nicotiana attenuata</i>	3938	3,94E+03	86%	0.0	81.35%	934905	NW_017670343.1
monotypic-Ch06a_26	refeq-genome	8	Solanum pennellii chromosome 12, SPENN200	<i>Solanum pennellii</i>	3934	3,81E+04	85%	0.0	81.45%	83305730	NC_028648.1

monotypic-Ch06a_26	refeq-genome	8	Solanum pennellii chromosome 11, SPENN200	<i>Solanum pennellii</i>	3928	4,61E+04	86%	0.0	81.45%	66223686	NC_028647.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3923	3,92E+03	85%	0.0	81.31%	1217564	NW_017670290.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3919	6,10E+03	85%	0.0	81.38%	664384	NW_017670472.1
monotypic-Ch06a_26	refeq-genome	8	Solanum pennellii chromosome 1, SPENN200	<i>Solanum pennellii</i>	3917	7,15E+04	86%	0.0	81.42%	109333515	NC_028637.1
monotypic-Ch06a_26	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	<i>Solanum lycopersicum</i>	3914	4,66E+04	86%	0.0	81.30%	56597135	NC_015448.3
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3904	3,90E+03	85%	0.0	81.27%	467702	NW_017670748.1
monotypic-Ch06a_26	refeq-genome	8	Solanum pennellii chromosome 5, SPENN200	<i>Solanum pennellii</i>	3903	4,50E+04	86%	0.0	81.46%	77991103	NC_028641.1
monotypic-Ch06a_26	refeq-genome	8	Solanum pennellii chromosome 4, SPENN200	<i>Solanum pennellii</i>	3901	6,33E+04	86%	0.0	81.36%	77197300	NC_028640.1
monotypic-Ch06a_26	refeq-genome	8	Solanum pennellii chromosome 6, SPENN200	<i>Solanum pennellii</i>	3886	4,27E+04	85%	0.0	81.41%	60730942	NC_028642.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3871	3,87E+03	85%	0.0	81.15%	177091	NW_017672673.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3868	3,87E+03	85%	0.0	81.13%	418852	NW_017670889.1
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4912	<i>Capsicum annuum</i>	3868	4,98E+03	85%	0.0	85.80%	178551	NW_015960903.1
monotypic-Ch06a_26	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	<i>Solanum lycopersicum</i>	3840	3,12E+04	86%	0.0	81.21%	55977580	NC_015439.3
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3829	3,83E+03	86%	0.0	80.92%	177020	NW_017672675.1
monotypic-Ch06a_26	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	<i>Solanum lycopersicum</i>	3829	4,28E+04	86%	0.0	81.18%	65633393	NC_015447.3
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT chromosome 12, NIATTR2	<i>Nicotiana attenuata</i>	3814	1,76E+04	86%	0.0	80.91%	53491057	NC_032000.1
monotypic-Ch06a_26	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	<i>Solanum lycopersicum</i>	3810	5,16E+04	86%	0.0	81.05%	72290146	NC_015440.3
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3803	3,80E+03	85%	0.0	80.97%	351307	NW_017671127.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT chromosome 7, NIATTR2	<i>Nicotiana attenuata</i>	3797	4,65E+04	86%	0.0	80.95%	114765206	NC_031995.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3795	3,80E+03	85%	0.0	80.88%	303526	NW_017671390.1
monotypic-Ch06a_26	refeq-genome	8	Solanum pennellii chromosome 8, SPENN200	<i>Solanum pennellii</i>	3740	3,03E+04	86%	0.0	80.88%	70546659	NC_028644.1
monotypic-Ch06a_26	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	<i>Solanum lycopersicum</i>	3729	3,01E+04	86%	0.0	80.83%	72906345	NC_015446.3
monotypic-Ch06a_26	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	<i>Solanum lycopersicum</i>	3723	4,56E+04	86%	0.0	81.12%	98455869	NC_015438.3
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT chromosome 2, NIATTR2	<i>Nicotiana attenuata</i>	3720	4,72E+04	86%	0.0	80.84%	96359640	NC_031990.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3716	3,72E+03	85%	0.0	80.58%	397494	NW_017670957.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3698	3,70E+03	85%	0.0	80.64%	43788	NW_017676498.1
monotypic-Ch06a_26	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00046	<i>Solanum tuberosum</i>	3679	4,74E+03	86%	0.0	83.71%	2411080	NW_006238971.1
monotypic-Ch06a_26	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00210	<i>Solanum tuberosum</i>	3679	4,74E+03	86%	0.0	83.72%	1116674	NW_006239135.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT chromosome 5, NIATTR2	<i>Nicotiana attenuata</i>	3605	1,26E+04	85%	0.0	80.28%	57012777	NC_031993.1
monotypic-Ch06a_26	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	<i>Solanum lycopersicum</i>	3592	2,52E+04	85%	0.0	80.27%	66723567	NC_015442.3
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3589	3,59E+03	85%	0.0	80.28%	477375	NW_017670730.1
monotypic-Ch06a_26	refeq-genome	8	Solanum pennellii chromosome 2, SPENN200	<i>Solanum pennellii</i>	3589	3,06E+04	85%	0.0	80.33%	59803892	NC_028638.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold17550	<i>Nicotiana tomentosiformis</i>	3557	3,56E+03	85%	0.0	80.06%	68912	NW_008896431.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold124000	<i>Nicotiana tabacum</i>	3539	3,54E+03	85%	0.0	80.05%	21940	NW_015805202.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3491	3,49E+03	81%	0.0	80.60%	11477	NW_017681333.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold36325	<i>Nicotiana sylvestris</i>	3482	3,48E+03	85%	0.0	79.85%	52737	NW_009521971.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3463	3,46E+03	85%	0.0	79.71%	286131	NW_017671511.1
monotypic-Ch06a_26	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	<i>Solanum lycopersicum</i>	3452	3,28E+04	86%	0.0	81.06%	68126176	NC_015449.3
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1354	<i>Capsicum annuum</i>	3441	6,84E+03	97%	0.0	89.07%	1998780	NW_015960394.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT chromosome 1, NIATTR2	<i>Nicotiana attenuata</i>	3428	2,11E+04	86%	0.0	79.94%	110538649	NC_031989.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	<i>Nicotiana attenuata</i>	3428	3,43E+03	85%	0.0	79.56%	36286	NW_017676975.1
monotypic-Ch06a_26	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00313	<i>Solanum tuberosum</i>	3424	9,09E+03	86%	0.0	83.32%	714955	NW_006239248.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold92947	<i>Nicotiana tabacum</i>	3422	3,42E+03	85%	0.0	79.61%	29023	NW_01594907.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold26998	<i>Nicotiana sylvestris</i>	3417	3,42E+03	85%	0.0	79.58%	127419	NW_009511606.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13082	<i>Nicotiana tabacum</i>	3415	3,42E+03	82%	0.0	80.06%	32879	NW_015810064.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold103063	<i>Nicotiana tabacum</i>	3410	3,41E+03	85%	0.0	79.56%	98024	NW_015789620.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold2376	<i>Nicotiana sylvestris</i>	3410	3,41E+03	82%	0.0	80.05%	108876	NW_009491691.1
monotypic-Ch06a_26	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	<i>Solanum lycopersicum</i>	3402	4,22E+04	86%	0.0	79.67%	66557038	NC_015441.3
monotypic-Ch06a_26	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	<i>Solanum lycopersicum</i>	3402	4,12E+04	86%	0.0	79.58%	68175699	NC_015444.3
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT chromosome 3, NIATTR2	<i>Nicotiana attenuata</i>	3363	1,72E+04	86%	0.0	83.38%	72893213	NC_031991.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold13159	<i>Nicotiana sylvestris</i>	3310	3,31E+03	85%	0.0	79.36%	124446	NW_009373903.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold25416	<i>Nicotiana sylvestris</i>	3302	3,30E+03	86%	0.0	79.09%	71009	NW_009509850.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT chromosome 10, NIATTR2	<i>Nicotiana attenuata</i>	3288	9,33E+03	86%	0.0	83.44%	51082206	NC_031998.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1606	<i>Nicotiana tabacum</i>	3273	3,27E+03	85%	0.0	79.21%	85121	NW_015829570.1
monotypic-Ch06a_26	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	<i>Solanum lycopersicum</i>	3265	7,96E+03	85%	0.0	79.14%	49794276	NC_015443.3
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17796	<i>Nicotiana tabacum</i>	3264	3,26E+03	86%	0.0	78.93%	194812	NW_015839666.1

monotypic-Ch06a_26	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold76388	<i>Nicotiana tabacum</i>	3260	3,26E+03	85%	0.0	79.03%	40455	NW_015936080.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold152014	<i>Nicotiana tabacum</i>	3177	3,18E+03	86%	0.0	78.62%	9702	NW_015824184.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	<i>Nicotiana attenuata</i>	3162	3,16E+03	85%	0.0	78.78%	127487	NW_017673598.1
monotypic-Ch06a_26	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00160	<i>Solanum tuberosum</i>	3158	6,34E+03	86%	0.0	82.34%	1367927	NW_006239085.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold506	<i>Nicotiana sylvestris</i>	3121	3,12E+03	85%	0.0	78.57%	206678	NW_009537831.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold158954	<i>Nicotiana tabacum</i>	3118	3,12E+03	85%	0.0	78.51%	52167	NW_015828538.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold37387	<i>Nicotiana sylvestris</i>	3114	3,11E+03	85%	0.0	78.61%	100328	NW_009523150.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold100834	<i>Nicotiana sylvestris</i>	3109	3,11E+03	86%	0.0	78.38%	16882	NW_009339731.1
monotypic-Ch06a_26	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00019	<i>Solanum tuberosum</i>	3099	6,95E+03	86%	0.0	82.43%	3257788	NW_006238945.1
monotypic-Ch06a_26	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00108	<i>Solanum tuberosum</i>	3096	5,59E+03	83%	0.0	81.99%	1692442	NW_006239033.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	<i>Nicotiana attenuata</i>	3092	3,09E+03	80%	0.0	79.23%	115557	NW_017673916.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7235	<i>Nicotiana tabacum</i>	3070	3,07E+03	81%	0.0	78.88%	201617	NW_015932627.1
monotypic-Ch06a_26	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00001	<i>Solanum tuberosum</i>	3059	7,18E+03	85%	0.0	81.85%	7090834	NW_006238927.1
monotypic-Ch06a_26	refeq-genome	8	Solanum pennellii chromosome 9, SPENN9V200	<i>Solanum pennellii</i>	3053	3,26E+04	86%	0.0	79.86%	84057508	NC_028645.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold7876	<i>Nicotiana sylvestris</i>	2990	2,99E+03	86%	0.0	78.01%	125558	NW_009569119.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold8085	<i>Nicotiana sylvestris</i>	2922	2,92E+03	80%	0.0	78.37%	71323	NW_009517442.1
monotypic-Ch06a_26	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	<i>Solanum lycopersicum</i>	2915	2,14E+04	85%	0.0	81.24%	65987440	NC_015445.3
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	<i>Nicotiana attenuata</i>	2881	3,25E+03	82%	0.0	79.64%	529766	NW_017670632.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	<i>Nicotiana attenuata</i>	2787	3,61E+03	86%	0.0	79.80%	221937	NW_017672070.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	<i>Nicotiana attenuata</i>	2784	4,49E+03	86%	0.0	83.36%	52867	NW_017676063.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold88443	<i>Nicotiana tabacum</i>	2760	2,76E+03	81%	0.0	77.83%	50891	NW_015946186.1
monotypic-Ch06a_26	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00200	<i>Solanum tuberosum</i>	2715	4,52E+03	85%	0.0	80.23%	1164273	NW_006239124.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold40969	<i>Nicotiana sylvestris</i>	2713	3,52E+03	84%	0.0	79.60%	73657	NW_009527131.0
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10766	<i>Nicotiana tabacum</i>	2708	3,51E+03	84%	0.0	79.57%	73266	NW_015793144.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	<i>Nicotiana attenuata</i>	2610	3,55E+03	84%	0.0	82.54%	278571	NW_017671558.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1102	<i>Nicotiana sylvestris</i>	2566	3,43E+03	85%	0.0	79.19%	63895	NW_009350137.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	<i>Nicotiana attenuata</i>	2549	3,83E+03	86%	0.0	80.46%	766046	NW_017670410.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7837	<i>Nicotiana tabacum</i>	2549	3,40E+03	85%	0.0	79.12%	64018	NW_015937790.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold36477	<i>Nicotiana tabacum</i>	2531	3,35E+03	85%	0.0	80.34%	51295	NW_015900243.1
monotypic-Ch06a_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold680	<i>Capsicum annuum</i>	2507	4,13E+03	81%	0.0	82.41%	788567	NW_015961649.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold41956	<i>Nicotiana tabacum</i>	2444	3,24E+03	86%	0.0	81.07%	73326	NW_015905338.1
monotypic-Ch06a_26	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00570	<i>Solanum tuberosum</i>	2442	4,67E+03	86%	0.0	85.79%	294773	NW_006239488.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5390	<i>Nicotiana tomentosiformis</i>	2401	2,94E+03	85%	0.0	79.59%	147114	NW_008936819.1
monotypic-Ch06a_26	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00297	<i>Solanum tuberosum</i>	2350	3,70E+03	86%	0.0	81.25%	814182	NW_006239220.1
monotypic-Ch06a_26	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold51791	<i>Nicotiana tabacum</i>	2218	3,94E+03	86%	0.0	79.84%	61637	NW_015914371.1
monotypic-Ch06a_26	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00553	<i>Solanum tuberosum</i>	2217	3,65E+03	82%	0.0	82.84%	279692	NW_006239507.1
monotypic-Ch06a_93	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10159	1,18E+04	100%	0.0	100.00%	219521584	NC_029982.1
monotypic-Ch06a_93	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4464	1,63E+04	100%	0.0	82.94%	215701946	NC_029980.1
monotypic-Ch06a_93	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4303	1,65E+04	100%	0.0	90.47%	221112641	NC_029983.1
monotypic-Ch06a_93	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3493	4,07E+04	100%	0.0	91.06%	217274494	NC_029981.1
monotypic-Ch06a_93	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2100	5,63E+03	88%	0.0	79.91%	301019445	NC_029977.1
monotypic-Ch06a_93	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2056	5,40E+03	91%	0.0	87.32%	220335243	NC_029987.1
monotypic-Ch06a_93	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	848	5,65E+03	93%	0.0	89.21%	261510930	NC_029979.1
monotypic-Ch07e_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9430	1,92E+04	100%	0.0	100.00%	221112641	NC_029983.1
monotypic-Ch07e_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7788	9,41E+03	99%	0.0	94.28%	163962470	NC_029978.1
monotypic-Ch07e_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6861	1,58E+04	100%	0.0	95.33%	220335243	NC_029987.1
monotypic-Ch07e_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold32	<i>Capsicum annuum</i>	6109	6,56E+03	86%	0.0	93.69%	160053	NW_015960960.1
monotypic-Ch07e_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5592	7,26E+03	96%	0.0	90.91%	238794889	NC_029985.1
monotypic-Ch07e_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3910	8,19E+03	98%	0.0	92.97%	153299543	NC_029984.1
monotypic-Ch07e_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3371	8,97E+03	100%	0.0	91.74%	217274494	NC_029981.1
monotypic-Ch07e_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4495	<i>Capsicum annuum</i>	2543	6,58E+03	99%	0.0	89.02%	406145	NW_015960864.1
monotypic-Ch07e_22	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10076	1,09E+04	100%	0.0	100.00%	221112641	NC_029983.1
monotypic-Ch07e_22	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2724	4,87E+03	91%	0.0	84.48%	163962470	NC_029978.1
monotypic-Ch08d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9319	1,22E+04	100%	0.0	100.00%	153299543	NC_029984.1
monotypic-Ch08d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6927	1,52E+04	100%	0.0	91.60%	229934170	NC_029988.1
monotypic-Ch08d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6878	1,34E+04	100%	0.0	91.26%	219521584	NC_029982.1

monotypic-Ch08d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6844	1,40E+04	100%	0.0	91.20%	220335243	NC_029987.1
monotypic-Ch08d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6279	9,66E+03	100%	0.0	89.26%	261510930	NC_029979.1
monotypic-Ch08d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5493	1,82E+04	100%	0.0	91.70%	301019445	NC_029977.1
monotypic-Ch08d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4942	1,71E+04	100%	0.0	87.79%	222112641	NC_029983.1
monotypic-Ch08d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	1853	7,44E+03	84%	0.0	79.92%	217274494	NC_029981.1
monotypic-Ch09a_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9088	1,61E+04	100%	0.0	100.00%	238794889	NC_029985.1
monotypic-Ch09a_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1399	<i>Capsicum annuum</i>	6239	8,45E+03	99%	0.0	94.80%	1742971	NW_015961630.1
monotypic-Ch09a_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6150	9,76E+03	99%	0.0	94.51%	229934170	NC_029988.1
monotypic-Ch09a_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5363	1,05E+04	99%	0.0	92.61%	219521584	NC_029982.1
monotypic-Ch09a_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5072	9,92E+03	99%	0.0	93.11%	261510930	NC_029979.1
monotypic-Ch09a_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4285	4,29E+03	81%	0.0	86.11%	301019445	NC_029977.1
monotypic-Ch10c_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10152	1,15E+04	100%	0.0	100.00%	205736368	NC_029986.1
monotypic-Ch10c_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3984	2,89E+04	98%	0.0	90.58%	220335243	NC_029987.1
monotypic-Ch10c_26	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3476	4,90E+03	97%	0.0	83.39%	238794889	NC_029985.1
monotypic-Ch10f_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9780	1,93E+04	100%	0.0	100.00%	205736368	NC_029986.1
monotypic-Ch10f_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3326	7,39E+03	87%	0.0	91.22%	222112641	NC_029983.1
monotypic-Ch10f_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10074	1,47E+04	100%	0.0	100.00%	205736368	NC_029986.1
monotypic-Ch10f_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5110	8,25E+03	99%	0.0	90.10%	261510930	NC_029979.1
monotypic-Ch10f_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3210	5,42E+03	93%	0.0	85.67%	238794889	NC_029985.1
monotypic-Ch10f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9424	1,04E+04	100%	0.0	100.00%	205736368	NC_029986.1
monotypic-Ch10f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5753	6,07E+03	100%	0.0	87.18%	215701946	NC_029980.1
monotypic-Ch10f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2501	9,01E+03	99%	0.0	88.52%	261510930	NC_029979.1
monotypic-Ch10f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	1788	6,32E+03	83%	0.0	87.75%	229934170	NC_029988.1
monotypic-Ch11a_12	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10006	2,21E+04	100%	0.0	100.00%	220335243	NC_029987.1
monotypic-Ch11a_12	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6124	2,02E+04	98%	0.0	91.12%	217274494	NC_029981.1
monotypic-Ch11a_12	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5216	1,38E+04	97%	0.0	91.34%	238794889	NC_029985.1
monotypic-Ch11a_12	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4643	1,49E+04	99%	0.0	92.31%	261510930	NC_029979.1
monotypic-Ch11a_12	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4043	1,36E+04	96%	0.0	91.94%	229934170	NC_029988.1
monotypic-Ch11a_12	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1399	<i>Capsicum annuum</i>	3430	9,43E+03	93%	0.0	94.53%	1742971	NW_015961630.1
monotypic-Ch11a_12	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3144	1,04E+04	86%	0.0	90.11%	205736368	NC_029986.1
monotypic-Ch11a_12	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	1858	1,22E+04	83%	0.0	75.33%	215701946	NC_029980.1
monotypic-Ch11a_12	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	1696	9,42E+03	85%	0.0	74.83%	301019445	NC_029977.1
monotypic-Ch11a_12	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	1600	5,29E+03	82%	0.0	74.26%	163962470	NC_029978.1
monotypic-Ch11a_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8879	1,03E+04	100%	0.0	100.00%	220335243	NC_029987.1
monotypic-Ch11a_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5890	1,19E+04	100%	0.0	90.11%	217274494	NC_029981.1
monotypic-Ch11a_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5603	9,70E+03	99%	0.0	91.61%	222112641	NC_029983.1
monotypic-Ch11c_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10497	1,28E+04	100%	0.0	100.00%	220335243	NC_029987.1
monotypic-Ch11c_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5227	1,15E+04	100%	0.0	91.93%	219521584	NC_029982.1
monotypic-Ch11c_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4126	1,18E+04	98%	0.0	92.60%	261510930	NC_029979.1
monotypic-Ch11c_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2857	7,34E+03	98%	0.0	82.45%	215701946	NC_029980.1
monotypic-Ch11c_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2512	9,66E+03	97%	0.0	91.93%	153299543	NC_029984.1
monotypic-Ch12a_47	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10000	1,11E+04	100%	0.0	100.00%	229934170	NC_029988.1
monotypic-Ch12a_47	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8161	9,62E+03	100%	0.0	93.86%	301019445	NC_029977.1
monotypic-Ch12a_47	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8028	9,40E+03	100%	0.0	93.39%	205736368	NC_029986.1
monotypic-Ch12a_47	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7456	1,35E+04	100%	0.0	93.59%	261510930	NC_029979.1
monotypic-Ch12a_47	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6968	8,08E+03	99%	0.0	89.96%	219521584	NC_029982.1
monotypic-Ch12a_47	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5622	9,21E+03	100%	0.0	89.89%	215701946	NC_029980.1
monotypic-Ch12e_38	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	12019	1,81E+04	100%	0.0	100.00%	229934170	NC_029988.1
monotypic-Ch12e_38	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4089	9,36E+03	89%	0.0	86.44%	215701946	NC_029980.1
monotypic-Ch12e_38	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3969	9,61E+03	86%	0.0	86.82%	220335243	NC_029987.1
monotypic-Ch12e_38	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3831	7,15E+03	99%	0.0	85.24%	163962470	NC_029978.1
monotypic-Ch12e_38	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3777	6,27E+03	89%	0.0	85.17%	238794889	NC_029985.1
monotypic-Ch12e_38	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2959	6,77E+03	98%	0.0	84.65%	205736368	NC_029986.1
monotypic-Ch12e_38	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2896	7,37E+03	89%	0.0	80.82%	222112641	NC_029983.1
monotypic-Ch12e_38	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	1840	5,61E+03	90%	0.0	83.19%	219521584	NC_029982.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	12201	2,49E+04	100%	0.0	100.00%	153299543	NC_029984.1

GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6927	9,66E+04	100%	0.0	93.61%	205736368	NC_029986.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5866	1,71E+08	100%	0.0	93.05%	261510930	NC_029979.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5760	8,12E+04	100%	0.0	89.31%	301019445	NC_029977.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5729	4,57E+04	100%	0.0	90.67%	238794889	NC_029985.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5701	8,87E+04	100%	0.0	88.99%	219521584	NC_029982.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5672	9,46E+04	100%	0.0	89.72%	220335243	NC_029987.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2138	<i>Capsicum annuum</i>	5646	8,70E+03	99%	0.0	88.81%	681955	NW_015960370.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5603	1,07E+08	100%	0.0	88.70%	217274494	NC_029981.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5578	9,85E+04	100%	0.0	88.53%	222112641	NC_029983.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4841	5,52E+04	100%	0.0	85.78%	229934170	NC_029988.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1514	<i>Capsicum annuum</i>	4839	7,20E+03	100%	0.0	85.76%	427058	NW_015960414.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold189	<i>Capsicum annuum</i>	4697	1,47E+04	100%	0.0	86.20%	3791111	NW_015960691.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4590	7,87E+04	100%	0.0	89.33%	215701946	NC_029980.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold924	<i>Capsicum annuum</i>	4253	1,02E+04	86%	0.0	92.73%	1342734	NW_015960575.1
GypsyCa_01	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00013	<i>Solanum tuberosum</i>	3934	6,11E+03	83%	0.0	86.64%	3422333	NW_006238940.1
GypsyCa_01	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00067	<i>Solanum tuberosum</i>	3773	6,04E+03	86%	0.0	85.72%	2156588	NW_006238991.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3400	4,51E+04	100%	0.0	88.24%	163962470	NC_029978.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold24528	<i>Capsicum annuum</i>	2682	5,18E+03	80%	0.0	83.15%	145047	NW_015960905.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9	<i>Capsicum annuum</i>	2591	7,56E+03	97%	0.0	87.76%	6567951	NW_015961815.1
GypsyCa_01	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold148	<i>Capsicum annuum</i>	2209	6,39E+03	93%	0.0	87.03%	1323090	NW_015961290.1
GypsyCa_01	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00009	<i>Solanum tuberosum</i>	2069	5,76E+03	85%	0.0	85.84%	3955149	NW_006238935.1
GypsyCa_01	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00001	<i>Solanum tuberosum</i>	1441	7,26E+03	89%	0.0	88.45%	7090834	NW_006238927.1
GypsyCa_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10864	6,52E+04	100%	0.0	100.00%	219521584	NC_029982.1
GypsyCa_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5792	7,79E+04	100%	0.0	93.39%	301019445	NC_029977.1
GypsyCa_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5694	5,09E+04	100%	0.0	92.92%	261510930	NC_029979.1
GypsyCa_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	5675	1,00E+04	98%	0.0	92.84%	2470984	NW_015961229.1
GypsyCa_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5670	7,56E+04	100%	0.0	92.84%	163962470	NC_029978.1
GypsyCa_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5620	4,02E+04	100%	0.0	92.58%	205736368	NC_029986.1
GypsyCa_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5426	5,52E+04	100%	0.0	91.77%	222112641	NC_029983.1
GypsyCa_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5310	3,64E+04	100%	0.0	91.21%	238794889	NC_029985.1
GypsyCa_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5227	3,67E+04	100%	0.0	90.79%	215701946	NC_029980.1
GypsyCa_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5118	3,37E+04	100%	0.0	90.30%	229934170	NC_029988.1
GypsyCa_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4841	3,16E+04	100%	0.0	92.13%	220335243	NC_029987.1
GypsyCa_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2719	<i>Capsicum annuum</i>	4667	7,99E+03	99%	0.0	92.11%	277434	NW_015960900.1
GypsyCa_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4582	2,23E+04	100%	0.0	89.39%	153299543	NC_029984.1
GypsyCa_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4250	2,43E+04	100%	0.0	86.38%	217274494	NC_029981.1
GypsyCa_02	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold35263	<i>Capsicum annuum</i>	1923	5,74E+03	82%	0.0	87.18%	18848	NW_015960802.1
GypsyCa_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	14063	8,93E+04	100%	0.0	100.00%	215701946	NC_029980.1
GypsyCa_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5494	9,58E+04	99%	0.0	85.71%	220335243	NC_029987.1
GypsyCa_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3792	1,37E+08	96%	0.0	86.20%	261510930	NC_029979.1
GypsyCa_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3746	7,66E+04	97%	0.0	83.58%	222112641	NC_029983.1
GypsyCa_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3626	7,95E+04	97%	0.0	85.22%	238794889	NC_029985.1
GypsyCa_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3446	8,41E+04	97%	0.0	82.20%	163962470	NC_029978.1
GypsyCa_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2918	7,76E+04	90%	0.0	81.60%	219521584	NC_029982.1
GypsyCa_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2736	1,15E+08	81%	0.0	81.12%	301019445	NC_029977.1
GypsyCa_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2669	9,07E+04	99%	0.0	80.90%	217274494	NC_029981.1
GypsyCa_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2353	8,33E+04	97%	0.0	79.30%	229934170	NC_029988.1
GypsyCa_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2202	7,26E+04	98%	0.0	81.08%	205736368	NC_029986.1
GypsyCa_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2115	6,40E+04	97%	0.0	77.93%	153299543	NC_029984.1
GypsyCa_03	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4183	<i>Capsicum annuum</i>	1378	9,94E+03	81%	0.0	75.62%	1223518	NW_015961565.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	17014	1,20E+09	100%	0.0	99.90%	205736368	NC_029986.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	12451	1,51E+09	100%	0.0	90.99%	217274494	NC_029981.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11524	1,17E+09	100%	0.0	90.21%	222112641	NC_029983.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11444	1,18E+09	100%	0.0	90.05%	229934170	NC_029988.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10957	1,43E+09	100%	0.0	90.87%	238794889	NC_029985.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10545	7,18E+08	100%	0.0	90.14%	163962470	NC_029978.1

GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10442	1,29E+09	100%	0.0	87.97%	261510930	NC_029979.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9978	7,50E+08	100%	0.0	89.15%	153299543	NC_029984.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2274	<i>Capsicum annuum</i>	9734	2,42E+04	100%	0.0	90.17%	1395869	NW_015961152.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9635	1,17E+09	100%	0.0	88.93%	219521584	NC_029982.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9522	1,45E+09	100%	0.0	89.32%	301019445	NC_029977.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8996	1,23E+09	100%	0.0	88.45%	215701946	NC_029980.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	8820	2,83E+04	100%	0.0	88.43%	7059680	NW_015960855.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1722	<i>Capsicum annuum</i>	8678	1,61E+04	100%	0.0	88.59%	153596	NW_015960391.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8288	1,05E+09	100%	0.0	89.56%	220335243	NC_029987.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold8281	<i>Capsicum annuum</i>	7627	1,20E+04	95%	0.0	86.75%	213152	NW_015961792.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold188	<i>Capsicum annuum</i>	6885	1,38E+04	89%	0.0	84.60%	980346	NW_015960557.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1083	<i>Capsicum annuum</i>	6673	2,42E+04	99%	0.0	85.42%	207146	NW_015960434.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold395	<i>Capsicum annuum</i>	6488	1,24E+04	82%	0.0	84.86%	1727395	NW_015960459.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold170	<i>Capsicum annuum</i>	6447	1,31E+04	86%	0.0	85.72%	2217302	NW_015960390.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2375	<i>Capsicum annuum</i>	6381	2,76E+04	90%	0.0	85.42%	1171512	NW_015961637.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	6301	2,00E+04	95%	0.0	85.17%	3311724	NW_015960635.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold516	<i>Capsicum annuum</i>	6156	1,16E+04	80%	0.0	84.96%	4299619	NW_015961344.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1422	<i>Capsicum annuum</i>	6111	4,17E+04	100%	0.0	84.76%	802096	NW_015960747.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold90	<i>Capsicum annuum</i>	6109	6,70E+04	100%	0.0	91.70%	6074945	NW_015960458.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1894	<i>Capsicum annuum</i>	6109	2,54E+04	96%	0.0	85.52%	2334823	NW_015961490.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1741	<i>Capsicum annuum</i>	6085	2,08E+04	89%	0.0	84.69%	1897442	NW_015960687.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2400	<i>Capsicum annuum</i>	5947	1,52E+04	95%	0.0	84.15%	1353284	NW_015960968.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1973	<i>Capsicum annuum</i>	5930	3,59E+04	100%	0.0	86.10%	1714326	NW_015961422.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2948	<i>Capsicum annuum</i>	5879	7,53E+03	81%	0.0	83.13%	898408	NW_015961258.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold555	<i>Capsicum annuum</i>	5533	1,14E+04	85%	0.0	88.25%	1396435	NW_015960461.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold349	<i>Capsicum annuum</i>	5533	1,37E+04	90%	0.0	86.64%	2130419	NW_015960729.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1354	<i>Capsicum annuum</i>	5391	1,67E+04	84%	0.0	84.41%	1998780	NW_015960394.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold949	<i>Capsicum annuum</i>	5219	5,31E+04	100%	0.0	90.33%	3128562	NW_015961487.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1399	<i>Capsicum annuum</i>	5090	5,29E+04	100%	0.0	82.97%	1742971	NW_015961630.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2523	<i>Capsicum annuum</i>	5073	1,12E+04	94%	0.0	87.32%	735434	NW_015960464.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold181	<i>Capsicum annuum</i>	4959	1,22E+04	84%	0.0	82.43%	2703691	NW_015961388.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1339	<i>Capsicum annuum</i>	4870	2,51E+04	96%	0.0	85.60%	928404	NW_015960451.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2059	<i>Capsicum annuum</i>	4608	2,71E+04	95%	0.0	87.69%	416460	NW_015960523.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold16	<i>Capsicum annuum</i>	4423	3,68E+04	91%	0.0	86.32%	2023789	NW_015960959.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2866	<i>Capsicum annuum</i>	4386	7,30E+03	81%	0.0	78.94%	1254371	NW_015961127.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	4141	2,28E+04	96%	0.0	84.42%	2470984	NW_015961229.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1110	<i>Capsicum annuum</i>	3856	1,64E+04	96%	0.0	85.69%	740557	NW_015961474.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold558	<i>Capsicum annuum</i>	3530	1,70E+04	84%	0.0	82.45%	309877	NW_015960759.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold7976	<i>Capsicum annuum</i>	3452	1,15E+04	89%	0.0	89.31%	57623	NW_015961638.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1194	<i>Capsicum annuum</i>	3286	2,26E+04	100%	0.0	83.18%	61568	NW_015960372.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1765	<i>Capsicum annuum</i>	3014	2,20E+04	94%	0.0	82.77%	289511	NW_015961400.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1929	<i>Capsicum annuum</i>	2998	1,56E+04	95%	0.0	76.70%	996492	NW_015961795.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold450	<i>Capsicum annuum</i>	2737	1,56E+04	94%	0.0	89.93%	175795	NW_015961071.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold891	<i>Capsicum annuum</i>	2726	9,51E+03	81%	0.0	80.50%	1431046	NW_015960469.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold229	<i>Capsicum annuum</i>	2647	1,31E+04	97%	0.0	80.21%	174263	NW_015961369.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capsicum annuum</i>	2377	1,27E+04	92%	0.0	78.73%	2397553	NW_015961318.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3026	<i>Capsicum annuum</i>	2342	2,05E+04	91%	0.0	78.74%	1386929	NW_015961102.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3027	<i>Capsicum annuum</i>	2331	1,00E+04	88%	0.0	77.00%	193869	NW_015961172.1
GypsyCa_04	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold171	<i>Capsicum annuum</i>	1820	1,43E+04	82%	0.0	79.76%	887051	NW_015960473.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	19028	2,34E+09	100%	0.0	100.00%	163962470	NC_029978.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	19023	1,71E+09	100%	0.0	99.99%	222112641	NC_029983.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	19023	1,81E+09	100%	0.0	99.99%	220335243	NC_029987.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	19023	1,78E+09	100%	0.0	99.99%	229934170	NC_029988.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	19017	1,56E+09	100%	0.0	99.98%	238794889	NC_029985.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold277	<i>Capsicum annuum</i>	19017	4,30E+04	100%	0.0	99.98%	448122	NW_015961390.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	19012	2,10E+09	100%	0.0	99.97%	219521584	NC_029982.1

GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	19006	2,44E+09	100%	0.0	99.96%	261510930	NC_029979.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	18979	2,08E+09	100%	0.0	99.91%	215701946	NC_029980.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	18975	2,61E+09	100%	0.0	99.90%	301019445	NC_029977.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	18975	1,58E+09	100%	0.0	99.90%	205736368	NC_029986.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	18962	1,64E+09	100%	0.0	99.87%	153299543	NC_029984.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold985	<i>Capsicum annuum</i>	18787	3,19E+04	100%	0.0	99.58%	1349800	NW_015961816.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3202	<i>Capsicum annuum</i>	18768	2,86E+04	100%	0.0	99.56%	302159	NW_015961119.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	18702	1,63E+09	100%	0.0	99.63%	217274494	NC_029981.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold258	<i>Capsicum annuum</i>	17758	8,70E+04	100%	0.0	99.76%	3098259	NW_015961180.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2057	<i>Capsicum annuum</i>	17302	3,71E+04	100%	0.0	97.44%	330742	NW_015961751.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3026	<i>Capsicum annuum</i>	16872	2,55E+04	100%	0.0	98.56%	1386929	NW_015961102.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9433	<i>Capsicum annuum</i>	16183	3,26E+04	100%	0.0	99.98%	23376	NW_015961766.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3745	<i>Capsicum annuum</i>	15439	3,14E+04	100%	0.0	99.93%	97749	NW_015961051.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold561	<i>Capsicum annuum</i>	15343	3,19E+04	100%	0.0	99.94%	1145873	NW_015961012.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2639	<i>Capsicum annuum</i>	15073	2,17E+04	100%	0.0	98.68%	306189	NW_015960882.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3079	<i>Capsicum annuum</i>	14233	2,19E+04	100%	0.0	98.61%	299956	NW_015961555.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2298	<i>Capsicum annuum</i>	14011	3,98E+04	99%	0.0	99.51%	1494726	NW_015960454.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3859	<i>Capsicum annuum</i>	13882	2,17E+04	100%	0.0	98.33%	99728	NW_015961156.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold171	<i>Capsicum annuum</i>	13871	2,02E+04	100%	0.0	99.80%	887051	NW_015960473.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1124	<i>Capsicum annuum</i>	13834	5,43E+04	100%	0.0	99.97%	2110277	NW_015961370.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1651	<i>Capsicum annuum</i>	13730	2,86E+04	97%	0.0	99.97%	331682	NW_015961206.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold433	<i>Capsicum annuum</i>	13503	2,96E+04	97%	0.0	99.80%	1277097	NW_015961081.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1390	<i>Capsicum annuum</i>	13298	5,06E+04	96%	0.0	99.79%	1730405	NW_015960783.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold949	<i>Capsicum annuum</i>	10739	5,70E+04	100%	0.0	99.69%	3128562	NW_015961487.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1037	<i>Capsicum annuum</i>	10635	2,72E+04	91%	0.0	98.21%	2186250	NW_015960681.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1246	<i>Capsicum annuum</i>	10549	2,88E+04	88%	0.0	98.87%	3388803	NW_015960834.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4971	<i>Capsicum annuum</i>	10462	2,03E+04	98%	0.0	99.74%	81603	NW_015960449.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2938	<i>Capsicum annuum</i>	9049	2,01E+04	97%	0.0	97.54%	524377	NW_015960410.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold775	<i>Capsicum annuum</i>	9014	1,44E+04	88%	0.0	92.52%	696435	NW_015961719.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	8695	2,75E+04	100%	0.0	98.97%	2470984	NW_015961229.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold619	<i>Capsicum annuum</i>	8329	4,08E+04	100%	0.0	97.29%	2512642	NW_015960498.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5089	<i>Capsicum annuum</i>	8224	4,25E+04	100%	0.0	97.26%	522036	NW_015961006.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold658	<i>Capsicum annuum</i>	7904	1,37E+04	82%	0.0	89.66%	613851	NW_015961079.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold175	<i>Capsicum annuum</i>	7668	2,28E+04	86%	0.0	95.41%	655560	NW_015960845.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold519	<i>Capsicum annuum</i>	7518	1,99E+04	86%	0.0	88.59%	1101621	NW_015961578.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold90	<i>Capsicum annuum</i>	7339	8,12E+04	86%	0.0	89.09%	6074945	NW_015960458.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3695	<i>Capsicum annuum</i>	6684	1,14E+04	84%	0.0	91.33%	350731	NW_015960809.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold349	<i>Capsicum annuum</i>	6601	1,78E+04	80%	0.0	90.33%	2130419	NW_015960729.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1872	<i>Capsicum annuum</i>	6503	4,50E+04	100%	0.0	99.58%	241253	NW_015960977.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold181	<i>Capsicum annuum</i>	6176	7,76E+04	100%	0.0	95.04%	2703691	NW_015961388.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4063	<i>Capsicum annuum</i>	5055	2,16E+04	85%	0.0	96.98%	1657713	NW_015960826.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1463	<i>Capsicum annuum</i>	4964	1,73E+04	82%	0.0	94.90%	343407	NW_015961513.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1307	<i>Capsicum annuum</i>	4747	1,62E+04	86%	0.0	90.69%	1549638	NW_015960466.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold26496	<i>Capsicum annuum</i>	4721	1,89E+04	85%	0.0	98.68%	65566	NW_015960947.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold6075	<i>Capsicum annuum</i>	4582	1,20E+04	85%	0.0	89.81%	316563	NW_015960475.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold189	<i>Capsicum annuum</i>	4543	1,91E+04	83%	0.0	80.91%	3791111	NW_015960691.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3499	<i>Capsicum annuum</i>	4536	1,99E+04	99%	0.0	98.45%	681687	NW_015960538.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1633	<i>Capsicum annuum</i>	4530	2,57E+04	97%	0.0	98.41%	1620172	NW_015961041.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold170	<i>Capsicum annuum</i>	4403	2,18E+04	80%	0.0	97.42%	2217302	NW_015960390.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2343	<i>Capsicum annuum</i>	4320	1,63E+04	86%	0.0	81.61%	440837	NW_015961556.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold937	<i>Capsicum annuum</i>	4226	1,96E+04	92%	0.0	96.60%	1465884	NW_015961782.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	3958	2,55E+04	86%	0.0	79.63%	3493078	NW_015960517.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2322	<i>Capsicum annuum</i>	2859	1,19E+04	83%	0.0	89.41%	1668020	NW_015961155.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold454	<i>Capsicum annuum</i>	2815	1,23E+04	83%	0.0	77.12%	3392484	NW_015961509.1
GypsyCa_09	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2274	<i>Capsicum annuum</i>	2558	1,52E+04	85%	0.0	93.09%	1395869	NW_015961152.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	18148	1,92E+09	100%	0.0	100.00%	219521584	NC_029982.1

GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11952	1,24E+09	100%	0.0	88.79%	163962470	NC_029978.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10818	2,20E+09	100%	0.0	86.70%	301019445	NC_029977.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9897	1,20E+09	100%	0.0	85.01%	153299543	NC_029984.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold551	<i>Capsicum annuum</i>	9751	2,72E+04	100%	0.0	89.01%	761547	NW_015961519.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold619	<i>Capsicum annuum</i>	9657	2,95E+04	100%	0.0	84.67%	2512642	NW_015960498.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9012	1,87E+09	100%	0.0	83.63%	215701946	NC_029980.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold603	<i>Capsicum annuum</i>	8927	2,05E+04	100%	0.0	85.46%	1800753	NW_015960521.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8776	1,84E+09	100%	0.0	83.02%	220335243	NC_029987.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8776	1,66E+09	100%	0.0	83.02%	229934170	NC_029988.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8516	1,80E+09	100%	0.0	82.57%	222112641	NC_029983.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8484	1,64E+09	100%	0.0	83.34%	217274494	NC_029981.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8390	2,15E+09	100%	0.0	82.31%	238794889	NC_029985.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7958	1,85E+09	100%	0.0	82.60%	261510930	NC_029979.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold189	<i>Capsicum annuum</i>	7936	2,28E+04	100%	0.0	81.51%	3791111	NW_015960691.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7744	1,70E+09	100%	0.0	82.36%	205736368	NC_029986.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1670	<i>Capsicum annuum</i>	7400	3,10E+04	100%	0.0	81.83%	1202455	NW_015961465.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1923	<i>Capsicum annuum</i>	7234	3,86E+04	99%	0.0	88.74%	2480791	NW_015961228.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2569	<i>Capsicum annuum</i>	7203	1,63E+04	100%	0.0	80.22%	624013	NW_015961724.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4467	<i>Capsicum annuum</i>	6549	1,41E+04	100%	0.0	81.84%	686372	NW_015961169.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1038	<i>Capsicum annuum</i>	6466	1,72E+04	97%	0.0	82.30%	2164954	NW_015960831.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2788	<i>Capsicum annuum</i>	6381	1,45E+04	97%	0.0	82.74%	433018	NW_015961312.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold742	<i>Capsicum annuum</i>	6296	4,76E+04	100%	0.0	81.95%	2382255	NW_015961471.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2348	<i>Capsicum annuum</i>	6048	7,63E+03	81%	0.0	83.93%	373378	NW_015960596.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold301	<i>Capsicum annuum</i>	6019	2,87E+04	100%	0.0	81.49%	1226190	NW_015960697.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1620	<i>Capsicum annuum</i>	5853	9,76E+03	95%	0.0	81.52%	500299	NW_015961331.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold181	<i>Capsicum annuum</i>	5674	2,52E+04	99%	0.0	82.65%	2703691	NW_015961388.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold516	<i>Capsicum annuum</i>	5483	3,21E+04	100%	0.0	82.33%	4299619	NW_015961344.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold90	<i>Capsicum annuum</i>	5328	4,84E+04	89%	0.0	82.87%	6074945	NW_015960458.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	5193	8,78E+04	100%	0.0	84.94%	7059680	NW_015960855.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	5086	3,85E+04	100%	0.0	82.61%	3311724	NW_015960635.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2189	<i>Capsicum annuum</i>	4975	1,98E+04	98%	0.0	82.33%	1569264	NW_015960648.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9	<i>Capsicum annuum</i>	4698	7,68E+04	95%	0.0	83.50%	6567951	NW_015961815.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold296	<i>Capsicum annuum</i>	4643	2,03E+04	93%	0.0	82.52%	1995018	NW_015961641.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold171	<i>Capsicum annuum</i>	4265	3,06E+04	100%	0.0	86.91%	887051	NW_015960473.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1124	<i>Capsicum annuum</i>	4159	2,26E+04	97%	0.0	82.59%	2110277	NW_015961370.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold846	<i>Capsicum annuum</i>	3980	1,54E+04	93%	0.0	81.67%	1232610	NW_015960818.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4747	<i>Capsicum annuum</i>	3716	1,90E+04	96%	0.0	80.30%	1151343	NW_015960460.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1973	<i>Capsicum annuum</i>	3221	2,12E+04	87%	0.0	76.71%	1714326	NW_015961422.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold448	<i>Capsicum annuum</i>	3192	2,41E+04	95%	0.0	75.91%	1323857	NW_015960958.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3313	<i>Capsicum annuum</i>	3125	6,97E+03	87%	0.0	76.97%	770237	NW_015961029.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold170	<i>Capsicum annuum</i>	3118	1,48E+04	81%	0.0	83.65%	2217302	NW_015960390.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2707	<i>Capsicum annuum</i>	3072	1,08E+04	83%	0.0	83.63%	1896504	NW_015961173.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold660	<i>Capsicum annuum</i>	3062	8,93E+03	80%	0.0	77.01%	663463	NW_015961299.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold933	<i>Capsicum annuum</i>	3013	2,01E+04	99%	0.0	84.41%	1853301	NW_015961435.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1633	<i>Capsicum annuum</i>	2872	1,30E+04	91%	0.0	77.15%	1620172	NW_015961041.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1587	<i>Capsicum annuum</i>	2780	2,15E+04	85%	0.0	77.32%	1400685	NW_015961196.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3339	<i>Capsicum annuum</i>	2721	1,88E+04	98%	0.0	87.45%	1234997	NW_015960491.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold949	<i>Capsicum annuum</i>	2649	2,47E+04	96%	0.0	76.59%	3128562	NW_015961487.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2298	<i>Capsicum annuum</i>	2623	1,33E+04	92%	0.0	76.51%	1494726	NW_015960454.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	2586	1,77E+04	92%	0.0	76.43%	3493078	NW_015960517.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold891	<i>Capsicum annuum</i>	2542	1,41E+04	100%	0.0	83.15%	1431046	NW_015960469.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2948	<i>Capsicum annuum</i>	2516	1,70E+04	90%	0.0	76.28%	898408	NW_015961258.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2400	<i>Capsicum annuum</i>	2501	7,76E+03	96%	0.0	76.12%	1353284	NW_015960968.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1894	<i>Capsicum annuum</i>	2475	1,38E+04	97%	0.0	76.10%	2334823	NW_015961490.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1647	<i>Capsicum annuum</i>	2464	6,72E+03	81%	0.0	76.08%	1362880	NW_015960892.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3037	<i>Capsicum annuum</i>	2444	1,35E+04	84%	0.0	75.95%	1979607	NW_015960688.1

GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capsicum annuum</i>	2422	1,34E+04	91%	0.0	76.12%	2397553	NW_015961318.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1754	<i>Capsicum annuum</i>	2420	1,09E+04	91%	0.0	77.92%	1029277	NW_015960393.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1341	<i>Capsicum annuum</i>	2418	2,91E+04	100%	0.0	76.75%	1906221	NW_015960665.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4063	<i>Capsicum annuum</i>	2359	1,53E+04	95%	0.0	75.71%	1657713	NW_015960261.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold969	<i>Capsicum annuum</i>	2357	8,52E+03	97%	0.0	75.70%	1557263	NW_015961740.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3224	<i>Capsicum annuum</i>	2043	6,86E+03	93%	0.0	82.48%	891864	NW_015961692.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold454	<i>Capsicum annuum</i>	1945	2,30E+04	98%	0.0	74.23%	3392484	NW_015961509.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4118	<i>Capsicum annuum</i>	1934	6,36E+03	80%	0.0	78.18%	593898	NW_015961463.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2523	<i>Capsicum annuum</i>	1842	9,01E+03	93%	0.0	77.94%	735434	NW_015960464.1
GypsyCa_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9842	<i>Capsicum annuum</i>	1703	5,54E+03	80%	0.0	79.96%	206340	NW_015961066.1
GypsyCa_14	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9225	1,15E+04	100%	0.0	100.00%	163962470	NC_029978.1
GypsyCa_14	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9149	1,03E+04	100%	0.0	99.72%	219521584	NC_029982.1
GypsyCa_14	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9121	1,13E+04	100%	0.0	99.62%	238794889	NC_029985.1
GypsyCa_14	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9119	3,09E+04	100%	0.0	99.64%	261510930	NC_029979.1
GypsyCa_14	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9099	1,44E+04	100%	0.0	99.56%	301019445	NC_029977.1
GypsyCa_14	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9071	1,95E+04	100%	0.0	99.46%	229934170	NC_029988.1
GypsyCa_14	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9049	2,28E+04	100%	0.0	99.38%	153299543	NC_029984.1
GypsyCa_14	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7788	1,01E+04	97%	0.0	99.88%	222112641	NC_029983.1
GypsyCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9701	1,98E+04	100%	0.0	100.00%	261510930	NC_029979.1
GypsyCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7969	2,74E+04	100%	0.0	93.99%	301019445	NC_029977.1
GypsyCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7753	8,86E+03	100%	0.0	93.25%	205736368	NC_029986.1
GypsyCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7661	1,40E+04	100%	0.0	93.01%	219521584	NC_029982.1
GypsyCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7633	2,34E+04	100%	0.0	92.86%	153299543	NC_029984.1
GypsyCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold688	<i>Capsicum annuum</i>	6231	7,68E+03	100%	0.0	90.63%	1322070	NW_015960950.1
GypsyCa_15	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00322	<i>Solanum tuberosum</i>	5718	1,08E+04	96%	0.0	87.41%	720736	NW_006239245.1
GypsyCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1653	<i>Capsicum annuum</i>	5059	8,61E+03	99%	0.0	92.08%	913988	NW_015961387.1
GypsyCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4933	1,34E+04	100%	0.0	88.83%	163962470	NC_029978.1
GypsyCa_15	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold186433	<i>Nicotiana tabacum</i>	4353	4,76E+03	94%	0.0	83.25%	22458	NW_015844369.1
GypsyCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4217	8,09E+03	100%	0.0	87.87%	215701946	NC_029980.1
GypsyCa_15	refeq-genome	8	Solanum pennellii chromosome 10, SPENN200	<i>Solanum pennellii</i>	4209	1,57E+04	96%	0.0	82.77%	82529941	NC_028646.1
GypsyCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4161	9,60E+03	97%	0.0	90.28%	229934170	NC_029988.1
GypsyCa_15	refeq-genome	8	Solanum pennellii chromosome 7, SPENN200	<i>Solanum pennellii</i>	4089	5,62E+03	94%	0.0	82.38%	79292169	NC_028643.1
GypsyCa_15	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00078	<i>Solanum tuberosum</i>	3954	8,18E+03	96%	0.0	81.17%	83305730	NC_028648.1
GypsyCa_15	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold75504	<i>Nicotiana tabacum</i>	3864	6,27E+03	96%	0.0	88.61%	1956963	NW_006239005.1
GypsyCa_15	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold33719	<i>Nicotiana sylvestris</i>	3747	4,32E+03	93%	0.0	82.87%	79457	NW_015935321.1
GypsyCa_15	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00424	<i>Solanum tuberosum</i>	3742	4,32E+03	93%	0.0	82.85%	96466	NW_009519075.1
GypsyCa_15	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	<i>Solanum lycopersicum</i>	3714	6,20E+03	92%	0.0	88.61%	499964	NW_006239350.1
GypsyCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3703	3,77E+03	90%	0.0	80.94%	68175699	NC_015444.3
GypsyCa_15	refeq-genome	8	Nicotiana attenuata strain UT chromosome 3, NIATT2	<i>Nicotiana attenuata</i>	3561	1,44E+04	100%	0.0	92.64%	222112641	NC_029983.1
GypsyCa_15	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00212	<i>Solanum tuberosum</i>	3378	3,50E+03	88%	0.0	80.30%	72893213	NC_031991.1
GypsyCa_15	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	<i>Solanum lycopersicum</i>	3081	4,96E+03	89%	0.0	85.73%	1104745	NW_006239138.1
GypsyCa_15	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	<i>Solanum lycopersicum</i>	2928	3,51E+03	80%	0.0	81.51%	68126176	NC_015449.3
GypsyCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2902	6,73E+03	96%	0.0	81.76%	65633393	NC_015447.3
GypsyCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2811	1,28E+04	96%	0.0	89.40%	220335243	NC_029987.1
GypsyCa_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold7810	<i>Capsicum annuum</i>	2427	8,15E+03	86%	0.0	91.80%	116558	NW_015961118.1
GypsyCa_15	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold41550	<i>Nicotiana tabacum</i>	2189	4,52E+03	94%	0.0	83.75%	160711	NW_015904978.1
GypsyCa_15	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold11955	<i>Nicotiana tomentosiformis</i>	2189	4,52E+03	93%	0.0	83.95%	349838	NW_008850213.1
GypsyCa_15	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold309	<i>Nicotiana tabacum</i>	1552	4,13E+03	93%	0.0	82.65%	169657	NW_015888989.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23385	2,29E+09	100%	0.0	100.00%	163962470	NC_029978.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23385	3,11E+09	100%	0.0	100.00%	261510930	NC_029979.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23385	2,24E+09	100%	0.0	100.00%	215701946	NC_029980.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23385	2,41E+09	100%	0.0	100.00%	217274494	NC_029981.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23385	3,15E+09	100%	0.0	100.00%	219521584	NC_029982.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23385	1,32E+09	100%	0.0	100.00%	153299543	NC_029984.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23385	3,46E+09	100%	0.0	100.00%	238794889	NC_029985.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23385	1,27E+09	100%	0.0	100.00%	220335243	NC_029987.1

GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23385	3,45E+09	100%	0.0	100.00%	229934170	NC_029988.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1647	<i>Capsicum annuum</i>	23385	2,89E+04	100%	0.0	100.00%	1362880	NW_015960892.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3478	<i>Capsicum annuum</i>	23385	5,67E+04	100%	0.0	100.00%	211087	NW_015961510.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9337	<i>Capsicum annuum</i>	23385	2,89E+04	100%	0.0	100.00%	397715	NW_015961734.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23379	9,53E+08	100%	0.0	99.99%	222112641	NC_029983.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23379	1,88E+09	100%	0.0	99.99%	205736368	NC_029986.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3026	<i>Capsicum annuum</i>	23379	9,33E+04	100%	0.0	99.99%	1386929	NW_015961102.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23374	1,52E+09	100%	0.0	99.98%	301019445	NC_029977.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1434	<i>Capsicum annuum</i>	23374	3,38E+04	100%	0.0	99.98%	1100793	NW_015960373.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2635	<i>Capsicum annuum</i>	23374	2,90E+04	100%	0.0	99.98%	699991	NW_015960547.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1341	<i>Capsicum annuum</i>	23374	3,02E+04	100%	0.0	99.98%	1906221	NW_015960665.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold640	<i>Capsicum annuum</i>	23374	2,93E+04	100%	0.0	99.98%	1087533	NW_015960952.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold258	<i>Capsicum annuum</i>	23374	6,45E+04	100%	0.0	99.98%	3098259	NW_015961180.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1923	<i>Capsicum annuum</i>	23374	5,70E+04	100%	0.0	99.98%	2480791	NW_015961228.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1124	<i>Capsicum annuum</i>	23374	8,26E+04	100%	0.0	99.98%	2110277	NW_015961370.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3592	<i>Capsicum annuum</i>	23374	5,98E+04	100%	0.0	99.98%	707215	NW_015961728.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9	<i>Capsicum annuum</i>	23374	8,69E+04	100%	0.0	99.98%	6567951	NW_015961815.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	23368	6,25E+04	100%	0.0	99.98%	3493078	NW_015960517.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	23368	5,74E+04	100%	0.0	99.98%	3311724	NW_015960635.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2189	<i>Capsicum annuum</i>	23368	2,89E+04	100%	0.0	99.98%	1569264	NW_015960648.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold181	<i>Capsicum annuum</i>	23368	2,90E+04	100%	0.0	99.98%	2703691	NW_015961388.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold438	<i>Capsicum annuum</i>	23363	5,96E+04	100%	0.0	99.97%	1337825	NW_015961507.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2715	<i>Capsicum annuum</i>	23357	2,88E+04	100%	0.0	99.96%	316776	NW_015960519.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold846	<i>Capsicum annuum</i>	23357	2,90E+04	100%	0.0	99.97%	1232610	NW_015960818.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold8606	<i>Capsicum annuum</i>	23355	2,90E+04	100%	0.0	99.95%	138485	NW_015960788.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2274	<i>Capsicum annuum</i>	23353	1,09E+08	100%	0.0	99.95%	1395869	NW_015961152.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold775	<i>Capsicum annuum</i>	23353	3,02E+04	100%	0.0	99.95%	696435	NW_015961719.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4183	<i>Capsicum annuum</i>	23348	8,25E+04	100%	0.0	99.94%	1223518	NW_015961565.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4429	<i>Capsicum annuum</i>	23346	2,88E+04	100%	0.0	99.94%	158156	NW_015960693.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold90	<i>Capsicum annuum</i>	23342	3,21E+04	100%	0.0	99.94%	6074945	NW_015960458.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold516	<i>Capsicum annuum</i>	23342	5,88E+04	100%	0.0	99.93%	4299619	NW_015961344.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5219	<i>Capsicum annuum</i>	23340	3,14E+04	100%	0.0	99.93%	107289	NW_015961153.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2375	<i>Capsicum annuum</i>	23335	5,08E+04	100%	0.0	99.92%	1171512	NW_015961637.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold8691	<i>Capsicum annuum</i>	23326	2,89E+04	100%	0.0	99.91%	220109	NW_015961223.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4239	<i>Capsicum annuum</i>	23324	2,88E+04	100%	0.0	99.90%	167789	NW_015960814.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4695	<i>Capsicum annuum</i>	23322	2,88E+04	100%	0.0	99.90%	183029	NW_015961566.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold7284	<i>Capsicum annuum</i>	23315	2,88E+04	100%	0.0	99.88%	144137	NW_015961274.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2793	<i>Capsicum annuum</i>	23313	2,87E+04	100%	0.0	99.89%	203362	NW_015961784.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4365	<i>Capsicum annuum</i>	23309	3,24E+04	100%	0.0	99.87%	116290	NW_015960713.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5622	<i>Capsicum annuum</i>	23300	2,87E+04	100%	0.0	99.86%	419150	NW_015961511.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2298	<i>Capsicum annuum</i>	23287	3,23E+04	100%	0.0	99.84%	1494726	NW_015960454.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold619	<i>Capsicum annuum</i>	22990	2,93E+04	100%	0.0	99.98%	2512642	NW_015960498.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1754	<i>Capsicum annuum</i>	13821	2,69E+04	91%	0.0	99.95%	1029277	NW_015960393.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold827	<i>Capsicum annuum</i>	13821	2,68E+04	91%	0.0	99.95%	186540	NW_015960457.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold555	<i>Capsicum annuum</i>	13821	3,33E+04	91%	0.0	99.95%	1396435	NW_015960461.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold171	<i>Capsicum annuum</i>	13821	5,82E+04	91%	0.0	99.95%	887051	NW_015960473.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1963	<i>Capsicum annuum</i>	13821	3,30E+04	91%	0.0	99.95%	732782	NW_015960474.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1741	<i>Capsicum annuum</i>	13821	2,72E+04	91%	0.0	99.95%	1897442	NW_015960687.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold301	<i>Capsicum annuum</i>	13821	2,68E+04	91%	0.0	99.95%	1226190	NW_015960697.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold46	<i>Capsicum annuum</i>	13821	2,92E+04	91%	0.0	99.95%	941867	NW_015960787.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold15566	<i>Capsicum annuum</i>	13821	2,98E+04	91%	0.0	99.95%	78250	NW_015960799.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1038	<i>Capsicum annuum</i>	13821	4,91E+04	91%	0.0	99.95%	2164954	NW_015960831.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold175	<i>Capsicum annuum</i>	13821	5,37E+04	91%	0.0	99.95%	655560	NW_015960845.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	13821	8,80E+04	91%	0.0	99.95%	7059680	NW_015960855.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold433	<i>Capsicum annuum</i>	13821	2,80E+04	91%	0.0	99.95%	1277097	NW_015961081.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold578	<i>Capsicum annuum</i>	13821	4,25E+04	91%	0.0	99.95%	1213777	NW_015961091.1

GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3714	<i>Capsicum annuum</i>	13821	5,20E+04	91%	0.0	99.95%	935490	NW_015961128.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2707	<i>Capsicum annuum</i>	13821	2,68E+04	91%	0.0	99.95%	1896504	NW_015961173.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2003	<i>Capsicum annuum</i>	13821	2,68E+04	91%	0.0	99.95%	314382	NW_015961213.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2756	<i>Capsicum annuum</i>	13821	3,74E+04	91%	0.0	99.95%	64827	NW_015961263.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold660	<i>Capsicum annuum</i>	13821	2,68E+04	91%	0.0	99.95%	663463	NW_015961299.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capsicum annuum</i>	13821	3,13E+04	91%	0.0	99.95%	2397553	NW_015961318.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2901	<i>Capsicum annuum</i>	13821	2,68E+04	91%	0.0	99.95%	1161691	NW_015961380.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold933	<i>Capsicum annuum</i>	13821	3,14E+04	91%	0.0	99.95%	1853301	NW_015961435.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold454	<i>Capsicum annuum</i>	13821	2,96E+04	91%	0.0	99.95%	3392484	NW_015961509.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3928	<i>Capsicum annuum</i>	13821	3,53E+04	91%	0.0	99.95%	385536	NW_015961713.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold969	<i>Capsicum annuum</i>	13821	2,98E+04	91%	0.0	99.95%	1557263	NW_015961740.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold306	<i>Capsicum annuum</i>	13815	2,68E+04	91%	0.0	99.93%	749978	NW_015961154.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	13815	2,88E+04	91%	0.0	99.93%	2470984	NW_015961229.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2726	<i>Capsicum annuum</i>	13795	2,66E+04	91%	0.0	99.88%	212873	NW_015961432.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2102	<i>Capsicum annuum</i>	13778	2,67E+04	91%	0.0	99.84%	627640	NW_015961440.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3416	<i>Capsicum annuum</i>	13754	2,67E+04	91%	0.0	99.95%	264140	NW_015961685.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5097	<i>Capsicum annuum</i>	13581	1,96E+04	86%	0.0	99.44%	302211	NW_015961801.1
GypsyCa_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2567	<i>Capsicum annuum</i>	13224	2,75E+04	91%	0.0	99.94%	379486	NW_015961593.1
GypsyCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	21433	2,23E+09	100%	0.0	100.00%	163962470	NC_029978.1
GypsyCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	21433	3,03E+09	100%	0.0	100.00%	261510930	NC_029979.1
GypsyCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	21433	2,18E+09	100%	0.0	100.00%	215701946	NC_029980.1
GypsyCa_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	21433	2,35E+09	100%	0.0	100.00%	217274494	NC_029981.1
GypsyCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	15446	7,20E+04	100%	0.0	100.00%	229934170	NC_029988.1
GypsyCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	12013	1,12E+08	100%	0.0	92.77%	220335243	NC_029987.1
GypsyCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11570	7,26E+04	100%	0.0	91.71%	238794889	NC_029985.1
GypsyCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1339	<i>Capsicum annuum</i>	8973	1,36E+04	83%	0.0	92.59%	928404	NW_015960451.1
GypsyCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1107	<i>Capsicum annuum</i>	7197	9,68E+03	85%	0.0	87.19%	318496	NW_015961239.1
GypsyCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7047	8,22E+04	100%	0.0	86.94%	219521584	NC_029982.1
GypsyCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6715	6,45E+04	86%	0.0	88.82%	217274494	NC_029981.1
GypsyCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6637	6,80E+04	86%	0.0	86.02%	163962470	NC_029978.1
GypsyCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6617	1,06E+08	86%	0.0	85.31%	261510930	NC_029979.1
GypsyCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5734	5,12E+04	100%	0.0	84.27%	215701946	NC_029980.1
GypsyCa_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5602	2,20E+04	98%	0.0	83.61%	153299543	NC_029984.1
monotypic-Ch01e_38	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	NA	4,07E+03	100%	0.0	100.00%	301019445	NC_029977.1
monotypic-Ch01e_38	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	NA	4,07E+03	100%	0.0	90.49%	261510930	NC_029979.1
monotypic-Ch01e_38	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	NA	4,07E+03	99%	0.0	93.21%	220335243	NC_029987.1
monotypic-Ch01e_38	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	NA	4,07E+03	98%	0.0	93.08%	153299543	NC_029984.1
monotypic-Ch01e_38	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	NA	4,07E+03	93%	0.0	88.29%	238794889	NC_029985.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	25233	5,97E+08	100%	0.0	100.00%	301019445	NC_029977.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11601	4,93E+08	100%	0.0	85.24%	229934170	NC_029988.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11520	3,83E+08	100%	0.0	82.06%	222112641	NC_029983.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11334	4,16E+08	100%	0.0	81.84%	205736368	NC_029986.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11241	5,19E+08	100%	0.0	82.91%	238794889	NC_029985.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9391	4,33E+08	100%	0.0	82.79%	220335243	NC_029987.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7474	6,28E+08	100%	0.0	81.02%	261510930	NC_029979.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6903	3,12E+08	100%	0.0	84.19%	153299543	NC_029984.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold814	<i>Capsicum annuum</i>	6889	1,47E+04	89%	0.0	81.72%	1128876	NW_015960740.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6852	3,05E+08	100%	0.0	84.48%	163962470	NC_029978.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6782	3,93E+08	100%	0.0	81.20%	219521584	NC_029982.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6410	4,56E+08	100%	0.0	82.44%	215701946	NC_029980.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3339	<i>Capsicum annuum</i>	5853	1,42E+04	94%	0.0	82.93%	1234997	NW_015960491.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold90	<i>Capsicum annuum</i>	5773	2,99E+04	100%	0.0	78.63%	6074945	NW_015960458.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5760	4,37E+08	100%	0.0	81.05%	217274494	NC_029981.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1664	<i>Capsicum annuum</i>	4929	1,31E+04	98%	0.0	81.56%	131442	NW_015960953.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold658	<i>Capsicum annuum</i>	3397	1,19E+04	84%	0.0	80.81%	613851	NW_015961079.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold516	<i>Capsicum annuum</i>	3380	1,80E+04	89%	0.0	80.26%	4299619	NW_015961344.1

monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold21075	<i>Capsicum annuum</i>	3262	8,64E+03	81%	0.0	80.56%	40025	NW_015961234.1
monotypic-Ch01f_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	2955	2,32E+04	82%	0.0	83.34%	7059680	NW_015960855.1
monotypic-Ch01h_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	30498	6,13E+08	100%	0.0	100.00%	301019445	NC_029977.1
monotypic-Ch01h_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5945	3,74E+08	94%	0.0	83.91%	222112641	NC_029983.1
monotypic-Ch01h_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5792	4,43E+08	94%	0.0	84.12%	219521584	NC_029982.1
monotypic-Ch01h_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5690	5,06E+08	99%	0.0	84.36%	229934170	NC_029988.1
monotypic-Ch01h_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5631	3,84E+08	99%	0.0	83.05%	163962470	NC_029978.1
monotypic-Ch01h_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5293	5,08E+08	96%	0.0	82.05%	238794889	NC_029985.1
monotypic-Ch01h_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5252	3,96E+08	93%	0.0	82.71%	220335243	NC_029987.1
monotypic-Ch01h_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5188	4,59E+08	94%	0.0	82.44%	215701946	NC_029980.1
monotypic-Ch01h_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5035	4,20E+08	96%	0.0	81.89%	205736368	NC_029986.1
monotypic-Ch01h_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4946	4,74E+08	99%	0.0	81.84%	217274494	NC_029981.1
monotypic-Ch01h_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4924	4,07E+08	94%	0.0	81.32%	261510930	NC_029979.1
monotypic-Ch01h_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4676	4,16E+08	93%	0.0	80.37%	153299543	NC_029984.1
monotypic-Ch01h_17	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	3219	3,24E+04	89%	0.0	85.13%	7059680	NW_015960855.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	18526	7,28E+08	100%	0.0	99.94%	163962470	NC_029978.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11889	1,00E+09	100%	0.0	88.01%	217274494	NC_029981.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8990	9,61E+08	100%	0.0	83.00%	229934170	NC_029988.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8383	9,68E+08	100%	0.0	83.25%	219521584	NC_029982.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8043	1,38E+09	100%	0.0	83.69%	301019445	NC_029977.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7664	6,69E+08	100%	0.0	81.56%	153299543	NC_029984.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7574	1,26E+09	100%	0.0	84.01%	261510930	NC_029979.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1651	<i>Capsicum annuum</i>	7424	1,11E+04	97%	0.0	84.09%	331682	NW_015961206.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7422	9,13E+08	100%	0.0	83.57%	215701946	NC_029980.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7406	1,24E+09	100%	0.0	80.93%	222112641	NC_029983.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7385	1,08E+09	100%	0.0	84.73%	238794889	NC_029985.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1118	<i>Capsicum annuum</i>	7234	8,81E+03	88%	0.0	82.97%	528969	NW_015960738.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1923	<i>Capsicum annuum</i>	7197	3,15E+04	100%	0.0	80.71%	2480791	NW_015961228.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7153	8,27E+08	100%	0.0	79.77%	220335243	NC_029987.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold296	<i>Capsicum annuum</i>	6983	1,31E+04	91%	0.0	84.12%	1995018	NW_015961641.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold454	<i>Capsicum annuum</i>	6968	2,53E+04	99%	0.0	83.01%	3392484	NW_015961509.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6961	1,03E+09	100%	0.0	81.58%	205736368	NC_029986.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold303	<i>Capsicum annuum</i>	6913	6,91E+03	100%	0.0	79.39%	285559	NW_015960893.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1390	<i>Capsicum annuum</i>	6878	1,13E+04	100%	0.0	83.47%	1730405	NW_015960783.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold985	<i>Capsicum annuum</i>	6855	4,13E+04	100%	0.0	80.19%	1349800	NW_015961816.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold349	<i>Capsicum annuum</i>	6839	2,22E+04	99%	0.0	79.28%	2130419	NW_015960729.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9337	<i>Capsicum annuum</i>	6711	9,07E+03	94%	0.0	80.47%	397715	NW_015961734.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1124	<i>Capsicum annuum</i>	6671	8,17E+04	100%	0.0	81.08%	2110277	NW_015961370.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3202	<i>Capsicum annuum</i>	6637	9,71E+03	88%	0.0	82.49%	302159	NW_015961119.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3611	<i>Capsicum annuum</i>	6421	7,26E+03	91%	0.0	81.74%	177705	NW_015960430.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2451	<i>Capsicum annuum</i>	6318	1,08E+04	94%	0.0	82.11%	364752	NW_015961179.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2157	<i>Capsicum annuum</i>	6309	8,76E+03	95%	0.0	82.05%	158560	NW_015960692.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4747	<i>Capsicum annuum</i>	6285	1,72E+04	100%	0.0	78.38%	1151343	NW_015960460.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold669	<i>Capsicum annuum</i>	6144	1,28E+04	96%	0.0	80.72%	640007	NW_015960674.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold90	<i>Capsicum annuum</i>	6069	3,88E+04	100%	0.0	81.41%	6074945	NW_015960458.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold516	<i>Capsicum annuum</i>	5655	3,58E+04	100%	0.0	82.02%	4296919	NW_015961344.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	5528	1,65E+04	92%	0.0	78.14%	3311724	NW_015960635.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3714	<i>Capsicum annuum</i>	5494	9,26E+03	90%	0.0	82.70%	935490	NW_015961128.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1038	<i>Capsicum annuum</i>	5480	2,06E+04	99%	0.0	83.53%	2164954	NW_015960831.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	5422	5,20E+04	99%	0.0	81.13%	7059680	NW_015960855.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold983	<i>Capsicum annuum</i>	5393	8,60E+03	99%	0.0	79.95%	627343	NW_015961594.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2102	<i>Capsicum annuum</i>	5216	6,15E+03	82%	0.0	80.57%	627640	NW_015961440.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold438	<i>Capsicum annuum</i>	5051	1,18E+04	100%	0.0	80.35%	1337825	NW_015961507.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold969	<i>Capsicum annuum</i>	4992	1,58E+04	99%	0.0	83.97%	1557263	NW_015961740.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold189	<i>Capsicum annuum</i>	4957	1,99E+04	99%	0.0	78.07%	3791111	NW_015960691.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3026	<i>Capsicum annuum</i>	4865	1,24E+04	90%	0.0	82.06%	1386929	NW_015961102.1

monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold170	<i>Capsicum annuum</i>	4813	3,17E+04	100%	0.0	79.70%	2217302	NW_015960390.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold933	<i>Capsicum annuum</i>	4763	1,86E+04	91%	0.0	79.58%	1853301	NW_015961435.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold24	<i>Capsicum annuum</i>	4706	6,49E+03	98%	0.0	78.53%	1258456	NW_015961700.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2707	<i>Capsicum annuum</i>	4639	1,45E+04	83%	0.0	78.32%	1896504	NW_015961173.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1341	<i>Capsicum annuum</i>	4551	1,83E+04	100%	0.0	82.91%	1906221	NW_015960665.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold619	<i>Capsicum annuum</i>	4379	8,20E+03	81%	0.0	79.06%	2512642	NW_015960498.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	4372	1,29E+04	94%	0.0	80.68%	2470984	NW_015961229.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1246	<i>Capsicum annuum</i>	4331	1,59E+04	91%	0.0	81.02%	3388803	NW_015960834.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3843	<i>Capsicum annuum</i>	4161	1,18E+04	99%	0.0	80.30%	465776	NW_015961174.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold231	<i>Capsicum annuum</i>	4126	6,96E+03	81%	0.0	78.01%	1482909	NW_015961568.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3499	<i>Capsicum annuum</i>	4076	1,21E+04	99%	0.0	85.49%	681687	NW_015960538.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1973	<i>Capsicum annuum</i>	3567	1,15E+04	96%	0.0	82.70%	1714326	NW_015961422.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold578	<i>Capsicum annuum</i>	3229	1,60E+04	94%	0.0	81.00%	1213777	NW_015961091.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold188	<i>Capsicum annuum</i>	3046	4,48E+03	80%	0.0	76.73%	980346	NW_015960557.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9	<i>Capsicum annuum</i>	3037	2,21E+04	100%	0.0	76.31%	6567951	NW_015961815.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2298	<i>Capsicum annuum</i>	2839	1,07E+04	92%	0.0	78.62%	1494726	NW_015960454.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1741	<i>Capsicum annuum</i>	2728	1,42E+04	100%	0.0	83.65%	1897442	NW_015960687.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold46	<i>Capsicum annuum</i>	2643	5,99E+03	88%	0.0	76.63%	941867	NW_015960787.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capsicum annuum</i>	2351	6,80E+03	81%	0.0	77.89%	2397553	NW_015961318.1
monotypic-Ch02c_27	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold6075	<i>Capsicum annuum</i>	2350	1,05E+04	94%	0.0	84.23%	316563	NW_015960475.1
monotypic-Ch02e_84	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9441	1,82E+04	100%	0.0	100.00%	163962470	NC_029978.1
monotypic-Ch02e_84	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3101	8,54E+03	92%	0.0	94.83%	261510930	NC_029979.1
monotypic-Ch02e_84	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2612	7,29E+03	83%	0.0	90.59%	301019445	NC_029977.1
monotypic-Ch03a_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10048	2,10E+04	100%	0.0	100.00%	261510930	NC_029979.1
monotypic-Ch03a_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7145	9,91E+03	98%	0.0	95.37%	222112641	NC_029983.1
monotypic-Ch03a_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6900	3,34E+04	100%	0.0	93.34%	217274494	NC_029981.1
monotypic-Ch03a_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6833	9,17E+03	100%	0.0	93.15%	238794889	NC_029985.1
monotypic-Ch03a_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5175	1,10E+04	96%	0.0	95.21%	301019445	NC_029977.1
monotypic-Ch03a_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5007	9,49E+03	100%	0.0	94.14%	229934170	NC_029988.1
monotypic-Ch03a_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3967	8,69E+03	98%	0.0	91.99%	220335243	NC_029987.1
monotypic-Ch03a_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10033	2,99E+04	100%	0.0	100.00%	261510930	NC_029979.1
monotypic-Ch03a_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8327	1,94E+04	100%	0.0	94.38%	153299543	NC_029984.1
monotypic-Ch03a_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8041	2,55E+04	100%	0.0	93.42%	205736368	NC_029986.1
monotypic-Ch03a_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7875	9,36E+03	100%	0.0	92.86%	229934170	NC_029988.1
monotypic-Ch03a_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7681	8,97E+03	100%	0.0	92.22%	301019445	NC_029977.1
monotypic-Ch03a_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5089	<i>Capsicum annuum</i>	7396	9,10E+03	100%	0.0	93.04%	522036	NW_015961006.1
monotypic-Ch03a_23	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7382	8,67E+03	100%	0.0	91.27%	219521584	NC_029982.1
monotypic-Ch03a_23	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00019	<i>Solanum tuberosum</i>	3099	3,10E+03	100%	0.0	77.29%	3257788	NW_006238945.1
monotypic-Ch03a_23	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00002	<i>Solanum tuberosum</i>	3096	3,10E+03	100%	0.0	77.28%	6551949	NW_006238928.1
monotypic-Ch03a_41	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10054	1,75E+04	100%	0.0	100.00%	261510930	NC_029979.1
monotypic-Ch03a_41	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8035	8,97E+03	100%	0.0	93.41%	205736368	NC_029986.1
monotypic-Ch03a_41	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7914	8,67E+03	100%	0.0	93.05%	222112641	NC_029983.1
monotypic-Ch03a_41	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6682	6,68E+03	96%	0.0	89.50%	220335243	NC_029987.1
monotypic-Ch03a_41	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5249	8,46E+03	100%	0.0	93.66%	163962470	NC_029978.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23941	2,86E+08	100%	0.0	100.00%	261510930	NC_029979.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10113	3,64E+08	100%	0.0	90.62%	217274494	NC_029981.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8794	2,86E+08	100%	0.0	87.54%	229934170	NC_029988.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8503	2,68E+08	100%	0.0	86.92%	220335243	NC_029987.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold277	<i>Capsicum annuum</i>	6970	1,97E+04	100%	0.0	89.50%	448122	NW_015961390.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6623	2,84E+08	100%	0.0	88.37%	153299543	NC_029984.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6516	2,02E+08	100%	0.0	88.80%	222112641	NC_029983.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6433	2,08E+08	99%	0.0	88.50%	219521584	NC_029982.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	6115	1,88E+04	97%	0.0	87.50%	3311724	NW_015960635.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6095	3,83E+08	100%	0.0	87.40%	301019445	NC_029977.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6074	2,17E+08	100%	0.0	87.29%	205736368	NC_029986.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold985	<i>Capsicum annuum</i>	6004	1,89E+04	99%	0.0	87.12%	1349800	NW_015961816.1

monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	5947	1,43E+04	87%	0.0	86.83%	3493078	NW_015960517.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold891	<i>Capsicum annuum</i>	5873	1,87E+04	94%	0.0	86.59%	1431046	NW_015960469.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5853	1,85E+08	98%	0.0	86.61%	163962470	NC_029978.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold949	<i>Capsicum annuum</i>	5812	1,58E+04	92%	0.0	86.38%	3128562	NW_015961487.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5716	2,52E+08	100%	0.0	85.48%	215701946	NC_029980.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5317	2,84E+08	95%	0.0	87.03%	238794889	NC_029985.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold189	<i>Capsicum annuum</i>	4584	1,74E+04	93%	0.0	84.63%	3791111	NW_015960691.1
monotypic-Ch03b_33	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3339	<i>Capsicum annuum</i>	2394	1,21E+04	88%	0.0	87.06%	1234997	NW_015960491.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23148	8,89E+08	100%	0.0	100.00%	261510930	NC_029979.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	16643	6,83E+08	100%	0.0	90.62%	215701946	NC_029980.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	16292	6,02E+08	100%	0.0	91.08%	229934170	NC_029988.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	15998	7,26E+08	100%	0.0	89.72%	205736368	NC_029986.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold742	<i>Capsicum annuum</i>	15813	2,90E+04	100%	0.0	89.49%	2382255	NW_015961471.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	13786	4,74E+08	100%	0.0	88.42%	153299543	NC_029984.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	12990	7,82E+08	100%	0.0	89.85%	217274494	NC_029981.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	12728	9,07E+08	100%	0.0	89.19%	301019445	NC_029977.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10944	1,14E+09	100%	0.0	89.41%	238794889	NC_029985.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10536	6,22E+08	100%	0.0	91.26%	220335243	NC_029987.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9747	4,13E+08	100%	0.0	90.18%	163962470	NC_029978.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9646	6,80E+08	100%	0.0	88.82%	222112641	NC_029983.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9023	5,84E+08	100%	0.0	89.27%	219521584	NC_029982.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	8589	1,80E+04	94%	0.0	86.46%	3311724	NW_015960635.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	7978	7,82E+04	98%	0.0	86.85%	7059680	NW_015960855.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2707	<i>Capsicum annuum</i>	7967	1,83E+04	98%	0.0	88.02%	1896504	NW_015961173.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold519	<i>Capsicum annuum</i>	5899	1,37E+04	86%	0.0	85.83%	1101621	NW_015961578.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1587	<i>Capsicum annuum</i>	5864	1,21E+04	81%	0.0	88.32%	1400685	NW_015961196.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1038	<i>Capsicum annuum</i>	5753	1,82E+04	92%	0.0	87.63%	2164954	NW_015960831.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1620	<i>Capsicum annuum</i>	4972	1,29E+04	89%	0.0	85.66%	500299	NW_015961331.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2697	<i>Capsicum annuum</i>	4876	2,15E+04	86%	0.0	86.52%	139253	NW_015961756.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	4854	1,73E+04	89%	0.0	85.78%	3493078	NW_015960517.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold454	<i>Capsicum annuum</i>	4767	2,52E+04	93%	0.0	86.54%	3392484	NW_015961509.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1074	<i>Capsicum annuum</i>	4277	2,05E+04	93%	0.0	88.03%	1095636	NW_015961132.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4204	<i>Capsicum annuum</i>	4259	2,66E+04	93%	0.0	85.04%	323636	NW_015960601.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold603	<i>Capsicum annuum</i>	4237	1,82E+04	95%	0.0	79.69%	1800753	NW_015960521.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2298	<i>Capsicum annuum</i>	4060	1,59E+04	92%	0.0	88.08%	1494726	NW_015960454.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3151	<i>Capsicum annuum</i>	3964	1,35E+04	92%	0.0	88.22%	198665	NW_015960810.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1307	<i>Capsicum annuum</i>	3773	1,38E+04	93%	0.0	87.68%	1549638	NW_015960466.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold983	<i>Capsicum annuum</i>	3240	1,66E+04	89%	0.0	88.42%	627343	NW_015961594.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold846	<i>Capsicum annuum</i>	3000	9,64E+03	83%	0.0	82.47%	1232610	NW_015960818.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1923	<i>Capsicum annuum</i>	2407	1,62E+04	86%	0.0	87.23%	2480791	NW_015961228.1
monotypic-Ch03e_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	1927	1,23E+04	82%	0.0	89.18%	2470984	NW_015961229.1
monotypic-Ch03g_96	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9640	1,06E+04	100%	0.0	100.00%	261510930	NC_029979.1
monotypic-Ch03g_96	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6658	2,08E+04	99%	0.0	90.91%	219521584	NC_029982.1
monotypic-Ch03g_96	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6089	1,24E+04	99%	0.0	89.63%	301019445	NC_029977.1
monotypic-Ch03g_96	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5598	9,32E+03	99%	0.0	92.05%	238794889	NC_029985.1
monotypic-Ch03g_96	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2516	6,98E+03	95%	0.0	89.65%	220335243	NC_029987.1
monotypic-Ch03g_96	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold7892	<i>Capsicum annuum</i>	2447	2,97E+03	97%	0.0	76.51%	83640	NW_015961298.1
monotypic-Ch03g_96	refeq-genome	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	<i>Solanum lycopersicum</i>	1349	1,35E+03	80%	0.0	72.99%	68126176	NC_015449.3
monotypic-Ch04a_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9564	1,23E+04	100%	0.0	100.00%	215701946	NC_029980.1
monotypic-Ch04a_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6104	3,04E+04	100%	0.0	87.97%	163962470	NC_029978.1
monotypic-Ch04a_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5984	3,41E+04	100%	0.0	87.54%	301019445	NC_029977.1
monotypic-Ch04a_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5500	8,07E+03	99%	0.0	90.32%	222112641	NC_029983.1
monotypic-Ch04a_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5337	1,13E+04	100%	0.0	85.86%	217274494	NC_029981.1
monotypic-Ch04a_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3568	1,25E+04	100%	0.0	91.94%	220335243	NC_029987.1
monotypic-Ch04a_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3474	6,73E+03	99%	0.0	84.15%	261510930	NC_029979.1
monotypic-Ch04a_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold891	<i>Capsicum annuum</i>	3162	6,43E+03	99%	0.0	88.23%	1431046	NW_015960469.1

monotypic-Ch04a_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2940	5,00E+03	81%	0.0	88.28%	153299543	NC_029984.1
monotypic-Ch04a_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	2699	7,02E+03	98%	0.0	86.61%	238794889	NC_029985.1
monotypic-Ch04b_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9684	1,40E+04	100%	0.0	99.98%	215701946	NC_029980.1
monotypic-Ch04b_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6399	3,74E+04	100%	0.0	90.91%	229934170	NC_029988.1
monotypic-Ch04b_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5827	8,14E+03	100%	0.0	90.65%	163962470	NC_029978.1
monotypic-Ch04b_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5511	7,37E+03	99%	0.0	89.46%	222112641	NC_029983.1
monotypic-Ch04b_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4381	1,46E+04	100%	0.0	90.49%	261510930	NC_029979.1
monotypic-Ch04b_28	refeq-genome	8	Nicotiana attenuata strain UT chromosome 11, NIATTr2	<i>Nicotiana attenuata</i>	4355	4,36E+03	98%	0.0	82.15%	61252062	NC_031999.1
monotypic-Ch04b_28	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold25671	<i>Nicotiana tomentosiformis</i>	3768	3,77E+03	88%	0.0	81.47%	242328	NW_008905454.1
monotypic-Ch04b_28	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold174119	<i>Nicotiana tabacum</i>	3757	3,76E+03	88%	0.0	81.43%	25118	NW_015837501.1
monotypic-Ch04b_28	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsy Nsy_scaffold29596	<i>Nicotiana sylvestris</i>	3661	3,66E+03	84%	0.0	81.84%	35257	NW_009514493.1
monotypic-Ch04b_28	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold102920	<i>Nicotiana tabacum</i>	3633	3,63E+03	84%	0.0	81.73%	27530	NW_015789514.1
monotypic-Ch04b_28	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold70857	<i>Nicotiana tabacum</i>	3620	3,62E+03	95%	0.0	79.97%	59007	NW_015931337.1
monotypic-Ch04b_28	refeq-genome	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold10355	<i>Nicotiana tomentosiformis</i>	3618	3,62E+03	95%	0.0	79.96%	83421	NW_008832438.1
monotypic-Ch04b_28	refeq-genome	8	Nicotiana sylvestris unplaced genomic scaffold, Nsy Nsy_scaffold12006	<i>Nicotiana sylvestris</i>	3615	4,21E+03	90%	0.0	83.44%	114056	NW_009361093.1
monotypic-Ch04b_28	refeq-genome	8	Nicotiana attenuata strain UT chromosome 6, NIATTr2	<i>Nicotiana attenuata</i>	3531	3,53E+03	84%	0.0	81.47%	97571839	NC_031994.1
monotypic-Ch04b_28	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold6534	<i>Nicotiana tabacum</i>	3360	3,36E+03	82%	0.0	80.97%	20057	NW_015926529.1
monotypic-Ch04b_28	refeq-genome	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	<i>Nicotiana attenuata</i>	2148	3,46E+03	82%	0.0	82.53%	466898	NW_017670751.1
monotypic-Ch04b_28	refeq-genome	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13981	<i>Nicotiana tabacum</i>	1906	3,96E+03	92%	0.0	80.39%	156831	NW_015816232.1
monotypic-Ch04b_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold454	<i>Capsicum annuum</i>	1766	4,58E+03	81%	0.0	86.81%	3392484	NW_015961509.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	18362	5,47E+08	100%	0.0	100.00%	215701946	NC_029980.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5097	<i>Capsicum annuum</i>	11114	1,48E+04	100%	0.0	86.95%	302211	NW_015961801.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10992	5,59E+08	100%	0.0	86.69%	229934170	NC_029988.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10918	6,38E+08	100%	0.0	88.02%	205736368	NC_029986.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10811	5,68E+08	100%	0.0	86.41%	217274494	NC_029981.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10595	6,22E+08	100%	0.0	87.38%	238794889	NC_029985.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8900	8,05E+08	100%	0.0	87.40%	301019445	NC_029977.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8863	5,48E+08	100%	0.0	87.22%	219521584	NC_029982.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8320	7,63E+08	100%	0.0	86.90%	222112641	NC_029983.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1341	<i>Capsicum annuum</i>	8237	1,74E+04	100%	0.0	88.50%	1906221	NW_015960665.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold438	<i>Capsicum annuum</i>	8019	1,25E+04	100%	0.0	83.72%	1337825	NW_015961507.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7234	4,01E+08	100%	0.0	82.41%	153299543	NC_029984.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold170	<i>Capsicum annuum</i>	7079	1,78E+04	95%	0.0	82.08%	2217302	NW_015960390.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold933	<i>Capsicum annuum</i>	6979	2,48E+04	100%	0.0	81.87%	1853301	NW_015961435.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3202	<i>Capsicum annuum</i>	6813	9,14E+03	91%	0.0	81.78%	302159	NW_015961119.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6807	4,95E+08	100%	0.0	81.68%	220335243	NC_029987.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2102	<i>Capsicum annuum</i>	6433	1,15E+04	99%	0.0	84.33%	627640	NW_015961440.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1118	<i>Capsicum annuum</i>	6377	9,38E+03	99%	0.0	82.41%	528969	NW_015960738.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6329	6,12E+08	100%	0.0	81.76%	261510930	NC_029979.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold619	<i>Capsicum annuum</i>	6189	1,11E+04	90%	0.0	81.03%	2512642	NW_015960498.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold90	<i>Capsicum annuum</i>	5908	3,00E+04	99%	0.0	79.85%	6074945	NW_015960458.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5710	3,93E+08	100%	0.0	80.79%	163962470	NC_029978.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	5469	1,42E+04	91%	0.0	83.99%	2470984	NW_015961229.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1124	<i>Capsicum annuum</i>	5398	5,13E+04	100%	0.0	80.04%	2110277	NW_015961370.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1038	<i>Capsicum annuum</i>	5251	1,02E+04	86%	0.0	82.13%	2164954	NW_015960831.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold516	<i>Capsicum annuum</i>	5208	2,22E+04	99%	0.0	79.98%	4299619	NW_015961344.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold231	<i>Capsicum annuum</i>	5208	7,72E+03	88%	0.0	80.01%	1482909	NW_015961568.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold669	<i>Capsicum annuum</i>	5197	1,29E+04	99%	0.0	80.49%	640007	NW_015960674.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1651	<i>Capsicum annuum</i>	4390	6,23E+03	86%	0.0	80.10%	331682	NW_015961206.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold296	<i>Capsicum annuum</i>	4244	6,26E+03	82%	0.0	79.58%	1995018	NW_015961641.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	4194	2,79E+04	100%	0.0	80.54%	7059680	NW_015960855.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold454	<i>Capsicum annuum</i>	4193	1,52E+04	90%	0.0	79.38%	3392484	NW_015961509.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1390	<i>Capsicum annuum</i>	4191	6,20E+03	83%	0.0	79.42%	1730405	NW_015960783.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold985	<i>Capsicum annuum</i>	4006	1,73E+04	87%	0.0	77.34%	1349800	NW_015961816.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2451	<i>Capsicum annuum</i>	3925	7,26E+03	84%	0.0	78.59%	364752	NW_015961179.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold578	<i>Capsicum annuum</i>	3775	1,58E+04	100%	0.0	83.30%	1213777	NW_015961091.1

monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold349	<i>Capsicum annuum</i>	3733	1,02E+04	88%	0.0	76.76%	2130419	NW_015960729.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3026	<i>Capsicum annuum</i>	3701	7,60E+03	86%	0.0	78.59%	1386929	NW_015961102.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2375	<i>Capsicum annuum</i>	3531	9,25E+03	88%	0.0	79.75%	1171512	NW_015961637.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9	<i>Capsicum annuum</i>	3522	2,12E+04	99%	0.0	81.55%	6567951	NW_015961815.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2707	<i>Capsicum annuum</i>	3367	1,09E+04	94%	0.0	76.96%	1896504	NW_015961173.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1246	<i>Capsicum annuum</i>	3138	8,85E+03	85%	0.0	80.01%	3388803	NW_015960834.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capsicum annuum</i>	3029	1,03E+04	89%	0.0	79.72%	2397553	NW_015961318.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold189	<i>Capsicum annuum</i>	2998	1,21E+04	94%	0.0	76.69%	3791111	NW_015960691.1
monotypic-Ch04e_05	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1973	<i>Capsicum annuum</i>	2850	6,72E+03	84%	0.0	79.63%	1714326	NW_015961422.1
monotypic-Ch05a_58	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8144	2,88E+04	100%	0.0	100.00%	217274494	NC_029981.1
monotypic-Ch05a_58	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6303	1,02E+04	98%	0.0	92.96%	222112641	NC_029983.1
monotypic-Ch05a_58	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6144	1,25E+04	86%	0.0	92.91%	261510930	NC_029979.1
monotypic-Ch05a_58	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6013	9,04E+03	100%	0.0	91.31%	238794889	NC_029985.1
monotypic-Ch05a_58	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4087	1,04E+04	96%	0.0	92.84%	301019445	NC_029977.1
monotypic-Ch05a_58	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3949	6,30E+03	86%	0.0	91.76%	229934170	NC_029988.1
monotypic-Ch05a_58	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3173	5,79E+03	84%	0.0	90.79%	220335243	NC_029987.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	15034	8,16E+08	100%	0.0	100.00%	217274494	NC_029981.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9928	8,47E+08	100%	0.0	88.78%	301019445	NC_029977.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9897	7,95E+08	100%	0.0	89.91%	222112641	NC_029983.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8990	8,40E+08	100%	0.0	86.66%	261510930	NC_029979.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8966	7,66E+08	100%	0.0	86.59%	229934170	NC_029988.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8896	7,04E+08	100%	0.0	87.19%	220335243	NC_029987.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8761	8,49E+08	100%	0.0	86.93%	215701946	NC_029980.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8696	6,48E+08	100%	0.0	88.53%	238794889	NC_029985.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold979	<i>Capsicum annuum</i>	8497	1,52E+04	98%	0.0	88.54%	1172749	NW_015961157.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2343	<i>Capsicum annuum</i>	8456	1,15E+04	92%	0.0	87.01%	440837	NW_015961556.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3339	<i>Capsicum annuum</i>	8366	1,07E+04	99%	0.0	85.38%	1234997	NW_015960491.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4043	<i>Capsicum annuum</i>	8200	1,39E+04	98%	0.0	85.42%	396497	NW_015960527.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8196	6,35E+08	100%	0.0	85.01%	163962470	NC_029978.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold90	<i>Capsicum annuum</i>	8113	9,72E+04	100%	0.0	84.82%	6074945	NW_015960458.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8041	6,93E+08	100%	0.0	84.62%	219521584	NC_029982.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8010	5,18E+08	100%	0.0	86.46%	153299543	NC_029984.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1754	<i>Capsicum annuum</i>	7987	1,02E+04	100%	0.0	86.60%	1029277	NW_015960393.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold189	<i>Capsicum annuum</i>	7718	1,67E+04	99%	0.0	84.43%	3791111	NW_015960691.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold181	<i>Capsicum annuum</i>	7631	1,68E+04	90%	0.0	85.51%	2703691	NW_015961388.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7594	5,36E+08	100%	0.0	85.52%	205736368	NC_029986.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	7483	3,11E+04	100%	0.0	87.32%	3311724	NW_015960635.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2189	<i>Capsicum annuum</i>	7402	5,33E+04	100%	0.0	85.00%	1569264	NW_015960648.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold296	<i>Capsicum annuum</i>	6896	6,90E+03	99%	0.0	82.30%	1995018	NW_015961411.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	6885	1,19E+04	99%	0.0	86.90%	2470984	NW_015961229.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2791	<i>Capsicum annuum</i>	6835	9,10E+03	99%	0.0	82.05%	158707	NW_015961540.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	6759	4,08E+04	100%	0.0	82.37%	3493078	NW_015960517.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3675	<i>Capsicum annuum</i>	6684	2,82E+04	100%	0.0	91.38%	103359	NW_015960497.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold619	<i>Capsicum annuum</i>	6527	1,43E+04	97%	0.0	81.88%	2512642	NW_015960498.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	6322	9,83E+03	100%	0.0	83.56%	624013	NW_015961724.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2791	<i>Capsicum annuum</i>	6013	6,01E+03	89%	0.0	81.79%	196436	NW_015961542.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	5808	5,89E+03	84%	0.0	82.06%	615095	NW_015960816.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1341	<i>Capsicum annuum</i>	5720	5,72E+03	100%	0.0	79.62%	1906221	NW_015960665.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4183	<i>Capsicum annuum</i>	5306	2,15E+04	97%	0.0	83.04%	1223518	NW_015961565.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9	<i>Capsicum annuum</i>	4436	6,25E+03	86%	0.0	84.10%	6567951	NW_015961815.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4118	<i>Capsicum annuum</i>	4176	1,21E+04	81%	0.0	83.35%	593898	NW_015961463.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold949	<i>Capsicum annuum</i>	4108	1,06E+04	90%	0.0	83.05%	3128562	NW_015961487.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold742	<i>Capsicum annuum</i>	4087	9,66E+03	99%	0.0	80.22%	2382255	NW_015961471.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	3999	1,15E+04	89%	0.0	85.54%	7059680	NW_015960855.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold933	<i>Capsicum annuum</i>	3964	6,22E+03	99%	0.0	78.15%	1853301	NW_015961435.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1336	<i>Capsicum annuum</i>	3637	5,52E+03	84%	0.0	80.08%	463180	NW_015961656.1

monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold7054	<i>Capsicum annuum</i>	2959	7,24E+03	98%	0.0	84.21%	311178	NW_015960778.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1312	<i>Capsicum annuum</i>	2569	2,53E+04	100%	0.0	86.15%	140791	NW_015960923.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold170	<i>Capsicum annuum</i>	2285	4,66E+03	80%	0.0	76.48%	2217302	NW_015960390.1
monotypic-Ch05d_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4063	<i>Capsicum annuum</i>	2043	6,42E+03	91%	0.0	84.40%	1657713	NW_015960826.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23996	3,02E+08	100%	0.0	100.00%	217274494	NC_029981.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	17167	2,44E+08	100%	0.0	90.66%	229934170	NC_029988.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7834	3,17E+08	100%	0.0	85.73%	301019445	NC_029977.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7769	1,51E+08	100%	0.0	85.49%	222112641	NC_029983.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7709	2,17E+08	100%	0.0	83.60%	261510930	NC_029979.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7686	2,18E+08	98%	0.0	84.78%	220335243	NC_029987.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold277	<i>Capsicum annuum</i>	7437	1,55E+04	100%	0.0	85.33%	448122	NW_015961390.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7334	1,55E+08	98%	0.0	84.44%	219521584	NC_029982.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7221	2,25E+08	100%	0.0	84.34%	153299543	NC_029984.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold985	<i>Capsicum annuum</i>	7182	1,76E+04	95%	0.0	84.24%	1349800	NW_015961816.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	7134	1,28E+04	92%	0.0	84.07%	3493078	NW_015960517.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7060	1,73E+08	98%	0.0	83.80%	205736368	NC_029986.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	7033	1,40E+04	94%	0.0	83.92%	3311724	NW_015960635.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold949	<i>Capsicum annuum</i>	6951	1,27E+04	82%	0.0	83.63%	3128562	NW_015961487.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6874	1,64E+08	95%	0.0	83.51%	163962470	NC_029978.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6259	2,02E+08	98%	0.0	82.00%	215701946	NC_029980.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6187	2,36E+08	94%	0.0	81.81%	238794889	NC_029985.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	3225	1,94E+04	85%	0.0	84.67%	7059680	NW_015960855.1
monotypic-Ch05e_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3339	<i>Capsicum annuum</i>	2730	9,25E+03	80%	0.0	80.34%	1234997	NW_015960491.1
monotypic-Ch05e_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9766	3,33E+04	100%	0.0	100.00%	217274494	NC_029981.1
monotypic-Ch05e_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7443	8,75E+03	100%	0.0	92.17%	238794889	NC_029985.1
monotypic-Ch05e_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6900	1,78E+04	100%	0.0	93.34%	261510930	NC_029979.1
monotypic-Ch05e_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6889	9,10E+03	98%	0.0	93.68%	222112641	NC_029983.1
monotypic-Ch05e_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4907	1,03E+04	96%	0.0	93.81%	301019445	NC_029977.1
monotypic-Ch05e_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4783	8,92E+03	100%	0.0	93.00%	229934170	NC_029988.1
monotypic-Ch05e_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3807	7,77E+03	100%	0.0	91.05%	220335243	NC_029987.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23368	3,92E+08	100%	0.0	100.00%	217274494	NC_029981.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8903	4,53E+08	100%	0.0	89.12%	222112641	NC_029983.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8530	5,86E+08	100%	0.0	88.20%	301019445	NC_029977.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold6454	<i>Capsicum annuum</i>	8493	1,75E+04	99%	0.0	88.06%	84551	NW_015961449.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8407	4,58E+08	100%	0.0	87.86%	215701946	NC_029980.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8078	3,92E+08	100%	0.0	87.28%	229934170	NC_029988.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8006	3,53E+08	100%	0.0	86.89%	153299543	NC_029984.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold454	<i>Capsicum annuum</i>	7854	2,00E+04	99%	0.0	86.46%	3392484	NW_015961509.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7805	5,02E+08	100%	0.0	86.91%	261510930	NC_029979.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7803	3,44E+08	100%	0.0	86.33%	219521584	NC_029982.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7733	3,38E+08	100%	0.0	86.36%	220335243	NC_029987.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7589	2,34E+08	100%	0.0	85.76%	163962470	NC_029978.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7566	4,29E+08	100%	0.0	85.76%	238794889	NC_029985.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold985	<i>Capsicum annuum</i>	5321	1,51E+04	85%	0.0	86.81%	1349800	NW_015961816.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4466	3,30E+08	100%	0.0	86.90%	205736368	NC_029986.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold603	<i>Capsicum annuum</i>	4257	1,03E+04	86%	0.0	77.65%	1800753	NW_015960521.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4183	<i>Capsicum annuum</i>	4237	1,27E+04	97%	0.0	78.61%	1223518	NW_015961565.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2202	<i>Capsicum annuum</i>	4128	1,67E+04	99%	0.0	88.41%	213497	NW_015960401.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold640	<i>Capsicum annuum</i>	3624	1,85E+04	99%	0.0	88.57%	1087533	NW_015960952.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1633	<i>Capsicum annuum</i>	2409	8,71E+03	85%	0.0	75.86%	1620172	NW_015961041.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	2039	1,99E+04	80%	0.0	75.14%	7059680	NW_015960855.1
monotypic-Ch05f_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2866	<i>Capsicum annuum</i>	1716	1,44E+04	83%	0.0	77.27%	1254371	NW_015961127.1
monotypic-Ch06e_21	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10089	1,11E+04	100%	0.0	100.00%	219521584	NC_029982.1
monotypic-Ch06e_21	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8161	9,21E+03	100%	0.0	93.66%	205736368	NC_029986.1
monotypic-Ch06e_21	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7293	9,01E+03	97%	0.0	94.22%	163962470	NC_029978.1
monotypic-Ch06e_21	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5258	6,31E+03	85%	0.0	92.03%	229934170	NC_029988.1

monotypic-Ch06e_21	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4392	7,79E+03	98%	0.0	88.61%	261510930	NC_029979.1
monotypic-Ch06e_21	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3474	8,61E+03	98%	0.0	92.39%	222112641	NC_029983.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	24184	2,65E+08	100%	0.0	100.00%	219521584	NC_029982.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	16676	3,18E+08	100%	0.0	89.76%	153299543	NC_029984.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	16430	4,06E+08	100%	0.0	89.51%	217274494	NC_029981.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold985	<i>Capsicum annuum</i>	15847	2,03E+04	99%	0.0	88.58%	1349800	NW_015961816.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	15271	2,28E+08	100%	0.0	89.29%	222112641	NC_029983.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold277	<i>Capsicum annuum</i>	14999	1,85E+04	100%	0.0	87.87%	448122	NW_015961390.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	14654	4,56E+08	100%	0.0	89.45%	301019445	NC_029977.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	14388	3,05E+08	100%	0.0	86.63%	261510930	NC_029979.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	13016	3,49E+08	100%	0.0	88.99%	229934170	NC_029988.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11699	2,68E+08	100%	0.0	87.53%	205736368	NC_029986.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10358	2,97E+08	100%	0.0	89.03%	215701946	NC_029980.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	9803	1,76E+04	97%	0.0	86.56%	3311724	NW_015960635.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9686	2,36E+08	100%	0.0	89.08%	163962470	NC_029978.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	9474	1,81E+04	94%	0.0	88.53%	3493078	NW_015960517.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold891	<i>Capsicum annuum</i>	9404	2,24E+04	98%	0.0	87.95%	1431046	NW_015960469.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9044	3,37E+08	100%	0.0	88.53%	238794889	NC_029985.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold949	<i>Capsicum annuum</i>	8883	1,81E+04	92%	0.0	88.25%	3128562	NW_015961487.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8817	3,00E+08	100%	0.0	86.53%	220335243	NC_029987.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1124	<i>Capsicum annuum</i>	5221	1,00E+04	89%	0.0	80.04%	2110277	NW_015961370.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold189	<i>Capsicum annuum</i>	4654	2,21E+04	99%	0.0	89.22%	3791111	NW_015960691.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	4183	2,73E+04	94%	0.0	82.05%	7059680	NW_015960855.1
monotypic-Ch06f_46	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3339	<i>Capsicum annuum</i>	2820	1,39E+04	92%	0.0	87.23%	1234997	NW_015960491.1
monotypic-Ch08d_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	18159	5,05E+04	100%	0.0	100.00%	153299543	NC_029984.1
monotypic-Ch08d_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10262	4,94E+04	99%	0.0	92.45%	222112641	NC_029983.1
monotypic-Ch08d_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9350	2,98E+04	98%	0.0	90.14%	261510930	NC_029979.1
monotypic-Ch08d_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8268	6,63E+04	99%	0.0	90.00%	219521584	NC_029982.1
monotypic-Ch08d_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8041	4,93E+04	99%	0.0	92.68%	217274494	NC_029981.1
monotypic-Ch08d_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7637	4,16E+04	99%	0.0	92.47%	215701946	NC_029980.1
monotypic-Ch08d_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6926	3,41E+04	99%	0.0	89.33%	205736368	NC_029986.1
monotypic-Ch08d_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold949	<i>Capsicum annuum</i>	6839	1,65E+04	99%	0.0	90.92%	3128562	NW_015961487.1
monotypic-Ch08d_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capsicum annuum</i>	5723	1,32E+04	99%	0.0	90.39%	2397553	NW_015961318.1
monotypic-Ch08d_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4272	4,32E+04	98%	0.0	90.57%	229934170	NC_029988.1
monotypic-Ch08d_10	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4126	5,47E+04	99%	0.0	82.74%	301019445	NC_029977.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	22445	1,33E+09	100%	0.0	100.00%	238794889	NC_029985.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6656	1,01E+09	100%	0.0	91.27%	261510930	NC_029979.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6610	5,09E+08	100%	0.0	92.57%	153299543	NC_029984.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6205	7,05E+08	100%	0.0	90.64%	222112641	NC_029983.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1038	<i>Capsicum annuum</i>	6107	2,34E+04	96%	0.0	90.32%	2164954	NW_015960831.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5993	9,67E+08	100%	0.0	90.21%	301019445	NC_029977.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	5932	9,07E+04	99%	0.0	90.40%	7059680	NW_015960855.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5880	8,81E+08	100%	0.0	91.49%	217274494	NC_029981.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5818	7,03E+08	100%	0.0	91.44%	229934170	NC_029988.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5747	7,56E+08	100%	0.0	89.49%	215701946	NC_029980.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1587	<i>Capsicum annuum</i>	5742	1,59E+04	90%	0.0	90.92%	1400685	NW_015961196.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5698	7,68E+08	100%	0.0	90.75%	205736368	NC_029986.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5677	7,05E+08	100%	0.0	90.69%	220335243	NC_029987.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5568	4,89E+08	100%	0.0	90.39%	219521584	NC_029982.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4985	2,40E+04	96%	0.0	91.01%	3493078	NW_015960517.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	4868	2,09E+04	89%	0.0	87.87%	2382255	NW_015961471.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold742	<i>Capsicum annuum</i>	4865	1,68E+04	95%	0.0	89.53%	1549638	NW_015960466.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1307	<i>Capsicum annuum</i>	4713	1,88E+04	94%	0.0	90.84%	1494726	NW_015960454.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2298	<i>Capsicum annuum</i>	4711	2,00E+04	96%	0.0	92.14%	1101621	NW_015961578.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold519	<i>Capsicum annuum</i>	4458	1,56E+04	95%	0.0	90.26%	198665	NW_015960810.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3151	<i>Capsicum annuum</i>							

monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1620	<i>Capsicum annuum</i>	4453	1,73E+04	93%	0.0	90.36%	500299	NW_015961331.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	4447	1,64E+04	82%	0.0	85.97%	3311724	NW_015960635.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4204	<i>Capsicum annuum</i>	4442	3,45E+04	98%	0.0	90.38%	323636	NW_015960601.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold454	<i>Capsicum annuum</i>	4401	2,43E+04	93%	0.0	90.06%	3392484	NW_015961509.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold325	<i>Capsicum annuum</i>	4235	1,78E+04	80%	0.0	92.03%	480952	NW_015961410.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	3762	1,77E+04	92%	0.0	90.39%	2470984	NW_015961229.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2697	<i>Capsicum annuum</i>	3675	2,84E+04	98%	0.0	90.02%	139253	NW_015961756.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold983	<i>Capsicum annuum</i>	3661	1,73E+04	90%	0.0	91.20%	627343	NW_015961594.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold46	<i>Capsicum annuum</i>	3232	1,88E+04	93%	0.0	90.13%	941867	NW_015960787.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1074	<i>Capsicum annuum</i>	3099	1,40E+04	87%	0.0	85.61%	1095636	NW_015961132.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold603	<i>Capsicum annuum</i>	2436	2,25E+04	94%	0.0	89.97%	1800753	NW_015960521.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1923	<i>Capsicum annuum</i>	2386	2,47E+04	96%	0.0	91.10%	2480791	NW_015961228.1
monotypic-Ch09a_15	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1647	<i>Capsicum annuum</i>	2307	1,41E+04	82%	0.0	92.20%	1362880	NW_015960892.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23693	1,83E+09	100%	0.0	99.94%	238794889	NC_029985.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10763	1,28E+09	99%	0.0	93.02%	301019445	NC_029977.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10621	1,14E+09	99%	0.0	92.82%	219521584	NC_029982.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10362	1,47E+09	100%	0.0	92.05%	261510930	NC_029979.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10205	1,22E+09	99%	0.0	91.85%	217274494	NC_029981.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	10059	1,12E+08	99%	0.0	91.32%	7059680	NW_015960855.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10035	7,22E+08	99%	0.0	92.30%	153299543	NC_029984.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10028	1,13E+09	99%	0.0	91.55%	229934170	NC_029988.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9947	9,46E+08	99%	0.0	91.47%	222112641	NC_029983.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9529	1,12E+09	99%	0.0	91.92%	215701946	NC_029980.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9378	1,08E+09	99%	0.0	91.93%	205736368	NC_029986.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9239	7,71E+08	99%	0.0	91.87%	163962470	NC_029978.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8962	9,01E+08	99%	0.0	89.34%	220335243	NC_029987.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold742	<i>Capsicum annuum</i>	8468	2,62E+04	96%	0.0	88.18%	2382255	NW_015961471.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	7797	2,50E+04	86%	0.0	86.46%	3311724	NW_015960635.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2707	<i>Capsicum annuum</i>	7426	1,96E+04	93%	0.0	86.58%	1896504	NW_015961173.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold519	<i>Capsicum annuum</i>	7254	1,81E+04	91%	0.0	90.13%	1101621	NW_015961578.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1587	<i>Capsicum annuum</i>	7040	1,63E+04	81%	0.0	91.96%	1400685	NW_015961196.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1038	<i>Capsicum annuum</i>	6870	2,91E+04	93%	0.0	89.53%	2164954	NW_015960831.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold454	<i>Capsicum annuum</i>	6421	3,71E+04	98%	0.0	93.29%	3392484	NW_015961509.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1620	<i>Capsicum annuum</i>	6176	1,74E+04	93%	0.0	90.17%	500299	NW_015961331.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	6107	3,24E+04	96%	0.0	90.62%	3493078	NW_015960517.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2697	<i>Capsicum annuum</i>	6028	2,92E+04	88%	0.0	91.15%	139253	NW_015961756.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4204	<i>Capsicum annuum</i>	5406	3,75E+04	97%	0.0	89.84%	323636	NW_015960601.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold46	<i>Capsicum annuum</i>	5306	2,46E+04	92%	0.0	91.02%	941867	NW_015960787.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2298	<i>Capsicum annuum</i>	4828	2,24E+04	90%	0.0	91.46%	1494726	NW_015960454.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold603	<i>Capsicum annuum</i>	4614	2,41E+04	92%	0.0	80.77%	1800753	NW_015960521.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3151	<i>Capsicum annuum</i>	4577	1,69E+04	93%	0.0	90.89%	198665	NW_015960810.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1307	<i>Capsicum annuum</i>	4464	1,96E+04	90%	0.0	91.26%	1549638	NW_015960466.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold983	<i>Capsicum annuum</i>	3794	1,88E+04	87%	0.0	92.09%	627343	NW_015961594.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1074	<i>Capsicum annuum</i>	3760	1,67E+04	88%	0.0	85.53%	1095636	NW_015961132.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2948	<i>Capsicum annuum</i>	3448	1,86E+04	82%	0.0	92.83%	898408	NW_015961258.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold846	<i>Capsicum annuum</i>	3282	1,56E+04	89%	0.0	83.91%	1232610	NW_015960818.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4183	<i>Capsicum annuum</i>	3149	2,79E+04	80%	0.0	79.11%	1223518	NW_015961565.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1923	<i>Capsicum annuum</i>	2808	2,95E+04	93%	0.0	90.65%	2480791	NW_015961228.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1647	<i>Capsicum annuum</i>	2462	1,77E+04	90%	0.0	91.34%	1362880	NW_015960892.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	2266	2,16E+04	88%	0.0	76.92%	2470984	NW_015961229.1
monotypic-Ch09a_16	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1434	<i>Capsicum annuum</i>	2261	9,53E+03	81%	0.0	76.79%	1100793	NW_015960373.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23028	9,26E+08	100%	0.0	100.00%	238794889	NC_029985.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold846	<i>Capsicum annuum</i>	11832	2,21E+04	99%	0.0	90.07%	1232610	NW_015960818.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8600	5,60E+08	100%	0.0	89.89%	229934170	NC_029988.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8475	5,07E+08	100%	0.0	88.24%	222112641	NC_029983.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7520	5,39E+08	100%	0.0	88.82%	215701946	NC_029980.1

monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7480	6,66E+08	97%	0.0	87.68%	301019445	NC_029977.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7476	4,85E+08	100%	0.0	86.09%	220335243	NC_029987.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6830	6,74E+08	100%	0.0	90.03%	261510930	NC_029979.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6589	3,80E+08	100%	0.0	89.27%	153299543	NC_029984.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6553	5,18E+08	92%	0.0	86.57%	219521584	NC_029982.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6279	5,64E+08	93%	0.0	85.75%	217274494	NC_029981.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold603	<i>Capsicum annuum</i>	6259	1,71E+04	89%	0.0	85.65%	1800753	NW_015960521.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6071	3,17E+08	92%	0.0	85.24%	163962470	NC_029978.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5415	5,47E+08	92%	0.0	83.20%	205736368	NC_029986.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	5223	4,89E+04	91%	0.0	80.30%	7059680	NW_015960855.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold742	<i>Capsicum annuum</i>	5051	1,34E+04	89%	0.0	80.01%	2382255	NW_015961471.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2707	<i>Capsicum annuum</i>	4447	9,55E+03	82%	0.0	79.14%	1896504	NW_015961173.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold454	<i>Capsicum annuum</i>	3938	1,29E+04	91%	0.0	83.84%	3392484	NW_015961509.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	3903	1,49E+04	85%	0.0	82.76%	3493078	NW_015960517.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4204	<i>Capsicum annuum</i>	3434	1,40E+04	80%	0.0	82.67%	323636	NW_015960601.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold46	<i>Capsicum annuum</i>	3245	1,20E+04	89%	0.0	81.66%	941867	NW_015960787.1
monotypic-Ch09b_19	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2298	<i>Capsicum annuum</i>	2545	9,74E+03	81%	0.0	85.61%	1494726	NW_015960454.1
monotypic-Ch09e_06	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	18295	1,49E+09	100%	0.0	100.00%	238794889	NC_029985.1
monotypic-Ch09e_06	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9943	7,74E+08	100%	0.0	88.57%	301019445	NC_029977.1
monotypic-Ch09e_06	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9814	6,08E+08	100%	0.0	88.35%	220335243	NC_029987.1
monotypic-Ch09e_06	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9409	9,68E+08	100%	0.0	90.17%	215701946	NC_029980.1
monotypic-Ch09e_06	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8630	6,59E+08	100%	0.0	85.85%	153299543	NC_029984.1
monotypic-Ch09e_06	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7672	1,33E+09	100%	0.0	88.70%	261510930	NC_029979.1
monotypic-Ch09e_06	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5939	4,65E+08	100%	0.0	88.78%	222112641	NC_029983.1
monotypic-Ch09e_06	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5432	1,33E+09	100%	0.0	87.17%	219521584	NC_029982.1
monotypic-Ch09e_06	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5411	8,32E+08	99%	0.0	86.76%	205736368	NC_029986.1
monotypic-Ch09e_06	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5260	1,11E+09	98%	0.0	86.53%	217274494	NC_029981.1
monotypic-Ch09e_06	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	5203	3,17E+04	91%	0.0	86.30%	3311724	NW_015960635.1
monotypic-Ch09e_06	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5179	1,52E+09	100%	0.0	86.30%	229934170	NC_029988.1
monotypic-Ch09e_06	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold985	<i>Capsicum annuum</i>	5127	1,19E+04	83%	0.0	85.67%	1349800	NW_015961816.1
monotypic-Ch09e_06	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	4820	3,67E+04	95%	0.0	85.87%	3493078	NW_015960517.1
monotypic-Ch09e_06	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4748	9,97E+08	96%	0.0	85.61%	163962470	NC_029978.1
monotypic-Ch09e_06	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	3528	4,61E+04	94%	0.0	81.34%	7059680	NW_015960855.1
monotypic-Ch09f_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9731	1,11E+04	100%	0.0	100.00%	238794889	NC_029985.1
monotypic-Ch09f_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7443	3,14E+04	100%	0.0	92.17%	217274494	NC_029981.1
monotypic-Ch09f_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6841	9,05E+03	98%	0.0	93.58%	222112641	NC_029983.1
monotypic-Ch09f_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6833	1,77E+04	100%	0.0	93.15%	261510930	NC_029979.1
monotypic-Ch09f_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4850	1,03E+04	96%	0.0	93.59%	301019445	NC_029977.1
monotypic-Ch09f_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4776	9,18E+03	100%	0.0	93.05%	229934170	NC_029988.1
monotypic-Ch09f_24	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3808	8,30E+03	100%	0.0	91.18%	220335243	NC_029987.1
monotypic-Ch09f_57	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10237	1,96E+04	100%	0.0	100.00%	238794889	NC_029985.1
monotypic-Ch09f_57	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7705	1,18E+04	100%	0.0	91.68%	219521584	NC_029982.1
monotypic-Ch09f_57	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7657	1,98E+04	100%	0.0	92.85%	301019445	NC_029977.1
monotypic-Ch09f_57	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7539	1,32E+04	100%	0.0	91.36%	261510930	NC_029979.1
monotypic-Ch09f_57	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6523	9,42E+03	99%	0.0	92.66%	163962470	NC_029978.1
monotypic-Ch09f_57	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4892	9,51E+03	99%	0.0	93.36%	153299543	NC_029984.1
monotypic-Ch09f_57	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2274	<i>Capsicum annuum</i>	3882	7,36E+03	100%	0.0	87.96%	1395869	NW_015961152.1
monotypic-Ch09f_70	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9286	1,75E+04	100%	0.0	100.00%	238794889	NC_029985.1
monotypic-Ch09f_70	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6536	9,22E+03	98%	0.0	94.63%	215701946	NC_029980.1
monotypic-Ch09f_70	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4863	1,33E+04	100%	0.0	90.14%	301019445	NC_029977.1
monotypic-Ch09f_70	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4615	6,96E+03	99%	0.0	88.41%	261510930	NC_029979.1
monotypic-Ch09f_70	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00548	<i>Solanum tuberosum</i>	3339	4,48E+03	98%	0.0	82.57%	317980	NW_006239470.1
monotypic-Ch09f_70	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3212	7,31E+03	91%	0.0	92.19%	222112641	NC_029983.1
monotypic-Ch09f_70	refeq-genome	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00532	<i>Solanum tuberosum</i>	2772	4,67E+03	96%	0.0	83.67%	325781	NW_006239461.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	18450	8,59E+08	100%	0.0	100.00%	205736368	NC_029986.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10237	9,19E+08	100%	0.0	85.94%	238794889	NC_029985.1

monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9677	1,13E+09	100%	0.0	86.78%	301019445	NC_029977.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9631	9,08E+08	100%	0.0	86.70%	261510930	NC_029979.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9426	9,95E+08	100%	0.0	86.26%	22112641	NC_029983.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9313	8,75E+08	100%	0.0	86.03%	217274494	NC_029981.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8726	7,24E+08	100%	0.0	84.88%	215701946	NC_029980.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8685	8,30E+08	100%	0.0	86.26%	219521584	NC_029982.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8671	5,62E+08	100%	0.0	86.49%	163962470	NC_029978.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8624	8,14E+08	100%	0.0	86.10%	229934170	NC_029988.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold669	<i>Capsicum annuum</i>	8477	2,04E+04	97%	0.0	84.55%	640007	NW_015960674.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8427	5,96E+08	100%	0.0	84.21%	153299543	NC_029984.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8371	7,09E+08	100%	0.0	85.56%	220335243	NC_029987.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold90	<i>Capsicum annuum</i>	7775	4,09E+04	99%	0.0	84.97%	6074945	NW_015960458.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold516	<i>Capsicum annuum</i>	7733	4,10E+04	100%	0.0	85.43%	4299619	NW_015961344.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1124	<i>Capsicum annuum</i>	7380	6,80E+04	100%	0.0	80.29%	2110277	NW_015961370.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1118	<i>Capsicum annuum</i>	7241	8,65E+03	88%	0.0	82.97%	528969	NW_015960738.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	7053	5,55E+04	100%	0.0	84.35%	7059680	NW_015960855.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3202	<i>Capsicum annuum</i>	6482	1,13E+04	96%	0.0	82.15%	302159	NW_015961119.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1246	<i>Capsicum annuum</i>	5962	2,27E+04	91%	0.0	86.56%	3388803	NW_015960834.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1651	<i>Capsicum annuum</i>	5869	8,42E+03	84%	0.0	81.57%	331682	NW_015961206.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold296	<i>Capsicum annuum</i>	5747	9,31E+03	90%	0.0	81.28%	1995018	NW_015961641.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1038	<i>Capsicum annuum</i>	5653	1,67E+04	99%	0.0	84.51%	2164954	NW_015960831.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold454	<i>Capsicum annuum</i>	5605	1,87E+04	90%	0.0	80.91%	3392484	NW_015961509.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1390	<i>Capsicum annuum</i>	5404	9,04E+03	93%	0.0	80.45%	1730405	NW_015960783.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1923	<i>Capsicum annuum</i>	5186	2,07E+04	87%	0.0	79.99%	2480791	NW_015961228.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3714	<i>Capsicum annuum</i>	5073	8,19E+03	85%	0.0	82.19%	935490	NW_015961128.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold985	<i>Capsicum annuum</i>	4998	2,46E+04	88%	0.0	79.43%	1349800	NW_015961816.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2102	<i>Capsicum annuum</i>	4994	7,05E+03	86%	0.0	80.94%	627640	NW_015961440.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold438	<i>Capsicum annuum</i>	4761	1,76E+04	100%	0.0	79.64%	1337825	NW_015961507.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5097	<i>Capsicum annuum</i>	4643	7,18E+03	89%	0.0	79.43%	302211	NW_015961801.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold933	<i>Capsicum annuum</i>	4610	1,90E+04	97%	0.0	79.86%	1853301	NW_015961435.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	4571	1,78E+04	99%	0.0	81.49%	2470984	NW_015961229.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold170	<i>Capsicum annuum</i>	4569	1,83E+04	96%	0.0	79.18%	2217302	NW_015960390.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold349	<i>Capsicum annuum</i>	4551	1,39E+04	91%	0.0	78.34%	2130419	NW_015960729.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold619	<i>Capsicum annuum</i>	4525	1,38E+04	100%	0.0	79.70%	2512642	NW_015960498.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1341	<i>Capsicum annuum</i>	4510	1,24E+04	95%	0.0	82.80%	1906221	NW_015960665.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3026	<i>Capsicum annuum</i>	4198	1,61E+04	97%	0.0	80.14%	1386929	NW_015961102.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold231	<i>Capsicum annuum</i>	3962	8,72E+03	93%	0.0	78.08%	1482909	NW_015961568.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2707	<i>Capsicum annuum</i>	3908	1,45E+04	99%	0.0	77.06%	1896504	NW_015961173.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold519	<i>Capsicum annuum</i>	3882	1,08E+04	89%	0.0	81.89%	1101621	NW_015961578.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2375	<i>Capsicum annuum</i>	3639	1,34E+04	94%	0.0	84.63%	1171512	NW_015961637.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold969	<i>Capsicum annuum</i>	3212	8,44E+03	87%	0.0	79.80%	1557263	NW_015961740.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1973	<i>Capsicum annuum</i>	3144	9,07E+03	80%	0.0	82.46%	1714326	NW_015961422.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold189	<i>Capsicum annuum</i>	3142	1,47E+04	95%	0.0	84.99%	3791111	NW_015960691.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold578	<i>Capsicum annuum</i>	3101	1,07E+04	87%	0.0	80.70%	1213777	NW_015961091.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3499	<i>Capsicum annuum</i>	2789	9,26E+03	87%	0.0	81.97%	681687	NW_015960538.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9	<i>Capsicum annuum</i>	2564	2,71E+04	100%	0.0	80.80%	6567951	NW_015961815.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1741	<i>Capsicum annuum</i>	2398	6,84E+03	83%	0.0	81.65%	1897442	NW_015960687.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	<i>Capsicum annuum</i>	2372	9,94E+03	99%	0.0	78.05%	2397553	NW_015961318.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold742	<i>Capsicum annuum</i>	2100	7,13E+03	91%	0.0	82.68%	2382255	NW_015961471.1
monotypic-Ch10c_34	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	1851	1,48E+04	94%	0.0	84.36%	3493078	NW_015960517.1
monotypic-Ch10f_29	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9939	1,13E+04	100%	0.0	100.00%	205736368	NC_029986.1
monotypic-Ch10f_29	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7155	8,26E+03	100%	0.0	90.74%	229934170	NC_029988.1
monotypic-Ch10f_29	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6191	6,61E+03	100%	0.0	87.57%	163962470	NC_029978.1
monotypic-Ch10f_29	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3229	1,01E+04	100%	0.0	86.86%	238794889	NC_029985.1
monotypic-Ch10f_29	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3005	1,62E+04	92%	0.0	90.75%	301019445	NC_029977.1
monotypic-Ch11a_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	18667	8,78E+04	100%	0.0	100.00%	220335243	NC_029987.1

monotypic-Ch11a_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9762	9,72E+04	100%	0.0	90.54%	217274494	NC_029981.1
monotypic-Ch11a_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8185	5,84E+04	100%	0.0	91.47%	153299543	NC_029984.1
monotypic-Ch11a_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6924	8,47E+04	87%	0.0	84.60%	219521584	NC_029982.1
monotypic-Ch11a_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6848	9,96E+04	100%	0.0	89.33%	261510930	NC_029979.1
monotypic-Ch11a_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6758	8,34E+04	92%	0.0	86.45%	222112641	NC_029983.1
monotypic-Ch11a_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6676	1,18E+08	93%	0.0	84.17%	238794889	NC_029985.1
monotypic-Ch11a_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6575	6,59E+04	90%	0.0	83.93%	229934170	NC_029988.1
monotypic-Ch11a_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6499	7,40E+04	91%	0.0	84.96%	205736368	NC_029986.1
monotypic-Ch11a_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6338	1,29E+08	100%	0.0	85.14%	301019445	NC_029977.1
monotypic-Ch11a_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6277	1,36E+08	100%	0.0	83.20%	163962470	NC_029978.1
monotypic-Ch11a_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold592	<i>Capsicum annuum</i>	5939	9,69E+03	85%	0.0	83.81%	1905997	NW_015960989.1
monotypic-Ch11a_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	5166	6,09E+04	92%	0.0	83.17%	215701946	NC_029980.1
monotypic-Ch11a_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1754	<i>Capsicum annuum</i>	4148	8,34E+03	87%	0.0	82.10%	1029277	NW_015960393.1
monotypic-Ch11a_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2635	<i>Capsicum annuum</i>	3472	1,39E+04	89%	0.0	81.82%	699991	NW_015960547.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	23979	6,70E+08	100%	0.0	100.00%	220335243	NC_029987.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10080	6,93E+08	99%	0.0	91.82%	229934170	NC_029988.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10028	7,49E+08	100%	0.0	89.47%	205736368	NC_029986.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9904	7,46E+08	99%	0.0	91.35%	215701946	NC_029980.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9762	4,61E+08	99%	0.0	91.12%	163962470	NC_029978.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9747	8,62E+08	99%	0.0	91.01%	217274494	NC_029981.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9655	9,75E+08	99%	0.0	90.75%	261510930	NC_029979.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold742	<i>Capsicum annuum</i>	9516	2,92E+04	99%	0.0	90.50%	2382255	NW_015961471.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9507	6,69E+08	99%	0.0	90.48%	219521584	NC_029982.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9487	1,23E+09	100%	0.0	90.36%	238794889	NC_029985.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9345	6,80E+08	99%	0.0	90.00%	222112641	NC_029983.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8927	9,12E+08	99%	0.0	89.25%	301019445	NC_029977.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8857	5,10E+08	100%	0.0	91.25%	153299543	NC_029984.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold454	<i>Capsicum annuum</i>	8399	2,47E+04	94%	0.0	88.67%	3392484	NW_015961509.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2707	<i>Capsicum annuum</i>	8347	1,86E+04	92%	0.0	89.03%	1896504	NW_015961173.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	8342	1,97E+04	92%	0.0	87.55%	3311724	NW_015960635.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	8329	8,23E+04	97%	0.0	87.72%	7059680	NW_015960855.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold519	<i>Capsicum annuum</i>	6288	1,45E+04	86%	0.0	87.80%	1101621	NW_015961578.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1038	<i>Capsicum annuum</i>	5976	1,92E+04	82%	0.0	88.50%	2164954	NW_015960831.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	5247	2,02E+04	87%	0.0	87.30%	3493078	NW_015960517.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1620	<i>Capsicum annuum</i>	5228	1,28E+04	84%	0.0	86.62%	500299	NW_015961331.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2697	<i>Capsicum annuum</i>	5118	2,31E+04	84%	0.0	87.48%	139253	NW_015961756.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold46	<i>Capsicum annuum</i>	4835	1,54E+04	80%	0.0	88.87%	941867	NW_015960787.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold603	<i>Capsicum annuum</i>	4623	1,94E+04	90%	0.0	80.82%	1800753	NW_015960521.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4204	<i>Capsicum annuum</i>	4593	2,73E+04	90%	0.0	86.43%	323636	NW_015960601.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1074	<i>Capsicum annuum</i>	4505	2,08E+04	92%	0.0	89.36%	1095636	NW_015961132.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2298	<i>Capsicum annuum</i>	4146	1,58E+04	86%	0.0	88.53%	1494726	NW_015960454.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3151	<i>Capsicum annuum</i>	3982	1,40E+04	92%	0.0	88.31%	198665	NW_015960810.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1307	<i>Capsicum annuum</i>	3962	1,47E+04	89%	0.0	88.58%	1549638	NW_015960466.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold983	<i>Capsicum annuum</i>	3308	1,57E+04	89%	0.0	88.86%	627343	NW_015961594.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold846	<i>Capsicum annuum</i>	3195	1,21E+04	88%	0.0	83.44%	1232610	NW_015960818.1
monotypic-Ch11a_42	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	2021	1,55E+04	85%	0.0	90.23%	2470984	NW_015961229.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	25601	3,95E+08	100%	0.0	100.00%	220335243	NC_029987.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	18306	4,06E+08	100%	0.0	90.58%	219521584	NC_029982.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	17669	5,38E+08	100%	0.0	89.75%	238794889	NC_029985.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	17451	3,70E+08	100%	0.0	89.91%	217274494	NC_029981.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	17403	3,73E+08	100%	0.0	89.96%	229934170	NC_029988.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	16814	5,26E+04	100%	0.0	90.49%	7059680	NW_015960855.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	15797	3,68E+08	100%	0.0	90.64%	222112641	NC_029983.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	15555	1,88E+08	100%	0.0	90.47%	163962470	NC_029978.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	14870	4,12E+08	100%	0.0	91.29%	215701946	NC_029980.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4660	<i>Capsicum annuum</i>	14796	2,35E+04	100%	0.0	91.46%	152652	NW_015961314.1

monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1963	<i>Capsicum annuum</i>	14565	2,03E+04	99%	0.0	88.74%	732782	NW_015960474.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	14103	5,00E+08	100%	0.0	88.94%	261510930	NC_029979.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	13985	3,16E+08	100%	0.0	88.93%	205736368	NC_029986.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	13941	2,57E+08	100%	0.0	89.17%	153299543	NC_029984.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	13298	5,83E+08	100%	0.0	88.92%	301019445	NC_029977.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1741	<i>Capsicum annuum</i>	9670	2,10E+04	98%	0.0	88.39%	1897442	NW_015960687.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2866	<i>Capsicum annuum</i>	8835	2,76E+04	81%	0.0	91.65%	1254371	NW_015961127.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4636	<i>Capsicum annuum</i>	8333	1,98E+04	87%	0.0	89.28%	463876	NW_015960556.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1633	<i>Capsicum annuum</i>	7300	1,14E+04	87%	0.0	83.51%	1620172	NW_015961041.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold603	<i>Capsicum annuum</i>	7295	1,65E+04	95%	0.0	83.50%	1800753	NW_015960521.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3026	<i>Capsicum annuum</i>	5433	2,41E+04	98%	0.0	84.25%	1386929	NW_015961102.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold572	<i>Capsicum annuum</i>	5384	1,02E+04	81%	0.0	81.25%	590163	NW_015960560.1
monotypic-Ch11a_51	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold6454	<i>Capsicum annuum</i>	3797	6,58E+03	84%	0.0	76.62%	84551	NW_015961449.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	18691	1,11E+09	100%	0.0	100.00%	220335243	NC_029987.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	12417	1,39E+09	100%	0.0	88.84%	238794889	NC_029985.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold303	<i>Capsicum annuum</i>	12233	1,61E+04	100%	0.0	88.57%	285559	NW_015960893.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11930	1,77E+09	100%	0.0	88.03%	301019445	NC_029977.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11836	1,28E+09	100%	0.0	87.82%	215701946	NC_029980.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11202	1,61E+09	100%	0.0	86.79%	222112641	NC_029983.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11047	8,34E+08	100%	0.0	86.47%	163962470	NC_029978.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10870	1,57E+09	100%	0.0	86.19%	261510930	NC_029979.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold349	<i>Capsicum annuum</i>	9827	3,83E+04	100%	0.0	84.33%	2130419	NW_015960729.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9736	1,25E+09	100%	0.0	84.11%	217274494	NC_029981.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9568	1,27E+09	100%	0.0	83.85%	205736368	NC_029986.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1923	<i>Capsicum annuum</i>	9258	3,33E+04	100%	0.0	83.39%	2480791	NW_015961228.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9193	1,32E+09	100%	0.0	83.19%	229934170	NC_029988.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold985	<i>Capsicum annuum</i>	8588	4,61E+04	100%	0.0	82.10%	1349800	NW_015961816.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8464	1,12E+09	100%	0.0	86.63%	219521584	NC_029982.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8436	7,41E+08	100%	0.0	84.08%	153299543	NC_029984.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	7967	3,46E+04	100%	0.0	82.87%	3311724	NW_015960635.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1341	<i>Capsicum annuum</i>	7779	2,31E+04	100%	0.0	81.74%	1906221	NW_015960665.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold516	<i>Capsicum annuum</i>	7533	5,88E+04	100%	0.0	81.24%	4299619	NW_015961344.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2157	<i>Capsicum annuum</i>	7522	1,35E+04	100%	0.0	82.33%	158560	NW_015960692.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9337	<i>Capsicum annuum</i>	7396	2,05E+04	100%	0.0	81.77%	397715	NW_015961734.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1124	<i>Capsicum annuum</i>	7278	7,22E+04	100%	0.0	79.90%	2110277	NW_015961370.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4747	<i>Capsicum annuum</i>	6966	1,92E+04	100%	0.0	79.50%	1151343	NW_015960460.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold189	<i>Capsicum annuum</i>	6927	2,85E+04	100%	0.0	82.14%	3791111	NW_015960691.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1670	<i>Capsicum annuum</i>	6684	4,16E+04	96%	0.0	88.74%	1202455	NW_015961465.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2451	<i>Capsicum annuum</i>	6678	1,88E+04	100%	0.0	81.82%	364752	NW_015961179.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold24	<i>Capsicum annuum</i>	6560	1,18E+04	95%	0.0	81.25%	1258456	NW_015961700.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3026	<i>Capsicum annuum</i>	6477	1,71E+04	99%	0.0	87.05%	1386929	NW_015961102.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3611	<i>Capsicum annuum</i>	6401	7,06E+03	85%	0.0	81.67%	177705	NW_015960430.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2707	<i>Capsicum annuum</i>	6381	1,79E+04	100%	0.0	82.26%	1896504	NW_015961173.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold454	<i>Capsicum annuum</i>	6226	3,79E+04	100%	0.0	80.11%	3392484	NW_015961509.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1651	<i>Capsicum annuum</i>	5930	6,52E+03	90%	0.0	79.80%	331682	NW_015961206.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold296	<i>Capsicum annuum</i>	5803	1,07E+04	97%	0.0	79.59%	1995018	NW_015961641.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1390	<i>Capsicum annuum</i>	5596	7,04E+03	84%	0.0	78.68%	1730405	NW_015960783.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold983	<i>Capsicum annuum</i>	5507	1,11E+04	99%	0.0	83.28%	627343	NW_015961594.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3714	<i>Capsicum annuum</i>	4741	1,07E+04	100%	0.0	80.49%	935490	NW_015961128.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1973	<i>Capsicum annuum</i>	4739	3,40E+04	100%	0.0	87.92%	1714326	NW_015961422.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2298	<i>Capsicum annuum</i>	4553	1,65E+04	100%	0.0	80.08%	1494726	NW_015960454.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold90	<i>Capsicum annuum</i>	4447	4,49E+04	100%	0.0	80.20%	6074945	NW_015960458.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2945	<i>Capsicum annuum</i>	4226	7,49E+03	98%	0.0	83.86%	320440	NW_015961001.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9	<i>Capsicum annuum</i>	4215	2,75E+04	96%	0.0	79.77%	6567951	NW_015961815.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold188	<i>Capsicum annuum</i>	4074	7,32E+03	81%	0.0	80.60%	980346	NW_015960557.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3843	<i>Capsicum annuum</i>	4043	2,15E+04	99%	0.0	79.94%	465776	NW_015961174.1

monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold170	<i>Capsicum annuum</i>	4002	3,45E+04	100%	0.0	80.39%	2217302	NW_015960390.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	3978	4,14E+04	100%	0.0	79.53%	7059680	NW_015960855.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold438	<i>Capsicum annuum</i>	3964	6,87E+03	96%	0.0	78.08%	1337825	NW_015961507.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1038	<i>Capsicum annuum</i>	3869	1,53E+04	100%	0.0	80.73%	2164954	NW_015960831.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold933	<i>Capsicum annuum</i>	3664	1,42E+04	100%	0.0	77.92%	1853301	NW_015961435.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	3659	1,02E+04	88%	0.0	78.69%	2470984	NW_015961229.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2400	<i>Capsicum annuum</i>	3605	1,48E+04	100%	0.0	89.27%	1353284	NW_015960968.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1246	<i>Capsicum annuum</i>	3491	1,48E+04	98%	0.0	78.38%	3388803	NW_015960834.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3224	<i>Capsicum annuum</i>	3435	1,45E+04	85%	0.0	85.59%	891864	NW_015961692.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold619	<i>Capsicum annuum</i>	3321	1,02E+04	93%	0.0	81.83%	2512642	NW_015960498.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold969	<i>Capsicum annuum</i>	3072	1,23E+04	100%	0.0	78.94%	1557263	NW_015961740.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1741	<i>Capsicum annuum</i>	2994	1,59E+04	100%	0.0	81.96%	1897442	NW_015960687.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4204	<i>Capsicum annuum</i>	2433	1,13E+04	82%	0.0	86.96%	323636	NW_015960601.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold578	<i>Capsicum annuum</i>	2425	1,72E+04	92%	0.0	80.53%	1213777	NW_015961091.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2239	<i>Capsicum annuum</i>	2276	9,61E+03	85%	0.0	86.18%	615095	NW_015960816.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2274	<i>Capsicum annuum</i>	2209	7,41E+03	93%	0.0	74.58%	1395869	NW_015961152.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3339	<i>Capsicum annuum</i>	2128	8,97E+03	99%	0.0	80.67%	1234997	NW_015960491.1
monotypic-Ch11d_13	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1434	<i>Capsicum annuum</i>	1687	8,53E+03	90%	0.0	74.87%	1100793	NW_015960373.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	18242	5,84E+08	100%	0.0	100.00%	220335243	NC_029987.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	12955	8,57E+08	100%	0.0	90.34%	238794889	NC_029985.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	12338	8,33E+08	100%	0.0	89.27%	301019445	NC_029977.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold985	<i>Capsicum annuum</i>	12050	4,34E+04	100%	0.0	88.74%	1349800	NW_015961816.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11675	6,61E+08	100%	0.0	88.07%	261510930	NC_029979.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11234	6,51E+08	100%	0.0	89.43%	222112641	NC_029983.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10624	6,64E+08	100%	0.0	86.16%	229934170	NC_029988.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10473	6,99E+08	100%	0.0	90.72%	217274494	NC_029981.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9640	4,45E+08	100%	0.0	88.88%	163962470	NC_029978.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8863	5,54E+08	100%	0.0	86.42%	215701946	NC_029980.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7821	6,40E+08	100%	0.0	88.30%	219521584	NC_029982.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4063	<i>Capsicum annuum</i>	7520	2,81E+04	99%	0.0	89.23%	1657713	NW_015960826.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	7247	5,97E+08	100%	0.0	83.29%	205736368	NC_029986.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6218	4,33E+08	100%	0.0	81.32%	153299543	NC_029984.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold231	<i>Capsicum annuum</i>	6011	1,29E+04	90%	0.0	82.33%	1482909	NW_015961568.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1894	<i>Capsicum annuum</i>	4826	1,42E+04	88%	0.0	83.24%	2334823	NW_015961490.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold170	<i>Capsicum annuum</i>	4250	1,18E+04	98%	0.0	78.90%	2217302	NW_015960390.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold519	<i>Capsicum annuum</i>	3615	7,79E+03	81%	0.0	79.91%	1101621	NW_015961578.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1124	<i>Capsicum annuum</i>	3535	3,56E+04	100%	0.0	78.69%	2110277	NW_015961370.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold296	<i>Capsicum annuum</i>	3171	6,12E+03	88%	0.0	75.52%	1995018	NW_015961641.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold516	<i>Capsicum annuum</i>	3031	1,52E+04	91%	0.0	76.29%	4299619	NW_015961344.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	2881	3,44E+04	100%	0.0	76.53%	7059680	NW_015960855.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3026	<i>Capsicum annuum</i>	2663	9,59E+03	82%	0.0	77.93%	1386929	NW_015961102.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4108	<i>Capsicum annuum</i>	2608	9,82E+03	85%	0.0	78.69%	259817	NW_015960574.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold578	<i>Capsicum annuum</i>	2534	9,79E+03	85%	0.0	79.33%	1213777	NW_015961091.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1246	<i>Capsicum annuum</i>	2283	1,20E+04	97%	0.0	76.62%	3388803	NW_015960834.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9	<i>Capsicum annuum</i>	2001	1,48E+04	91%	0.0	79.68%	6567951	NW_015961815.1
monotypic-Ch11f_28	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2298	<i>Capsicum annuum</i>	1657	8,50E+03	80%	0.0	75.59%	1494726	NW_015960454.1
monotypic-Ch12a_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9659	1,07E+04	100%	0.0	100.00%	229934170	NC_029988.1
monotypic-Ch12a_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	6144	6,68E+03	92%	0.0	91.37%	153299543	NC_029984.1
monotypic-Ch12a_43	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3725	9,53E+03	97%	0.0	82.11%	220335243	NC_029987.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	22774	8,51E+08	100%	0.0	100.00%	229934170	NC_029988.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	13594	1,15E+09	100%	0.0	90.16%	261510930	NC_029979.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10748	1,08E+09	100%	0.0	90.84%	301019445	NC_029977.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10501	1,01E+09	100%	0.0	90.64%	217274494	NC_029981.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10240	7,56E+08	100%	0.0	89.93%	220335243	NC_029987.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10028	1,53E+09	100%	0.0	91.55%	238794889	NC_029985.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9773	5,73E+08	100%	0.0	90.31%	153299543	NC_029984.1

monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold174	<i>Capsicum annuum</i>	9760	9,93E+04	99%	0.0	90.84%	7059680	NW_015960855.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9744	7,33E+08	100%	0.0	90.95%	222112641	NC_029983.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9668	8,52E+08	100%	0.0	90.69%	219521584	NC_029982.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9077	8,98E+08	100%	0.0	91.05%	205736368	NC_029986.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8903	8,86E+08	100%	0.0	90.25%	215701946	NC_029980.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	8492	6,02E+08	100%	0.0	89.73%	163962470	NC_029978.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold742	<i>Capsicum annuum</i>	7910	2,21E+04	97%	0.0	86.76%	2382255	NW_015961471.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold519	<i>Capsicum annuum</i>	7354	1,84E+04	94%	0.0	90.43%	1101621	NW_015961578.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1276	<i>Capsicum annuum</i>	7304	1,92E+04	88%	0.0	85.21%	3311724	NW_015960635.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1587	<i>Capsicum annuum</i>	7201	1,70E+04	87%	0.0	91.31%	1400685	NW_015961196.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2707	<i>Capsicum annuum</i>	6970	1,54E+04	94%	0.0	85.35%	1896504	NW_015961173.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1038	<i>Capsicum annuum</i>	6800	2,63E+04	95%	0.0	90.82%	2164954	NW_015960831.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1620	<i>Capsicum annuum</i>	6209	1,68E+04	94%	0.0	90.26%	500299	NW_015961331.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	6141	2,57E+04	97%	0.0	90.71%	3493078	NW_015960517.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2697	<i>Capsicum annuum</i>	5945	3,01E+04	94%	0.0	90.79%	139253	NW_015961756.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold454	<i>Capsicum annuum</i>	5849	2,50E+04	97%	0.0	90.96%	3392484	NW_015961509.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4204	<i>Capsicum annuum</i>	5500	3,51E+04	98%	0.0	90.21%	323636	NW_015960601.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold46	<i>Capsicum annuum</i>	5323	2,02E+04	97%	0.0	91.07%	941867	NW_015960787.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2298	<i>Capsicum annuum</i>	4787	2,14E+04	96%	0.0	89.99%	1494726	NW_015960454.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3151	<i>Capsicum annuum</i>	4667	1,72E+04	98%	0.0	91.36%	198665	NW_015960810.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1307	<i>Capsicum annuum</i>	4407	1,76E+04	95%	0.0	90.93%	1549638	NW_015960466.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold603	<i>Capsicum annuum</i>	4353	2,49E+04	97%	0.0	79.94%	1800753	NW_015960521.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold499	<i>Capsicum annuum</i>	4023	1,98E+04	92%	0.0	90.16%	2470984	NW_015961229.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold983	<i>Capsicum annuum</i>	3749	1,74E+04	87%	0.0	91.80%	627343	NW_015961594.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold325	<i>Capsicum annuum</i>	3733	1,80E+04	80%	0.0	89.34%	480952	NW_015961410.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1074	<i>Capsicum annuum</i>	3437	1,47E+04	86%	0.0	83.90%	1095636	NW_015961132.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2948	<i>Capsicum annuum</i>	3092	1,51E+04	80%	0.0	91.40%	898408	NW_015961258.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1923	<i>Capsicum annuum</i>	2918	2,60E+04	91%	0.0	91.59%	2480791	NW_015961228.1
monotypic-Ch12a_45	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1647	<i>Capsicum annuum</i>	2569	1,56E+04	80%	0.0	92.31%	1362880	NW_015960892.1
monotypic-Ch12f_40	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	19023	1,12E+09	100%	0.0	99.99%	229934170	NC_029988.1
monotypic-Ch12f_40	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4460	9,91E+08	100%	0.0	85.82%	215701946	NC_029980.1
monotypic-Ch12f_40	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4202	1,20E+09	99%	0.0	84.86%	22112641	NC_029983.1
monotypic-Ch12f_40	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4045	1,38E+09	99%	0.0	84.04%	301019445	NC_029977.1
monotypic-Ch12f_40	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	4004	1,11E+09	100%	0.0	83.92%	238794889	NC_029985.1
monotypic-Ch12f_40	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3967	1,38E+09	100%	0.0	89.26%	261510930	NC_029979.1
monotypic-Ch12f_40	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3965	8,92E+08	99%	0.0	89.38%	220335243	NC_029987.1
monotypic-Ch12f_40	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3951	9,86E+08	100%	0.0	89.15%	217274494	NC_029981.1
monotypic-Ch12f_40	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3879	8,46E+08	99%	0.0	88.76%	219521584	NC_029982.1
monotypic-Ch12f_40	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3827	9,80E+08	99%	0.0	88.41%	205736368	NC_029986.1
monotypic-Ch12f_40	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3722	6,26E+08	100%	0.0	88.77%	153299543	NC_029984.1
monotypic-Ch12f_40	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	3195	7,47E+08	99%	0.0	86.69%	163962470	NC_029978.1
monotypic-Ch12f_40	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1647	<i>Capsicum annuum</i>	2715	8,82E+03	97%	0.0	78.85%	1362880	NW_015960892.1
monotypic-Ch12f_40	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold258	<i>Capsicum annuum</i>	2113	2,44E+04	92%	0.0	83.80%	3098259	NW_015961180.1
monotypic-Ch08d_60	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	22075	7,36E+08	100%	0.0	100.00%	153299543	NC_029984.1
monotypic-Ch08d_60	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	12074	1,31E+09	99%	0.0	90.55%	261510930	NC_029979.1
monotypic-Ch08d_60	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11815	9,66E+08	99%	0.0	90.53%	229934170	NC_029988.1
monotypic-Ch08d_60	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	11114	7,23E+08	99%	0.0	91.63%	22112641	NC_029983.1
monotypic-Ch08d_60	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10848	1,42E+09	99%	0.0	91.17%	301019445	NC_029977.1
monotypic-Ch08d_60	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	10815	7,53E+08	99%	0.0	89.06%	219521584	NC_029982.1
monotypic-Ch08d_60	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9969	7,64E+08	99%	0.0	91.38%	215701946	NC_029980.1
monotypic-Ch08d_60	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9821	5,09E+08	99%	0.0	90.72%	220335243	NC_029987.1
monotypic-Ch08d_60	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9777	6,53E+08	99%	0.0	88.28%	205736368	NC_029986.1
monotypic-Ch08d_60	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold756	<i>Capsicum annuum</i>	9587	1,80E+04	99%	0.0	86.09%	884456	NW_015961254.1
monotypic-Ch08d_60	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9474	8,80E+08	99%	0.0	90.42%	163962470	NC_029978.1
monotypic-Ch08d_60	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9474	7,50E+08	99%	0.0	87.40%	238794889	NC_029985.1
monotypic-Ch08d_60	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4259	<i>Capsicum annuum</i>	9369	2,39E+04	96%	0.0	87.94%	78069	NW_015961216.1

monotypic-Ch08d_60	refeq-genome	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	<i>Capsicum annuum</i>	9358	6,80E+08	99%	0.0	87.88%	217274494	NC_029981.1
monotypic-Ch08d_60	refeq-genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold949	<i>Capsicum annuum</i>	8885	1,98E+04	97%	0.0	91.06%	3128562	NW_015961487.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4739	<i>Capsicum annuum</i>	8517	1,97E+04	97%	0.0	86.07%	195495	NW_015961149.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9	<i>Capsicum annuum</i>	8484	4,35E+04	99%	0.0	85.35%	6567951	NW_015961815.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1894	<i>Capsicum annuum</i>	8466	2,20E+04	97%	0.0	85.86%	2334823	NW_015961490.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold53	<i>Capsicum annuum</i>	8388	5,23E+04	98%	0.0	86.21%	1681775	NW_015961414.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold658	<i>Capsicum annuum</i>	8371	2,78E+04	99%	0.0	85.67%	613851	NW_015961079.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1888	<i>Capsicum annuum</i>	8144	1,92E+04	91%	0.0	89.32%	361877	NW_015960979.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2793	<i>Capsicum annuum</i>	8089	2,28E+04	96%	0.0	84.34%	203362	NW_015961784.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3946	<i>Capsicum annuum</i>	7958	1,13E+04	82%	0.0	84.86%	591642	NW_015960367.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5146	<i>Capsicum annuum</i>	7919	2,35E+04	84%	0.0	86.14%	70278	NW_015960368.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2595	<i>Capsicum annuum</i>	7398	1,17E+04	85%	0.0	83.57%	161504	NW_015961232.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1923	<i>Capsicum annuum</i>	7282	1,10E+04	80%	0.0	83.27%	2480791	NW_015961228.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold585	<i>Capsicum annuum</i>	6191	1,56E+04	89%	0.0	87.01%	313752	NW_015961764.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold979	<i>Capsicum annuum</i>	6050	2,77E+04	98%	0.0	84.76%	1172749	NW_015961157.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2901	<i>Capsicum annuum</i>	5829	1,89E+04	96%	0.0	85.53%	1161691	NW_015961380.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2239	<i>Capsicum annuum</i>	5494	1,52E+04	90%	0.0	84.43%	615095	NW_015960816.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1390	<i>Capsicum annuum</i>	5376	1,81E+04	81%	0.0	86.76%	1730405	NW_015960783.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3499	<i>Capsicum annuum</i>	5299	1,66E+04	96%	0.0	85.69%	681687	NW_015960538.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1038	<i>Capsicum annuum</i>	5081	1,59E+04	86%	0.0	86.68%	2164954	NW_015960831.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold301	<i>Capsicum annuum</i>	4907	8,27E+03	85%	0.0	79.45%	1226190	NW_015960697.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold920	<i>Capsicum annuum</i>	4571	2,05E+04	97%	0.0	85.45%	197492	NW_015961621.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2265	<i>Capsicum annuum</i>	4549	1,23E+04	82%	0.0	86.91%	1257389	NW_015961797.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2168	<i>Capsicum annuum</i>	3903	1,77E+04	85%	0.0	90.19%	89773	NW_015961676.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4518	<i>Capsicum annuum</i>	3705	2,29E+04	87%	0.0	92.19%	88275	NW_015961472.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1384	<i>Capsicum annuum</i>	3267	9,99E+03	81%	0.0	86.29%	533800	NW_015961668.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4257	<i>Capsicum annuum</i>	3057	1,65E+04	85%	0.0	91.22%	106013	NW_015961030.1
monotypic-Ch08d_60	refeq_genome	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	<i>Capsicum annuum</i>	3005	1,11E+04	83%	0.0	83.38%	3493078	NW_015960517.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold900	<i>Nicotiana tabacum</i>	9660	11886	100%	0.0	99.36%	218302	NW_015947482.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold24962	<i>Nicotiana sylvestris</i>	9625	11772	100%	0.0	99.27%	119121	NW_009505047.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold47941	<i>Nicotiana sylvestris</i>	9609	11753	100%	0.0	99.19%	106539	NW_009534877.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold43408	<i>Nicotiana tabacum</i>	9598	1,17E+04	100%	0.0	99.16%	70758	NW_015906690.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold20959	<i>Nicotiana tabacum</i>	9513	12014	100%	0.0	99.20%	211562	NW_015856489.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold941	<i>Nicotiana tabacum</i>	8957	1,13E+04	100%	0.0	98.56%	71070	NW_015950805.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold14911	<i>Nicotiana sylvestris</i>	8780	11966	100%	0.0	99.22%	94838	NW_009393370.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold23491	<i>Nicotiana sylvestris</i>	8780	12030	100%	0.0	99.22%	128539	NW_009488703.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7035	<i>Nicotiana tabacum</i>	8754	1,20E+04	100%	0.0	99.12%	113245	NW_015930881.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold888	<i>Nicotiana sylvestris</i>	8730	11881	100%	0.0	99.04%	85608	NW_009580274.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold8912	<i>Nicotiana sylvestris</i>	8691	11737	100%	0.0	98.89%	302464	NW_009580631.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold26600	<i>Nicotiana sylvestris</i>	8634	22006	100%	0.0	99.05%	221681	NW_009511166.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold64168	<i>Nicotiana sylvestris</i>	8619	1,67E+04	100%	0.0	98.98%	22405	NW_009552907.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold22238	<i>Nicotiana tabacum</i>	8597	21291	100%	0.0	98.90%	80703	NW_015862602.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5181	<i>Nicotiana tabacum</i>	8591	12040	100%	0.0	98.88%	94702	NW_015914389.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold6826	<i>Nicotiana sylvestris</i>	8543	1,12E+04	100%	0.0	98.55%	152392	NW_009557453.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold20247	<i>Nicotiana tabacum</i>	8423	1,13E+04	100%	0.0	98.26%	62189	NW_015852854.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold71206	<i>Nicotiana tabacum</i>	8298	11039	100%	0.0	99.18%	66708	NW_015931638.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1275	<i>Nicotiana sylvestris</i>	8268	19652	100%	0.0	98.09%	109494	NW_009369358.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold44244	<i>Nicotiana sylvestris</i>	8261	10507	100%	0.0	97.99%	88672	NW_009530770.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17315	<i>Nicotiana tabacum</i>	8070	1,08E+04	96%	0.0	99.28%	67960	NW_015836956.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold18407	<i>Nicotiana sylvestris</i>	7949	1,05E+04	100%	0.0	96.23%	59561	NW_009432214.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold20433	<i>Nicotiana tabacum</i>	7919	10962	100%	0.0	96.37%	144898	NW_015853831.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold43464	<i>Nicotiana tabacum</i>	7890	10487	100%	0.0	95.98%	51581	NW_015906741.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold781	<i>Nicotiana tabacum</i>	7843	1,05E+04	100%	0.0	96.12%	50117	NW_015937551.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold2145	<i>Nicotiana sylvestris</i>	7838	17810	100%	0.0	97.26%	71199	NW_009466025.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10934	<i>Nicotiana tabacum</i>	7334	14335	90%	0.0	96.25%	156973	NW_015794434.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold14856	<i>Nicotiana sylvestris</i>	7260	8,14E+03	90%	0.0	96.01%	193910	NW_009392758.1

Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 1, NIATTr2	Nicotiana attenuata	7188	29176	92%	0.0	95.29%	110538649	NC_031989.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold8378	Nicotiana tomentosiformis	7047	8372	99%	0.0	90.59%	108988	NW_008970018.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold40995	Nicotiana tomentosiformis	7035	9896	100%	0.0	90.82%	19427	NW_008922480.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	7033	8,84E+03	92%	0.0	94.70%	216675	NW_017672129.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold38738	Nicotiana tabacum	6992	8217	100%	0.0	90.34%	232414	NW_015902370.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10539	Nicotiana tabacum	6986	8278	99%	0.0	90.50%	92002	NW_015791427.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5911	Nicotiana tomentosiformis	6953	8167	100%	0.0	90.21%	517654	NW_008942608.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 3, NIATTr2	Nicotiana attenuata	6931	31512	94%	0.0	94.28%	72893213	NC_031991.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold43675	Nicotiana tabacum	6926	8286	90%	0.0	94.59%	101217	NW_015906932.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold6313	Nicotiana tomentosiformis	6926	8132	99%	0.0	90.16%	122694	NW_008947075.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold24995	Nicotiana sylvestris	6889	8541	91%	0.0	94.44%	101192	NW_009505413.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5983	Nicotiana tabacum	6885	8089	99%	0.0	90.13%	75022	NW_015921626.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold20220	Nicotiana tabacum	6854	7966	100%	0.0	89.87%	74730	NW_015852702.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold2163	Nicotiana sylvestris	6830	10796	100%	0.0	98.73%	53555	NW_009468025.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1033	Nicotiana tomentosiformis	6820	8038	99%	0.0	89.96%	196095	NW_008832160.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold41709	Nicotiana tabacum	6804	8032	99%	0.0	89.80%	75821	NW_015905124.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold115329	Nicotiana tabacum	6793	7276	89%	0.0	94.51%	25574	NW_015798935.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13838	Nicotiana tomentosiformis	6785	12996	99%	0.0	89.90%	179143	NW_008871135.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold15913	Nicotiana tabacum	6772	20314	100%	0.0	90.29%	302364	NW_015828652.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13941	Nicotiana tabacum	6765	7960	99%	0.0	89.77%	189720	NW_015815959.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10888	Nicotiana tabacum	6763	7990	99%	0.0	89.65%	69340	NW_015794085.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2507	Nicotiana tomentosiformis	6761	7966	99%	0.0	90.05%	150405	NW_008904786.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold75131	Nicotiana sylvestris	6754	8297	90%	0.0	93.84%	25769	NW_009565089.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold52439	Nicotiana tabacum	6734	8277	91%	0.0	93.73%	28490	NW_015914967.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold12665	Nicotiana sylvestris	6722	10431	98%	0.0	94.83%	62516	NW_009368414.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold47967	Nicotiana tabacum	6719	7953	99%	0.0	90.05%	136831	NW_015910891.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold11358	Nicotiana tabacum	6713	7925	99%	0.0	89.63%	29208	NW_015797613.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold49078	Nicotiana tomentosiformis	6706	7886	99%	0.0	90.02%	70781	NW_008931461.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold44401	Nicotiana sylvestris	6702	7975	88%	0.0	97.00%	85935	NW_009530945.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold77318	Nicotiana tabacum	6691	8070	90%	0.0	93.67%	72032	NW_015936870.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16478	Nicotiana tomentosiformis	6689	7921	99%	0.0	89.98%	70954	NW_008895239.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold50494	Nicotiana tabacum	6676	8170	91%	0.0	93.50%	60219	NW_015913188.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold39877	Nicotiana sylvestris	6669	8170	92%	0.0	93.31%	22882	NW_009525916.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold53158	Nicotiana tabacum	6665	8740	87%	0.0	96.39%	49428	NW_015915634.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold19127	Nicotiana tomentosiformis	6663	7907	99%	0.0	89.89%	81528	NW_008898183.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold136633	Nicotiana tabacum	6660	12342	99%	0.0	89.87%	49476	NW_015814088.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold52405	Nicotiana tabacum	6639	7530	99%	0.0	90.47%	79607	NW_015914937.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold22705	Nicotiana tomentosiformis	6626	7519	99%	0.0	90.25%	149011	NW_008902159.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold18357	Nicotiana tomentosiformis	6625	7792	100%	0.0	89.17%	18837	NW_008897327.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold45081	Nicotiana sylvestris	6623	8359	90%	0.0	93.39%	9376	NW_009531700.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold25368	Nicotiana tomentosiformis	6623	8119	100%	0.0	90.94%	144419	NW_008905117.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold24121	Nicotiana sylvestris	6615	7978	90%	0.0	93.35%	108052	NW_009495704.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold25494	Nicotiana tabacum	6612	8103	100%	0.0	90.90%	88174	NW_015877587.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold2989	Nicotiana sylvestris	6608	8009	91%	0.0	93.24%	34952	NW_009514819.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold4930	Nicotiana sylvestris	6606	9655	98%	0.0	93.33%	105967	NW_009536387.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold11177	Nicotiana tabacum	6604	8140	90%	0.0	93.30%	124530	NW_015796267.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16180	Nicotiana tomentosiformis	6604	7770	99%	0.0	89.79%	24397	NW_008894909.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	6602	8220	91%	0.0	93.12%	103397	NW_017674260.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold3719	Nicotiana tabacum	6601	7591	91%	0.0	93.22%	35027	NW_015900902.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7405	Nicotiana tabacum	6591	7976	91%	0.0	93.17%	56606	NW_015934073.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold68652	Nicotiana tabacum	6586	7759	99%	0.0	89.75%	72869	NW_015929427.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold76626	Nicotiana sylvestris	6586	7807	88%	0.0	96.78%	6175	NW_009566749.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold3226	Nicotiana tabacum	6584	7786	99%	0.0	89.32%	133034	NW_015890178.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold94042	Nicotiana tabacum	6584	8039	90%	0.0	93.24%	74818	NW_015950751.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold112542	Nicotiana tabacum	6577	7663	99%	0.0	89.96%	41180	NW_015796842.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold18637	Nicotiana tabacum	6565	8087	91%	0.0	93.02%	62391	NW_015844334.1

Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl_Nsyl_scaffold32482	Nicotiana sylvestris	6558	8015	91%	0.0	93.04%	116552	NW_009517701.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold20231	Nicotiana tabacum	6547	7399	99%	0.0	90.94%	7831	NW_015852764.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold6303	Nicotiana tabacum	6547	7729	99%	0.0	89.47%	119331	NW_015924496.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 6, NIATT2	Nicotiana attenuata	6545	74074	99%	0.0	93.04%	97571839	NC_031994.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl_Nsyl_scaffold4249	Nicotiana sylvestris	6529	7952	91%	0.0	92.96%	355138	NW_009528820.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 2, NIATT2	Nicotiana attenuata	6525	46561	94%	0.0	92.57%	96359640	NC_031990.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl_Nsyl_scaffold11822	Nicotiana sylvestris	6525	16225	91%	0.0	92.89%	232274	NW_009359048.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl_Nsyl_scaffold18120	Nicotiana sylvestris	6523	8196	94%	0.0	92.39%	190068	NW_009429026.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl_Nsyl_scaffold45677	Nicotiana sylvestris	6516	8514	91%	0.0	92.87%	113330	NW_009532361.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold24794	Nicotiana tomentosiformis	6516	8286	100%	0.0	91.15%	24987	NW_008904479.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold108203	Nicotiana tabacum	6512	8144	94%	0.0	92.56%	35212	NW_015793576.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold59508	Nicotiana tabacum	6506	7143	99%	0.0	91.01%	109704	NW_015921346.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold10551	Nicotiana tomentosiformis	6499	7574	97%	0.0	89.69%	74828	NW_008834616.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13993	Nicotiana tomentosiformis	6493	7683	99%	0.0	89.34%	172161	NW_008872857.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2558	Nicotiana tomentosiformis	6488	8319	92%	0.0	92.62%	65600	NW_008905352.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl_Nsyl_scaffold211	Nicotiana sylvestris	6482	8320	90%	0.0	92.83%	200149	NW_009462136.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl_Nsyl_scaffold16678	Nicotiana sylvestris	6479	7497	84%	0.0	97.43%	25570	NW_009413002.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold6678	Nicotiana tomentosiformis	6475	7609	99%	0.0	89.38%	209558	NW_008951129.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16653	Nicotiana tomentosiformis	6473	8449	99%	0.0	91.14%	68796	NW_008895434.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold28741	Nicotiana tabacum	6471	8325	99%	0.0	90.51%	30281	NW_015886859.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold78610	Nicotiana tabacum	6471	7679	99%	0.0	89.50%	26679	NW_015937987.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl_Nsyl_scaffold169896	Nicotiana sylvestris	6469	6879	91%	0.0	92.79%	9898	NW_009416464.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	6464	8074	91%	0.0	92.54%	586478	NW_017670551.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl_Nsyl_scaffold8223	Nicotiana sylvestris	6462	7937	90%	0.0	92.84%	102633	NW_009572976.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold12432	Nicotiana tabacum	6457	8520	99%	0.0	90.10%	124037	NW_015805427.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold31651	Nicotiana tabacum	6451	7915	91%	0.0	92.62%	52290	NW_015889605.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16694	Nicotiana tomentosiformis	6444	7748	93%	0.0	92.36%	25472	NW_008895479.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1183	Nicotiana tabacum	6436	7580	97%	0.0	89.23%	159099	NW_015801081.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	6421	8009	93%	0.0	92.09%	182132	NW_017672588.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13088	Nicotiana tomentosiformis	6420	7,55E+03	97%	0.0	89.16%	176428	NW_008862802.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold18452	Nicotiana tomentosiformis	6407	1,55E+04	100%	0.0	90.31%	77122	NW_008897433.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl_Nsyl_scaffold6510	Nicotiana sylvestris	6405	7,44E+03	90%	0.0	92.64%	107145	NW_009553943.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold103502	Nicotiana tabacum	6397	7,95E+03	99%	0.0	90.61%	39113	NW_015789959.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7978	Nicotiana tabacum	6396	8,19E+03	97%	0.0	94.71%	8160	NW_01593894.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold31852	Nicotiana tabacum	6386	7,10E+03	95%	0.0	92.42%	71837	NW_015889792.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold36118	Nicotiana tabacum	6381	8,10E+03	99%	0.0	90.49%	70952	NW_015899912.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl_Nsyl_scaffold29717	Nicotiana sylvestris	6381	15459	100%	0.0	94.52%	70310	NW_009514628.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 8, NIATT2	Nicotiana attenuata	6368	1,69E+04	98%	0.0	91.89%	47755514	NC_031996.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold34701	Nicotiana tabacum	6368	7196	99%	0.0	90.37%	75498	NW_015893859.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold37420	Nicotiana tabacum	6366	7,87E+03	91%	0.0	92.23%	164383	NW_015901126.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	6357	7893	91%	0.0	92.32%	335605	NW_017671207.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7194	Nicotiana tabacum	6351	7,53E+03	99%	0.0	88.61%	40111	NW_015932262.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	6335	7917	92%	0.0	91.85%	203160	NW_017672282.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31562	Nicotiana tomentosiformis	6333	8113	96%	0.0	91.20%	31162	NW_008912000.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	6331	6,88E+03	89%	0.0	92.77%	90879	NW_017674616.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold39248	Nicotiana tabacum	6327	7933	92%	0.0	91.96%	46196	NW_015902840.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13971	Nicotiana tomentosiformis	6327	21985	99%	0.0	90.06%	146979	NW_008872613.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold117472	Nicotiana tabacum	6316	7608	99%	0.0	89.99%	111611	NW_015800473.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	6305	7797	93%	0.0	91.67%	105537	NW_017674199.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold10777	Nicotiana tomentosiformis	6305	7882	94%	0.0	91.52%	258869	NW_008837126.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold73542	Nicotiana tabacum	6298	7788	90%	0.0	92.09%	16972	NW_015933632.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold40269	Nicotiana tomentosiformis	6298	6,92E+03	95%	0.0	92.11%	26048	NW_008921674.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold7307	Nicotiana tomentosiformis	6298	8,07E+03	99%	0.0	90.82%	59780	NW_008958119.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold112002	Nicotiana tabacum	6296	8,09E+03	99%	0.0	90.38%	51612	NW_015796444.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold65311	Nicotiana tabacum	6296	8159	99%	0.0	90.53%	132053	NW_015926505.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold7348	Nicotiana tomentosiformis	6294	8,28E+03	99%	0.0	89.48%	162329	NW_008958574.1

Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold65340	Nicotiana tabacum	6285	7756	92%	0.0	91.73%	35831	NW_015926530.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	6283	7762	93%	0.0	91.56%	415554	NW_017670898.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold3972	Nicotiana sylvestris	6279	7,84E+03	91%	0.0	92.00%	214752	NW_009525742.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3451	Nicotiana tomentosiformis	6279	8074	99%	0.0	90.32%	90093	NW_008915273.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold25240	Nicotiana tabacum	6277	7,88E+03	92%	0.0	91.71%	102696	NW_015876456.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1620	Nicotiana tomentosiformis	6277	7765	91%	0.0	91.90%	135579	NW_008894931.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold14749	Nicotiana tomentosiformis	6266	7096	97%	0.0	90.00%	71908	NW_008881257.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold59532	Nicotiana tabacum	6264	7719	91%	0.0	91.72%	111948	NW_015921368.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold30594	Nicotiana tabacum	6252	7852	96%	0.0	90.92%	92652	NW_01588604.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold29104	Nicotiana tabacum	6239	6720	91%	0.0	91.64%	4893	NW_015887196.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4588	Nicotiana tomentosiformis	6235	1,25E+04	89%	0.0	91.95%	93963	NW_008927907.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	6233	7,75E+03	92%	0.0	91.54%	370422	NW_017671056.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1530	Nicotiana sylvestris	6222	7251	82%	0.0	97.13%	173340	NW_009397692.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold43162	Nicotiana sylvestris	6222	8,37E+03	81%	0.0	96.96%	77093	NW_009529568.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	6216	6,22E+03	83%	0.0	91.92%	69192	NW_017675359.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9966	Nicotiana tabacum	6209	6,70E+03	80%	0.0	95.17%	45944	NW_015955195.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9431	Nicotiana tabacum	6205	7,72E+03	92%	0.0	91.48%	218470	NW_015950970.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13072	Nicotiana tabacum	6202	9,06E+03	91%	0.0	93.76%	42540	NW_015809998.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold81708	Nicotiana tabacum	6202	7,74E+03	91%	0.0	91.51%	129650	NW_015940616.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold11093	Nicotiana tabacum	6196	7,84E+03	99%	0.0	90.13%	28843	NW_015795636.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold129157	Nicotiana sylvestris	6192	7621	91%	0.0	91.64%	28956	NW_009371200.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold24190	Nicotiana tomentosiformis	6192	7,77E+03	93%	0.0	91.44%	30170	NW_008903089.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold40460	Nicotiana tomentosiformis	6192	7826	99%	0.0	90.13%	19217	NW_008921887.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold6079	Nicotiana sylvestris	6189	8258	89%	0.0	95.61%	533592	NW_009549153.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	6187	6536	92%	0.0	91.34%	228550	NW_017671987.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold6958	Nicotiana tabacum	6187	7,62E+03	91%	0.0	91.62%	23504	NW_015930225.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13421	Nicotiana tomentosiformis	6183	7,69E+03	91%	0.0	91.44%	128956	NW_008866503.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold28546	Nicotiana tomentosiformis	6180	7771	92%	0.0	91.29%	209861	NW_008908648.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold57259	Nicotiana tomentosiformis	6168	7980	99%	0.0	89.76%	34918	NW_008940551.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold15504	Nicotiana tabacum	6165	7632	91%	0.0	91.39%	101907	NW_015826108.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold35867	Nicotiana tomentosiformis	6159	7,62E+03	91%	0.0	91.32%	55044	NW_008916782.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold22374	Nicotiana tomentosiformis	6144	7,69E+03	99%	0.0	91.28%	114813	NW_008901791.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24408	Nicotiana tabacum	6137	7616	94%	0.0	90.92%	154488	NW_015872748.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold57738	Nicotiana sylvestris	6137	8,07E+03	96%	0.0	90.59%	29438	NW_009545762.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold22332	Nicotiana tabacum	6131	7592	80%	0.0	96.95%	133198	NW_015863028.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold94372	Nicotiana tabacum	6131	1,49E+04	91%	0.0	91.23%	78974	NW_015951020.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold68555	Nicotiana sylvestris	6130	9832	89%	0.0	97.78%	34226	NW_009557781.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold65799	Nicotiana tabacum	6128	7440	95%	0.0	91.21%	102478	NW_015926924.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold319	Nicotiana tomentosiformis	6128	7579	90%	0.0	91.46%	142124	NW_008912374.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4258	Nicotiana tomentosiformis	6126	7,54E+03	92%	0.0	91.06%	67558	NW_008924241.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold21998	Nicotiana tomentosiformis	6124	7,62E+03	91%	0.0	91.23%	185413	NW_008901372.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold102188	Nicotiana tabacum	6107	7,06E+03	92%	0.0	91.57%	37240	NW_015788939.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold51668	Nicotiana tabacum	6102	7195	99%	0.0	89.86%	84183	NW_015914252.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8505	Nicotiana tabacum	6089	7536	92%	0.0	90.87%	215517	NW_015943393.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold12955	Nicotiana tabacum	6085	7518	90%	0.0	91.30%	98519	NW_015809166.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold23616	Nicotiana tomentosiformis	6080	6,49E+03	89%	0.0	91.32%	80310	NW_008903171.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold8589	Nicotiana tomentosiformis	6078	7466	92%	0.0	90.99%	77132	NW_008972362.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	6072	7509	92%	0.0	90.82%	201149	NW_017672308.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	6072	18955	91%	0.0	91.19%	165435	NW_017672850.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	6072	6776	90%	0.0	91.19%	18689	NW_017679016.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold60854	Nicotiana tabacum	6069	6552	87%	0.0	92.09%	91511	NW_015922551.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5690	Nicotiana tomentosiformis	6069	7616	94%	0.0	90.60%	307960	NW_008940152.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9842	Nicotiana tomentosiformis	6069	7496	90%	0.0	91.24%	137779	NW_008986285.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13004	Nicotiana tabacum	6065	7319	89%	0.0	91.35%	129879	NW_015809526.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold16837	Nicotiana tabacum	6050	7668	93%	0.0	90.96%	100875	NW_015834189.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold30081	Nicotiana tomentosiformis	6050	7078	92%	0.0	91.19%	52569	NW_008910355.1

Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold49510	Nicotiana tabacum	6048	7960	91%	0.0	93.94%	9982	NW_015912287.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	6047	6,05E+03	83%	0.0	91.28%	539978	NW_017670619.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold21380	Nicotiana tomentosiformis	6045	7289	89%	0.0	91.24%	111797	NW_008900687.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31363	Nicotiana tomentosiformis	6041	6524	87%	0.0	91.97%	29413	NW_008911779.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 10, NIATTr2	Nicotiana attenuata	6039	1,98E+04	100%	0.0	91.02%	51082206	NC_031998.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold26705	Nicotiana tomentosiformis	6039	6,48E+03	89%	0.0	91.32%	41955	NW_008906603.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold100338	Nicotiana tabacum	6037	7398	92%	0.0	90.84%	73501	NW_015787497.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold27552	Nicotiana tabacum	6037	7118	97%	0.0	88.00%	82479	NW_015885731.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold38662	Nicotiana sylvestris	6037	7118	97%	0.0	88.00%	84807	NW_009524567.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold8433	Nicotiana tomentosiformis	6037	7241	89%	0.0	91.24%	146443	NW_008970630.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold12236	Nicotiana tabacum	6035	6035	83%	0.0	91.28%	139476	NW_015804041.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	6034	6332	89%	0.0	91.16%	435117	NW_017670843.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold14276	Nicotiana tomentosiformis	6032	7,37E+03	99%	0.0	89.25%	50827	NW_008876002.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold11122	Nicotiana tabacum	6030	7012	91%	0.0	91.16%	131331	NW_015795856.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31112	Nicotiana tomentosiformis	6030	7548	95%	0.0	90.80%	102337	NW_008911501.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold56126	Nicotiana tabacum	6026	7,95E+03	96%	0.0	90.32%	78728	NW_015918329.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold68745	Nicotiana tabacum	6017	7289	89%	0.0	91.18%	104675	NW_015929511.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold554	Nicotiana tomentosiformis	6015	7769	94%	0.0	90.51%	238694	NW_008938485.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold16073	Nicotiana tabacum	6010	7586	90%	0.0	92.34%	203079	NW_015829650.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold95853	Nicotiana tabacum	6010	7553	95%	0.0	90.71%	89290	NW_015952201.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold30202	Nicotiana tabacum	6008	7584	94%	0.0	90.35%	51425	NW_015888233.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1912	Nicotiana tomentosiformis	6008	7363	91%	0.0	90.84%	133118	NW_008898175.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold183409	Nicotiana tabacum	5995	7789	97%	0.0	89.96%	26684	NW_015842704.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold34464	Nicotiana tomentosiformis	5995	7404	92%	0.0	90.50%	45579	NW_008915224.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1130	Nicotiana tabacum	5982	8320	92%	0.0	90.53%	356733	NW_015797183.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13774	Nicotiana tomentosiformis	5980	7324	92%	0.0	90.96%	86629	NW_008870424.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	5973	7,53E+03	91%	0.0	90.82%	216689	NW_017672128.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold35281	Nicotiana tabacum	5967	7,70E+03	94%	0.0	90.32%	102698	NW_015899120.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold8132	Nicotiana tomentosiformis	5965	6947	91%	0.0	90.86%	161484	NW_008967286.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold32792	Nicotiana tabacum	5963	15434	91%	0.0	90.62%	39461	NW_015890673.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	5962	6182	92%	0.0	90.70%	205284	NW_017672254.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold20034	Nicotiana tomentosiformis	5962	6526	95%	0.0	89.93%	23404	NW_008899192.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold81582	Nicotiana tabacum	5958	6511	95%	0.0	89.94%	18140	NW_015940515.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold23869	Nicotiana tomentosiformis	5956	7476	94%	0.0	90.19%	97209	NW_008903451.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold914	Nicotiana tomentosiformis	5952	15554	91%	0.0	90.58%	109301	NW_008978485.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold8583	Nicotiana tomentosiformis	5949	7717	94%	0.0	90.21%	262523	NW_008972296.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold11214	Nicotiana tabacum	5947	7553	91%	0.0	90.93%	87770	NW_015796546.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold32936	Nicotiana tabacum	5945	7424	90%	0.0	90.58%	15184	NW_015890812.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold48732	Nicotiana tabacum	5945	7425	91%	0.0	90.54%	87162	NW_015911588.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold4431	Nicotiana sylvestris	5945	7415	91%	0.0	90.70%	157602	NW_009530843.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold2056	Nicotiana tabacum	5941	1,11E+04	99%	0.0	88.93%	58443	NW_015854471.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold97038	Nicotiana tabacum	5941	7202	89%	0.0	90.79%	118855	NW_015953126.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold19877	Nicotiana tomentosiformis	5941	6820	91%	0.0	91.73%	75032	NW_008899015.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17472	Nicotiana tabacum	5938	6495	99%	0.0	89.50%	344968	NW_015837843.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold33946	Nicotiana tabacum	5938	7,40E+03	90%	0.0	90.63%	68701	NW_015891773.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold14944	Nicotiana tomentosiformis	5930	7415	90%	0.0	90.59%	151478	NW_008883424.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23242	Nicotiana tabacum	5928	6791	90%	0.0	91.73%	354401	NW_015867302.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold107739	Nicotiana tabacum	5914	7935	90%	0.0	94.75%	18276	NW_015793205.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold26686	Nicotiana sylvestris	5914	9325	91%	0.0	94.29%	13361	NW_009511260.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1536	Nicotiana tomentosiformis	5910	7367	91%	0.0	90.49%	125125	NW_00888046.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16063	Nicotiana tomentosiformis	5910	6437	94%	0.0	90.13%	74560	NW_008894779.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold19286	Nicotiana tomentosiformis	5904	6989	94%	0.0	90.10%	173816	NW_008898359.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23242	Nicotiana tabacum	5903	7660	94%	0.0	90.22%	135222	NW_015918199.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold26686	Nicotiana tomentosiformis	5901	7493	91%	0.0	90.73%	87833	NW_008833582.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold10458	Nicotiana tomentosiformis	5901	6325	90%	0.0	90.47%	55036	NW_008984785.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9707	Nicotiana tabacum	5899	7287	90%	0.0	90.50%	76405	NW_015934850.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold74949								

Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold42735	Nicotiana tabacum	5897	7359	96%	0.0	90.46%	67689	NW_015906063.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold65632	Nicotiana tabacum	5895	7522	99%	0.0	89.07%	35386	NW_015926779.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold8635	Nicotiana tomentosiformis	5891	7459	92%	0.0	90.32%	95887	NW_008972874.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold40660	Nicotiana tomentosiformis	5890	6,92E+03	90%	0.0	90.36%	15493	NW_008922109.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold20595	Nicotiana tomentosiformis	5882	7046	95%	0.0	89.71%	37803	NW_008899814.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold39505	Nicotiana tabacum	5877	7171	87%	0.0	91.04%	194374	NW_015903077.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7593	Nicotiana tabacum	5871	6,94E+03	94%	0.0	89.97%	214177	NW_015935691.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16874	Nicotiana tomentosiformis	5871	7250	86%	0.0	91.71%	50659	NW_008895679.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold19276	Nicotiana tomentosiformis	5867	7152	89%	0.0	90.46%	112595	NW_008898348.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold61422	Nicotiana tabacum	5860	7834	92%	0.0	90.07%	11815	NW_008945178.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold64725	Nicotiana tabacum	5858	14713	91%	0.0	90.37%	256583	NW_015953907.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9802	Nicotiana tabacum	5856	7,29E+03	90%	0.0	90.39%	42538	NW_015848864.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold19469	Nicotiana tomentosiformis	5853	7197	90%	0.0	90.35%	43464	NW_008917316.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold36347	Nicotiana tabacum	5849	6,98E+03	89%	0.0	90.46%	70655	NW_015920349.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold58364	Nicotiana tomentosiformis	5847	7235	91%	0.0	90.19%	60363	NW_008895788.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16972	Nicotiana tomentosiformis	5847	7459	91%	0.0	90.28%	24557	NW_008912438.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31957	Nicotiana tomentosiformis	5830	8816	98%	0.0	90.22%	55974	NW_008911657.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31253	Nicotiana tabacum	5829	8226	89%	0.0	94.80%	120278	NW_015845192.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold18791	Nicotiana sylvestris	5821	9873	91%	0.0	95.21%	205743	NW_009385148.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold14171	Nicotiana tomentosiformis	5821	6288	92%	0.0	90.04%	29233	NW_008915006.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold34268	Nicotiana tomentosiformis	5819	10293	91%	0.0	90.05%	9089	NW_008945582.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold61787	Nicotiana tabacum	5810	6321	93%	0.0	89.76%	10998	NW_015873178.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24504	Nicotiana tabacum	5810	6,25E+03	89%	0.0	90.36%	71959	NW_015882390.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold2650	Nicotiana tabacum	5810	1,03E+04	92%	0.0	89.99%	51409	NW_015910398.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold21103	Nicotiana tomentosiformis	5801	6284	92%	0.0	89.89%	20769	NW_008900380.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3799	Nicotiana tomentosiformis	5799	7495	91%	0.0	91.82%	179455	NW_008919140.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2563	Nicotiana tomentosiformis	5794	7245	91%	0.0	90.02%	79364	NW_008905408.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold6751	Nicotiana tabacum	5790	7051	90%	0.0	90.12%	155870	NW_015928428.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5691	Nicotiana tomentosiformis	5786	7060	87%	0.0	90.72%	262724	NW_008940163.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold3770	Nicotiana tabacum	5782	7239	91%	0.0	89.98%	230784	NW_015901390.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4405	Nicotiana tabacum	5782	6830	94%	0.0	90.59%	116927	NW_015907286.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold25046	Nicotiana tomentosiformis	5781	8032	93%	0.0	92.25%	46165	NW_008904760.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold48093	Nicotiana tabacum	5777	6256	92%	0.0	89.70%	149912	NW_015911012.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9945	Nicotiana tabacum	5777	8,37E+03	96%	0.0	92.14%	48410	NW_015955036.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2417	Nicotiana tomentosiformis	5775	7241	90%	0.0	90.14%	275886	NW_008903786.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8507	Nicotiana tabacum	5766	6909	89%	0.0	90.90%	109085	NW_015943412.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold70944	Nicotiana sylvestris	5766	6256	92%	0.0	89.66%	77520	NW_009560436.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold171697	Nicotiana tabacum	5764	6765	89%	0.0	90.34%	53640	NW_015836116.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold21543	Nicotiana sylvestris	5753	10092	93%	0.0	89.49%	94545	NW_009467059.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4375	Nicotiana tomentosiformis	5753	6896	91%	0.0	90.36%	269466	NW_008925541.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTTr2	Nicotiana attenuata	5744	6626	89%	0.0	90.20%	398809	NW_017670952.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1008	Nicotiana tabacum	5738	10129	93%	0.0	89.43%	132392	NW_015787862.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5906	Nicotiana tabacum	5738	6122	91%	0.0	89.71%	229466	NW_015920954.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTTr2	Nicotiana attenuata	5733	7223	92%	0.0	89.81%	209876	NW_017672210.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold63112	Nicotiana tabacum	5731	7382	99%	0.0	91.20%	69490	NW_015924570.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold17962	Nicotiana tomentosiformis	5727	7231	99%	0.0	90.18%	144742	NW_008896888.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31593	Nicotiana tomentosiformis	5727	6786	94%	0.0	90.36%	58551	NW_008912034.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold18476	Nicotiana tomentosiformis	5716	6678	89%	0.0	90.14%	123680	NW_008897459.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24586	Nicotiana tabacum	5701	6855	91%	0.0	89.68%	22424	NW_015873563.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold50860	Nicotiana tabacum	5699	6724	88%	0.0	90.68%	87660	NW_015913518.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold59022	Nicotiana tabacum	5696	6496	92%	0.0	89.73%	42735	NW_015920924.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13171	Nicotiana tomentosiformis	5696	6243	85%	0.0	91.78%	137691	NW_008863725.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4794	Nicotiana tomentosiformis	5692	6164	91%	0.0	90.34%	80393	NW_008930196.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold51014	Nicotiana tabacum	5690	8162	90%	0.0	93.17%	28040	NW_015913659.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold58798	Nicotiana tomentosiformis	5688	6112	90%	0.0	89.77%	20384	NW_008942260.1

Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	5686	5686	83%	0.0	90.00%	609028	NW_017670521.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9832	Nicotiana tomentosiformis	5686	5686	85%	0.0	89.34%	40427	NW_008986174.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold71883	Nicotiana tabacum	5674	6221	85%	0.0	91.68%	135761	NW_015932208.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold64451	Nicotiana tabacum	5668	7324	88%	0.0	91.47%	83433	NW_015925768.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold6062	Nicotiana tomentosiformis	5668	7,50E+03	85%	0.0	91.37%	76416	NW_008944286.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9587	Nicotiana tomentosiformis	5668	6457	92%	0.0	89.77%	93896	NW_008983451.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold18106	Nicotiana tomentosiformis	5666	7139	91%	0.0	89.55%	22471	NW_008897049.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold104982	Nicotiana tabacum	5653	5653	83%	0.0	89.63%	16757	NW_015791105.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	5650	6885	89%	0.0	89.65%	30464	NW_017677444.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold7520	Nicotiana sylvestris	5648	8173	81%	0.0	95.83%	103998	NW_009565165.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4575	Nicotiana tabacum	5644	6253	84%	0.0	91.74%	193566	NW_015908835.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold29919	Nicotiana tabacum	5637	6977	92%	0.0	89.10%	18620	NW_015887964.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold18396	Nicotiana tabacum	5635	6059	90%	0.0	89.45%	56899	NW_015843010.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold8530	Nicotiana sylvestris	5635	9314	90%	0.0	94.83%	41440	NW_009576387.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold37079	Nicotiana tomentosiformis	5622	9333	83%	0.0	91.63%	29602	NW_008918129.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	5616	6293	89%	0.0	89.59%	65724	NW_017675503.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold147459	Nicotiana tabacum	5613	7058	85%	0.0	91.13%	15215	NW_015821291.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold160715	Nicotiana tabacum	5609	6406	83%	0.0	89.66%	29644	NW_015829643.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold70774	Nicotiana tabacum	5609	6936	92%	0.0	89.20%	46303	NW_015931260.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold36958	Nicotiana tabacum	5605	8969	83%	0.0	91.55%	25852	NW_015900682.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold4652	Nicotiana sylvestris	5603	6942	92%	0.0	89.18%	141735	NW_009533298.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5389	Nicotiana tomentosiformis	5581	7110	83%	0.0	91.33%	257796	NW_008936807.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31166	Nicotiana tomentosiformis	5579	8000	99%	0.0	90.78%	82530	NW_008911560.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold65714	Nicotiana sylvestris	5578	6,97E+03	91%	0.0	89.20%	93095	NW_009554625.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold41868	Nicotiana tabacum	5574	7,01E+03	92%	0.0	88.97%	32126	NW_015905266.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold81051	Nicotiana tabacum	5574	9941	97%	0.0	95.66%	19034	NW_015940058.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold50066	Nicotiana tabacum	5572	7707	90%	0.0	91.85%	19561	NW_015912806.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold93104	Nicotiana tabacum	5572	6971	91%	0.0	89.18%	82882	NW_015950030.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10248	Nicotiana tabacum	5570	5570	84%	0.0	89.09%	61358	NW_015789168.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24280	Nicotiana tabacum	5568	7741	91%	0.0	89.13%	52569	NW_015872155.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold47790	Nicotiana sylvestris	5568	6986	92%	0.0	88.95%	43786	NW_009534709.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3580	Nicotiana tomentosiformis	5565	5863	87%	0.0	89.97%	161351	NW_008916708.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold98834	Nicotiana sylvestris	5559	12870	100%	0.0	95.75%	50741	NW_009591424.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13662	Nicotiana tomentosiformis	5557	7741	91%	0.0	89.09%	53558	NW_008869180.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold70748	Nicotiana tomentosiformis	5550	6479	89%	0.0	91.70%	20671	NW_008955539.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13520	Nicotiana tabacum	5548	6399	87%	0.0	89.90%	110441	NW_015813061.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold63571	Nicotiana tabacum	5548	7066	83%	0.0	91.18%	201062	NW_015924977.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold15431	Nicotiana tabacum	5541	5541	82%	0.0	89.43%	54981	NW_015825658.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9395	Nicotiana tomentosiformis	5541	5,54E+03	82%	0.0	89.47%	108168	NW_008980818.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold29899	Nicotiana tomentosiformis	5531	8050	99%	0.0	90.03%	25575	NW_008910150.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5558	Nicotiana tomentosiformis	5520	9707	91%	0.0	91.15%	726744	NW_008938929.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8583	Nicotiana tabacum	5518	8054	100%	0.0	89.89%	160659	NW_015944044.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1315	Nicotiana sylvestris	5511	10321	100%	0.0	94.57%	72001	NW_009373803.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	5505	7924	92%	0.0	92.64%	271573	NW_017671610.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold26751	Nicotiana tabacum	5505	6758	94%	0.0	90.21%	165942	NW_015883831.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold54557	Nicotiana sylvestris	5505	5505	86%	0.0	88.43%	72065	NW_009542228.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold30103	Nicotiana tabacum	5500	6438	94%	0.0	90.56%	210528	NW_015888138.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold30412	Nicotiana tabacum	5500	7628	89%	0.0	93.15%	31389	NW_015888430.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold30069	Nicotiana tabacum	5494	15284	99%	0.0	88.79%	45921	NW_015888106.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold74241	Nicotiana tabacum	5494	5494	86%	0.0	88.39%	46557	NW_015934242.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold32257	Nicotiana tomentosiformis	5474	6381	93%	0.0	90.37%	17656	NW_008912772.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold24535	Nicotiana sylvestris	5469	6805	91%	0.0	88.64%	172148	NW_009500303.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold44741	Nicotiana tabacum	5467	11639	94%	0.0	97.08%	66850	NW_015907899.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3199	Nicotiana tomentosiformis	5465	8011	100%	0.0	89.61%	202801	NW_008912474.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold6107	Nicotiana sylvestris	5463	14194	92%	0.0	89.75%	36653	NW_009549465.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold9923	Nicotiana sylvestris	5459	7290	91%	0.0	90.75%	94979	NW_009591864.1

Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9418	Nicotiana tomentosiformis	5457	6533	91%	0.0	89.76%	40924	NW_008981574.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold60797	Nicotiana tabacum	5454	7285	91%	0.0	90.73%	69637	NW_015922499.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold377	Nicotiana sylvestris	5454	8766	91%	0.0	94.78%	31479	NW_009523497.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1471	Nicotiana tomentosiformis	5446	6446	91%	0.0	90.08%	26401	NW_00880824.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold59872	Nicotiana tomentosiformis	5445	6092	89%	0.0	90.17%	10814	NW_008943454.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1906	Nicotiana tomentosiformis	5437	7207	90%	0.0	90.58%	100193	NW_008898108.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold35539	Nicotiana tomentosiformis	5435	5641	88%	0.0	89.01%	56038	NW_008916418.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold41048	Nicotiana tomentosiformis	5433	6686	94%	0.0	89.87%	107544	NW_008922540.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold28037	Nicotiana sylvestris	5432	5432	83%	0.0	88.75%	19210	NW_009512762.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17152	Nicotiana tabacum	5424	6744	91%	0.0	88.46%	74361	NW_015836020.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold2128	Nicotiana tabacum	5421	7174	90%	0.0	90.51%	144595	NW_015857975.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold2919	Nicotiana tabacum	5409	7406	90%	0.0	91.09%	212745	NW_015887280.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold27417	Nicotiana tomentosiformis	5400	6742	91%	0.0	88.48%	64299	NW_008907394.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold91200	Nicotiana tabacum	5397	5397	83%	0.0	88.61%	65536	NW_015948478.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold114737	Nicotiana tabacum	5395	6312	91%	0.0	88.46%	18667	NW_015798493.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold22102	Nicotiana tomentosiformis	5391	15150	99%	0.0	88.39%	66189	NW_008901490.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	5387	6494	80%	0.0	94.52%	57170	NW_017675889.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1570	Nicotiana sylvestris	5387	6643	92%	0.0	88.25%	257076	NW_009402136.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16206	Nicotiana tomentosiformis	5384	6991	90%	0.0	89.74%	49266	NW_008894938.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold28304	Nicotiana tabacum	5382	6637	92%	0.0	88.22%	256825	NW_015886456.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold56230	Nicotiana tabacum	5376	8223	90%	0.0	95.12%	54476	NW_015918426.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1150	Nicotiana tomentosiformis	5374	7801	93%	0.0	90.58%	165743	NW_008845158.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold34776	Nicotiana tabacum	5371	5371	82%	0.0	88.79%	41998	NW_015894675.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold43688	Nicotiana tabacum	5369	6943	90%	0.0	89.70%	29373	NW_015906942.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold92338	Nicotiana tabacum	5360	7429	94%	0.0	89.54%	47373	NW_015949414.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 4, NIATT2	Nicotiana attenuata	5358	16029	91%	0.0	94.74%	12333206	NC_031992.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	5358	6545	92%	0.0	90.97%	96970	NW_017674432.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold95716	Nicotiana tabacum	5347	6150	91%	0.0	88.29%	156637	NW_015952090.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold32520	Nicotiana sylvestris	5337	6130	91%	0.0	88.28%	193941	NW_009517744.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	5326	7023	90%	0.0	89.75%	794788	NW_017670401.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold44798	Nicotiana sylvestris	5325	7988	89%	0.0	93.69%	97911	NW_009531184.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold44618	Nicotiana sylvestris	5321	6630	83%	0.0	90.67%	88679	NW_009531185.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	5317	5958	89%	0.0	88.21%	1653366	NW_017670255.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold52051	Nicotiana tabacum	5317	7525	99%	0.0	89.09%	96161	NW_015914613.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17004	Nicotiana tabacum	5299	7128	85%	0.0	89.18%	94556	NW_015835191.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold34455	Nicotiana sylvestris	5293	6661	86%	0.0	89.33%	47248	NW_009519893.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold55762	Nicotiana tabacum	5288	5288	83%	0.0	88.17%	16233	NW_015917998.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold5222	Nicotiana sylvestris	5282	5282	83%	0.0	88.15%	63977	NW_009542968.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold16464	Nicotiana tabacum	5276	6645	86%	0.0	89.26%	122971	NW_015831998.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold55560	Nicotiana tabacum	5273	5491	84%	0.0	88.15%	45686	NW_015917823.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold80979	Nicotiana sylvestris	5273	5273	83%	0.0	88.15%	31567	NW_009571585.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold29142	Nicotiana sylvestris	5267	7035	84%	0.0	95.30%	21086	NW_009513990.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold39729	Nicotiana tomentosiformis	5267	6137	89%	0.0	90.14%	64536	NW_008921073.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold14361	Nicotiana tabacum	5256	8237	90%	0.0	95.05%	48323	NW_015818785.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9005	Nicotiana tomentosiformis	5256	7476	99%	0.0	88.83%	347138	NW_008976986.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9507	Nicotiana tabacum	5251	6957	95%	0.0	91.64%	106418	NW_015951571.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1171	Nicotiana sylvestris	5245	6558	92%	0.0	87.66%	66843	NW_009357803.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold57812	Nicotiana tabacum	5234	6093	89%	0.0	89.99%	205811	NW_015919868.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold64459	Nicotiana tabacum	5234	5894	84%	0.0	90.69%	131598	NW_015925772.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10049	Nicotiana tabacum	5227	5227	83%	0.0	88.03%	75546	NW_015787621.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold130228	Nicotiana tabacum	5227	5769	84%	0.0	90.00%	7812	NW_015809656.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5281	Nicotiana tabacum	5227	6745	94%	0.0	91.59%	85240	NW_015915319.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold73913	Nicotiana tabacum	5225	8739	80%	0.0	95.73%	46711	NW_015933954.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13474	Nicotiana tabacum	5223	7061	97%	0.0	89.82%	102210	NW_015812762.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5623	Nicotiana tomentosiformis	5221	6703	94%	0.0	91.56%	25818	NW_008939408.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold70583	Nicotiana tabacum	5216	7816	91%	0.0	92.10%	36823	NW_015931092.1

Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1585	Nicotiana tomentosiformis	5193	6894	90%	0.0	91.38%	117667	NW_008893490.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10857	Nicotiana tabacum	5192	6,38E+03	90%	0.0	90.94%	172274	NW_015793863.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold48866	Nicotiana tabacum	5190	7,97E+03	99%	0.0	89.99%	66494	NW_015911712.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold11533	Nicotiana tomentosiformis	5190	7,81E+03	91%	0.0	91.98%	113970	NW_008845525.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold59680	Nicotiana tomentosiformis	5190	6,49E+03	82%	0.0	91.07%	56426	NW_008943241.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold6770	Nicotiana tomentosiformis	5188	7,29E+03	94%	0.0	90.05%	248095	NW_008952152.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold16999	Nicotiana tabacum	5186	7,20E+03	90%	0.0	91.47%	48105	NW_015835159.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold41084	Nicotiana tomentosiformis	5184	6,40E+03	90%	0.0	90.91%	61302	NW_008922580.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31284	Nicotiana tomentosiformis	5171	5,43E+03	85%	0.0	88.76%	143321	NW_008911691.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold35151	Nicotiana tabacum	5169	7,39E+03	99%	0.0	89.83%	49007	NW_015898773.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold51823	Nicotiana tabacum	5169	7543	94%	0.0	90.44%	57990	NW_015914400.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17820	Nicotiana tabacum	5168	7,27E+03	94%	0.0	89.98%	131673	NW_015839800.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9696	Nicotiana tomentosiformis	5168	7558	94%	0.0	90.44%	80470	NW_008984662.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold6593	Nicotiana tomentosiformis	5166	7177	90%	0.0	91.39%	64521	NW_008950185.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold19812	Nicotiana sylvestris	5155	5614	92%	0.0	87.34%	115706	NW_009447825.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16498	Nicotiana tomentosiformis	5155	83%	0.0	87.79%	196573	NW_008895261.1	
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold27847	Nicotiana tabacum	5147	6,26E+03	92%	0.0	88.27%	92593	NW_015886023.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold30764	Nicotiana tomentosiformis	5147	6223	92%	0.0	87.22%	65571	NW_008911113.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold73783	Nicotiana tabacum	5142	6,20E+03	92%	0.0	87.23%	64113	NW_015933843.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 7, NIATT2	Nicotiana attenuata	5136	49909	97%	0.0	87.65%	114765206	NC_031995.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9484	Nicotiana tomentosiformis	5134	7348	99%	0.0	89.65%	158425	NW_00892307.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold6090	Nicotiana sylvestris	5121	6458	91%	0.0	87.32%	273803	NW_009549276.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold29758	Nicotiana tabacum	5118	5,56E+03	92%	0.0	87.22%	104332	NW_015887820.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold29620	Nicotiana tomentosiformis	5114	8905	90%	0.0	91.07%	32394	NW_008909842.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8603	Nicotiana tabacum	5110	6,10E+03	90%	0.0	87.56%	410311	NW_015944214.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold10014	Nicotiana sylvestris	5107	10279	90%	0.0	94.74%	92590	NW_009338960.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold3997	Nicotiana sylvestris	5107	8,05E+03	90%	0.0	94.27%	66523	NW_009526019.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold12454	Nicotiana tomentosiformis	5107	6207	92%	0.0	88.14%	262243	NW_008855758.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold59851	Nicotiana tabacum	5105	6,36E+03	92%	0.0	87.21%	131180	NW_015921646.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold64399	Nicotiana sylvestris	5105	1,03E+04	83%	0.0	98.81%	40493	NW_009553163.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7770	Nicotiana tabacum	5096	6425	91%	0.0	87.25%	96594	NW_015937208.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold16847	Nicotiana tabacum	5066	11328	94%	0.0	87.33%	102024	NW_015834252.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	5062	5816	88%	0.0	90.19%	317457	NW_017671304.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold3371	Nicotiana tabacum	5057	13556	99%	0.0	89.97%	227154	NW_015891551.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9542	Nicotiana tomentosiformis	5055	5302	89%	0.0	87.30%	30946	NW_008982952.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	5051	5996	83%	0.0	90.20%	240938	NW_017671884.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold43524	Nicotiana tomentosiformis	5051	11821	94%	0.0	87.31%	23016	NW_008925291.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold6469	Nicotiana sylvestris	5049	7809	81%	0.0	95.11%	303965	NW_009553486.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold56527	Nicotiana sylvestris	5046	12077	95%	0.0	98.20%	27662	NW_009544147.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold6392	Nicotiana tomentosiformis	5046	5501	80%	0.0	89.92%	137286	NW_008947952.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold5662	Nicotiana sylvestris	5035	6640	92%	0.0	88.71%	256742	NW_009544520.1
Tnt1	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	5029	35344	89%	0.0	87.10%	77197300	NC_028640.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold39253	Nicotiana sylvestris	5027	8278	94%	0.0	93.14%	50535	NW_009525244.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold24713	Nicotiana sylvestris	5024	11035	84%	0.0	95.33%	39938	NW_009502281.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold53146	Nicotiana sylvestris	5024	11137	85%	0.0	98.69%	73043	NW_009540661.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold63764	Nicotiana tomentosiformis	5024	7682	91%	0.0	89.96%	5658	NW_008947779.1
Tnt1	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum pennellii	5022	43969	91%	0.0	86.92%	77991103	NC_028641.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold25007	Nicotiana tabacum	5022	7360	82%	0.0	94.95%	151099	NW_015875411.1
Tnt1	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	5018	59566	89%	0.0	87.06%	60730942	NC_028642.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold29524	Nicotiana tabacum	5016	7456	89%	0.0	91.48%	10402	NW_015887600.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1195	Nicotiana tomentosiformis	5014	7303	83%	0.0	91.85%	197884	NW_008850157.1
Tnt1	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	Solanum pennellii	5012	37139	89%	0.0	87.03%	70546659	NC_028644.1
Tnt1	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	Solanum pennellii	5012	55278	91%	0.0	87.03%	84057508	NC_028645.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold15046	Nicotiana tomentosiformis	5012	5713	88%	0.0	87.17%	116911	NW_008884558.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold89324	Nicotiana tabacum	5011	5302	82%	0.0	88.60%	26865	NW_015946922.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31268	Nicotiana tomentosiformis	5011	7450	89%	0.0	91.45%	43749	NW_008911673.1

Tnt1	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	Solanum pennellii	5009	32063	89%	0.0	87.07%	83305730	NC_028648.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold19423	Nicotiana tabacum	5009	6530	85%	0.0	89.49%	425230	NW_015848601.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5744	Nicotiana tabacum	5009	8093	99%	0.0	91.00%	216975	NW_015919530.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold70315	Nicotiana tabacum	5007	7896	91%	0.0	92.92%	54314	NW_015930853.1
Tnt1	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	5000	23202	89%	0.0	86.83%	59803892	NC_028638.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2792	Nicotiana tomentosiformis	5000	7022	86%	0.0	91.73%	210823	NW_008907952.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold642	Nicotiana tabacum	4988	6888	96%	0.0	87.27%	299001	NW_015925547.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold4323	Nicotiana sylvestris	4988	7843	88%	0.0	93.99%	63368	NW_009529643.1
Tnt1	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	4985	31383	91%	0.0	86.97%	75414019	NC_028639.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23142	Nicotiana tabacum	4985	6969	86%	0.0	91.64%	131057	NW_015866829.1
Tnt1	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum pennellii	4983	55157	91%	0.0	86.96%	82529941	NC_028646.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold22870	Nicotiana tomentosiformis	4977	5274	82%	0.0	88.45%	27703	NW_008902342.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold142	Nicotiana tabacum	4963	6465	89%	0.0	89.79%	66513	NW_015817726.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold14111	Nicotiana tomentosiformis	4959	8071	99%	0.0	90.76%	73162	NW_008874170.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24903	Nicotiana tabacum	4948	5479	82%	0.0	89.41%	72338	NW_015874945.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold11441	Nicotiana tomentosiformis	4933	6829	96%	0.0	87.04%	118416	NW_008844503.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4931	6032	91%	0.0	89.14%	395812	NW_017670962.1
Tnt1	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	4927	42273	91%	0.0	86.75%	109333515	NC_028637.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold25780	Nicotiana tabacum	4927	5169	86%	0.0	87.21%	39143	NW_015878929.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16076	Nicotiana tomentosiformis	4926	5451	82%	0.0	89.31%	28504	NW_008894793.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold68777	Nicotiana sylvestris	4924	11789	100%	0.0	95.78%	106987	NW_009558027.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13841	Nicotiana tabacum	4922	5992	82%	0.0	91.35%	61294	NW_015815290.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4918	5952	90%	0.0	86.66%	107418	NW_017674142.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16671	Nicotiana tomentosiformis	4918	4918	83%	0.0	86.81%	116571	NW_008895454.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold21251	Nicotiana tabacum	4900	4900	83%	0.0	86.71%	252328	NW_015857843.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5783	Nicotiana tabacum	4892	8170	94%	0.0	91.09%	199764	NW_015919844.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold78619	Nicotiana tabacum	4889	6278	82%	0.0	94.24%	26634	NW_015937995.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold21977	Nicotiana tabacum	4879	7120	82%	0.0	92.00%	26654	NW_015861371.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold29741	Nicotiana tomentosiformis	4868	7081	82%	0.0	91.94%	27061	NW_008909976.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13396	Nicotiana tabacum	4863	6382	92%	0.0	88.01%	247980	NW_015812226.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold31097	Nicotiana sylvestris	4854	8684	92%	0.0	94.30%	56767	NW_009516162.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold6426	Nicotiana tomentosiformis	4852	7247	91%	0.0	90.89%	113929	NW_008948330.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold17335	Nicotiana sylvestris	4826	10752	100%	0.0	95.94%	111540	NW_009420303.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold60020	Nicotiana tabacum	4815	6260	82%	0.0	91.05%	117055	NW_015921803.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13741	Nicotiana tomentosiformis	4809	6276	82%	0.0	91.02%	308773	NW_008870058.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold44457	Nicotiana tomentosiformis	4807	6004	89%	0.0	90.21%	63443	NW_008926327.1
Tnt1	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00684	Solanum tuberosum	4807	5119	89%	0.0	86.35%	199035	NW_006239598.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold33905	Nicotiana tomentosiformis	4802	7190	89%	0.0	91.07%	99512	NW_008914603.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold12251	Nicotiana tabacum	4796	7179	89%	0.0	91.06%	117754	NW_015804145.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4787	4787	81%	0.0	86.90%	79688	NW_017674964.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold49166	Nicotiana tabacum	4782	10901	100%	0.0	95.97%	99083	NW_015911986.1
Tnt1	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	Solanum pennellii	4780	30698	90%	0.0	86.13%	66223686	NC_028647.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold75918	Nicotiana tabacum	4780	7419	88%	0.0	91.42%	211019	NW_015935681.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold80642	Nicotiana tabacum	4776	5967	92%	0.0	85.90%	42565	NW_015939713.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold22530	Nicotiana sylvestris	4765	5145	92%	0.0	85.85%	14904	NW_009478026.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4761	5530	90%	0.0	86.21%	130124	NW_017673546.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold2956	Nicotiana sylvestris	4758	8068	91%	0.0	94.33%	20573	NW_009514453.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 5, NIATT2	Nicotiana attenuata	4752	26994	94%	0.0	91.85%	57012777	NC_031993.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold107752	Nicotiana tabacum	4752	9054	83%	0.0	95.77%	52021	NW_015793217.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold373	Nicotiana tabacum	4747	7208	93%	0.0	91.32%	326739	NW_015901010.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold16261	Nicotiana tabacum	4737	7057	89%	0.0	91.22%	63271	NW_015830791.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold545	Nicotiana sylvestris	4735	9281	91%	0.0	93.54%	250934	NW_009542164.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold6807	Nicotiana sylvestris	4732	4732	81%	0.0	86.32%	46912	NW_009557242.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold521	Nicotiana tomentosiformis	4726	7193	95%	0.0	91.22%	392850	NW_008934819.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold638	Nicotiana tabacum	4724	12192	98%	0.0	90.67%	95724	NW_015925192.1
Tnt1	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00673	Solanum tuberosum	4715	4715	83%	0.0	85.96%	182086	NW_006239623.1

Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold49643	Nicotiana tabacum	4713	7179	91%	0.0	90.47%	156996	NW_015912413.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold25466	Nicotiana tabacum	4704	7818	80%	0.0	95.94%	7262	NW_015877450.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold58725	Nicotiana tabacum	4691	7949	87%	0.0	93.59%	99986	NW_015920660.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13841	Nicotiana tomentosiformis	4691	9613	89%	0.0	89.58%	206941	NW_008871169.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold3277	Nicotiana tabacum	4686	7060	95%	0.0	89.26%	135543	NW_015890651.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold15302	Nicotiana tomentosiformis	4684	6851	86%	0.0	90.27%	86097	NW_008887403.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3495	Nicotiana tomentosiformis	4680	7098	92%	0.0	90.36%	253376	NW_008915763.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold46210	Nicotiana tabacum	4678	5992	92%	0.0	87.07%	96369	NW_015909277.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5462	Nicotiana tomentosiformis	4678	5992	92%	0.0	87.07%	87762	NW_008937619.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold125551	Nicotiana tabacum	4676	4879	87%	0.0	86.12%	53275	NW_015806344.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold2477	Nicotiana tabacum	4674	7080	80%	0.0	92.58%	158578	NW_015874368.1
Tnt1	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	4669	23621	88%	0.0	85.70%	65987440	NC_015445.3
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8404	Nicotiana tabacum	4667	6873	86%	0.0	90.19%	77498	NW_015942554.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold201213	Nicotiana tabacum	4663	4663	81%	0.0	86.07%	33078	NW_015852220.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold25606	Nicotiana sylvestris	4663	9480	84%	0.0	97.14%	19299	NW_009510061.1
Tnt1	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	4663	46356	91%	0.0	85.64%	65633393	NC_015447.3
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4506	Nicotiana tomentosiformis	4662	7789	95%	0.0	91.29%	139730	NW_008926997.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9369	Nicotiana tomentosiformis	4656	7389	92%	0.0	92.56%	55160	NW_008981029.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1545	Nicotiana sylvestris	4654	4654	82%	0.0	86.04%	60314	NW_009399358.1
Tnt1	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00302	Solanum tuberosum	4654	4956	89%	0.0	85.67%	758973	NW_006239238.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold109	Nicotiana tabacum	4641	7530	98%	0.0	89.77%	169686	NW_015794174.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4630	7672	80%	0.0	94.64%	757831	NW_017670414.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4630	6718	89%	0.0	89.73%	237845	NW_017671904.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold41462	Nicotiana sylvestris	4630	8149	88%	0.0	96.03%	16730	NW_009527679.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold12415	Nicotiana sylvestris	4628	6,76E+03	86%	0.0	93.50%	151871	NW_009365637.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold3281	Nicotiana sylvestris	4628	7840	87%	0.0	93.68%	114242	NW_009518065.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold61262	Nicotiana tabacum	4619	7,88E+03	95%	0.0	91.06%	117776	NW_015922915.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold7841	Nicotiana sylvestris	4588	8188	90%	0.0	93.43%	94394	NW_009568731.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold28530	Nicotiana tabacum	4582	7,65E+03	84%	0.0	92.07%	108497	NW_015886661.1
Tnt1	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	4580	2,01E+04	88%	0.0	85.31%	55977580	NC_015439.3
Tnt1	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	4580	1,60E+04	89%	0.0	85.36%	66557038	NC_015441.3
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold28736	Nicotiana tomentosiformis	4578	1,06E+04	98%	0.0	90.39%	28974	NW_008908859.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold100303	Nicotiana tabacum	4577	7,46E+03	97%	0.0	91.02%	111080	NW_015787473.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold81684	Nicotiana tabacum	4577	6670	80%	0.0	91.35%	44923	NW_015940596.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold30255	Nicotiana sylvestris	4566	9000	89%	0.0	97.33%	29426	NW_009515227.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold17473	Nicotiana tomentosiformis	4560	7,61E+03	97%	0.0	90.97%	178286	NW_008896345.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold30024	Nicotiana tabacum	4558	8,41E+03	90%	0.0	93.68%	76262	NW_015888064.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold15201	Nicotiana tomentosiformis	4558	7,44E+03	98%	0.0	89.39%	47518	NW_008886281.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold90456	Nicotiana tabacum	4554	1,49E+04	100%	0.0	94.81%	33497	NW_015947849.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold21362	Nicotiana tomentosiformis	4554	4554	84%	0.0	85.19%	130980	NW_008900667.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold316	Nicotiana tomentosiformis	4551	9,09E+03	91%	0.0	89.53%	23372	NW_008912041.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold8300	Nicotiana tomentosiformis	4551	8,02E+03	99%	0.0	90.76%	90653	NW_008969153.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold83	Nicotiana tomentosiformis	4549	8,68E+03	92%	0.0	92.00%	155576	NW_008969151.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold14673	Nicotiana tabacum	4540	9,01E+03	91%	0.0	89.47%	18380	NW_015820835.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4536	6166	90%	0.0	88.76%	658709	NW_017670476.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold18520	Nicotiana tabacum	4536	6232	95%	0.0	90.24%	328628	NW_015843687.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold195	Nicotiana tomentosiformis	4530	6,22E+03	95%	0.0	90.21%	67517	NW_008898596.1
Tnt1	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	4529	13247	89%	0.0	85.56%	68126176	NC_015449.3
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10176	Nicotiana tabacum	4527	4,53E+03	83%	0.0	85.30%	303100	NW_015788617.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold35904	Nicotiana sylvestris	4514	8182	90%	0.0	94.40%	31364	NW_009521503.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2994	Nicotiana tomentosiformis	4514	8,18E+03	98%	0.0	92.62%	171488	NW_008910196.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7359	Nicotiana tabacum	4508	7,86E+03	84%	0.0	95.50%	78256	NW_015933674.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold7568	Nicotiana tomentosiformis	4499	4499	83%	0.0	85.21%	95941	NW_008961018.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4494	7,07E+03	93%	0.0	90.09%	583767	NW_017670553.1
Tnt1	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	4473	2,51E+04	89%	0.0	84.96%	49794276	NC_015443.3
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold151436	Nicotiana tabacum	4470	4,47E+03	84%	0.0	84.74%	48833	NW_015823781.1

Tnt1	refseq_genomes	8	Capisicum annum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	Capisicum annum	4460	3,47E+04	88%	0.0	84.97%	205736368	NC_029986.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold383	Nicotiana tomentosiformis	4460	1,27E+04	94%	0.0	90.50%	295616	NW_008919485.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold38382	Nicotiana tabacum	4458	6,51E+03	90%	0.0	90.15%	165917	NW_015902036.1
Tnt1	refseq_genomes	8	Capisicum annum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	Capisicum annum	4457	2,99E+04	92%	0.0	85.01%	217274494	NC_029981.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4935	Nicotiana tabacum	4455	6854	90%	0.0	90.47%	44734	NW_015912146.1
Tnt1	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	Solanum pennellii	4453	1,90E+04	89%	0.0	84.77%	79292169	NC_028643.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold92017	Nicotiana tabacum	4451	8,29E+03	81%	0.0	98.88%	29324	NW_015949138.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold12875	Nicotiana tomentosiformis	4444	7422	99%	0.0	91.32%	132480	NW_008860435.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold32132	Nicotiana tabacum	4440	8008	98%	0.0	92.19%	83779	NW_015890061.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold41492	Nicotiana sylvestris	4440	6468	90%	0.0	90.06%	88890	NW_009527712.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold2343	Nicotiana tabacum	4438	7455	99%	0.0	91.29%	68530	NW_015868198.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold26334	Nicotiana tabacum	4434	9389	96%	0.0	91.86%	24328	NW_015881563.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold73667	Nicotiana tabacum	4425	8,27E+03	91%	0.0	93.96%	44534	NW_015933742.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold52447	Nicotiana tabacum	4421	5314	87%	0.0	87.12%	8946	NW_015914976.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4420	5345	80%	0.0	89.85%	556156	NW_017670591.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold89121	Nicotiana sylvestris	4418	5340	82%	0.0	87.32%	33088	NW_009580633.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4416	8149	90%	0.0	92.60%	202305	NW_017672291.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold31010	Nicotiana tabacum	4414	10675	100%	0.0	95.00%	60758	NW_015889002.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold120371	Nicotiana tabacum	4412	6264	85%	0.0	87.28%	34622	NW_015802584.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold33484	Nicotiana sylvestris	4409	5290	87%	0.0	87.07%	28573	NW_009518814.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4407	5886	83%	0.0	90.73%	232909	NW_017671950.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold14374	Nicotiana sylvestris	4403	9441	95%	0.0	93.09%	59061	NW_009387403.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold7145	Nicotiana tomentosiformis	4403	7,08E+03	92%	0.0	89.61%	51617	NW_008956319.1
Tnt1	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	4396	24590	89%	0.0	84.65%	72290146	NC_015440.3
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold44953	Nicotiana tabacum	4394	7478	92%	0.0	90.99%	13596	NW_015908089.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4392	8,04E+03	93%	0.0	92.90%	270725	NW_017671620.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4009	Nicotiana tabacum	4388	7541	91%	0.0	90.93%	103833	NW_015903615.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold97809	Nicotiana tabacum	4386	7057	92%	0.0	89.53%	61514	NW_015953746.1
Tnt1	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	4386	10287	84%	0.0	84.42%	56597135	NC_015448.3
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold8953	Nicotiana tomentosiformis	4370	7,73E+03	93%	0.0	91.09%	149722	NW_008976407.1
Tnt1	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	4362	29609	89%	0.0	84.35%	98455869	NC_015438.3
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold208371	Nicotiana tabacum	4361	7404	87%	0.0	92.30%	21834	NW_015855862.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold19913	Nicotiana tomentosiformis	4353	8100	100%	0.0	90.05%	145559	NW_008899056.1
Tnt1	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 S16 R44 unplaced genomic scaffold, SolTub_3.0 scf00337	Solanum tuberosum	4353	8,49E+03	90%	0.0	84.86%	664521	NW_006239266.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4278	Nicotiana tabacum	4348	7,73E+03	93%	0.0	90.97%	82534	NW_015906105.1
Tnt1	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	4348	40056	90%	0.0	84.41%	72906345	NC_015446.3
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold109560	Nicotiana sylvestris	4340	6873	83%	0.0	92.18%	19320	NW_009349426.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold36882	Nicotiana tomentosiformis	4340	5,40E+03	91%	0.0	87.75%	47244	NW_008917910.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold25586	Nicotiana tabacum	4337	8,07E+03	100%	0.0	89.96%	77139	NW_015878030.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold40766	Nicotiana tabacum	4331	9,20E+03	98%	0.0	91.04%	143100	NW_015904229.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold965	Nicotiana tabacum	4322	10560	100%	0.0	94.61%	163997	NW_015952705.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4313	8,75E+03	100%	0.0	92.81%	182969	NW_017672576.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8572	Nicotiana tabacum	4311	10134	99%	0.0	94.93%	32918	NW_015943952.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold26772	Nicotiana tomentosiformis	4311	7,98E+03	96%	0.0	91.39%	94554	NW_008906677.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold32717	Nicotiana tomentosiformis	4307	7432	87%	0.0	91.75%	103041	NW_008913283.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4305	7302	95%	0.0	90.99%	1050219	NW_017670312.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5409	Nicotiana tabacum	4305	7906	99%	0.0	90.49%	141880	NW_015916489.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold29340	Nicotiana tomentosiformis	4303	9199	98%	0.0	90.89%	56359	NW_008909531.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4852	Nicotiana tomentosiformis	4300	6405	89%	0.0	91.12%	19402	NW_008930841.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold6724	Nicotiana tabacum	4296	6819	92%	0.0	89.54%	155646	NW_015928195.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1241	Nicotiana tomentosiformis	4285	6883	91%	0.0	91.53%	189737	NW_008855269.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold33225	Nicotiana tabacum	4268	8652	99%	0.0	92.22%	78696	NW_015891089.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold14352	Nicotiana sylvestris	4265	6825	92%	0.0	89.36%	127016	NW_009387159.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold18953	Nicotiana tomentosiformis	4261	5219	82%	0.0	88.51%	56156	NW_008897989.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold11317	Nicotiana tomentosiformis	4244	7367	99%	0.0	89.02%	177377	NW_008843125.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold20699	Nicotiana tomentosiformis	4244	7,06E+03	94%	0.0	89.82%	42174	NW_008899929.1

Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold21750	Nicotiana tabacum	4242	8058	91%	0.0	94.73%	53630	NW_015860265.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold92074	Nicotiana tabacum	4222	7105	92%	0.0	89.22%	91110	NW_015949192.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold5319	Nicotiana sylvestris	4217	7100	92%	0.0	89.19%	97780	NW_009540709.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold34131	Nicotiana tomentosiformis	4209	7222	90%	0.0	90.30%	57011	NW_008914855.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold15284	Nicotiana tabacum	4207	7019	89%	0.0	89.61%	95461	NW_015824723.1
Tnt1	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	4189	20208	89%	0.0	84.12%	219521584	NC_029982.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold91491	Nicotiana sylvestris	4185	7911	91%	0.0	91.54%	8966	NW_009583266.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold51271	Nicotiana tomentosiformis	4185	5567	84%	0.0	89.49%	53471	NW_008933899.1
Tnt1	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	4178	37579	91%	0.0	83.38%	301019445	NC_029977.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10218	Nicotiana tabacum	4169	7747	97%	0.0	90.37%	90732	NW_015788933.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold27125	Nicotiana tabacum	4167	13350	91%	0.0	91.29%	73230	NW_015885333.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold33676	Nicotiana tabacum	4163	7565	97%	0.0	91.87%	150939	NW_015891517.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold40603	Nicotiana tabacum	4163	8426	100%	0.0	91.80%	41763	NW_015904078.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10608	Nicotiana tabacum	4157	8298	94%	0.0	93.27%	70778	NW_015791951.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2886	Nicotiana tomentosiformis	4135	15259	92%	0.0	91.15%	101759	NW_008908996.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4130	6024	84%	0.0	88.87%	241957	NW_017671873.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5395	Nicotiana tomentosiformis	4130	7692	98%	0.0	88.47%	154383	NW_008936874.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold244432	Nicotiana tabacum	4128	6683	92%	0.0	88.02%	30009	NW_015872910.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold7307	Nicotiana sylvestris	4117	6677	92%	0.0	87.97%	50604	NW_009562798.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1212	Nicotiana tomentosiformis	4115	15071	99%	0.0	92.87%	162405	NW_008852047.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9490	Nicotiana tomentosiformis	4113	7,40E+03	96%	0.0	91.55%	89622	NW_008928374.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold78822	Nicotiana tabacum	4111	7850	90%	0.0	94.07%	47477	NW_015938172.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold28590	Nicotiana tomentosiformis	4108	7592	97%	0.0	91.07%	37010	NW_008908697.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10329	Nicotiana tabacum	4104	13494	99%	0.0	92.80%	197271	NW_015789795.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold55772	Nicotiana tabacum	4102	7,76E+03	85%	0.0	93.97%	131411	NW_015918009.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold28812	Nicotiana sylvestris	4095	7695	85%	0.0	93.93%	135008	NW_009513623.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold15762	Nicotiana tomentosiformis	4069	8056	96%	0.0	90.83%	189671	NW_008892513.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold31027	Nicotiana tabacum	4067	5939	89%	0.0	89.44%	99240	NW_015889017.1
Tnt1	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	4067	14258	90%	0.0	83.22%	66723567	NC_015442.3
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4063	8986	91%	0.0	92.90%	269263	NW_017671635.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold84028	Nicotiana tomentosiformis	4056	6501	90%	0.0	88.71%	40603	NW_008970295.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold67005	Nicotiana tabacum	4052	6970	96%	0.0	90.73%	83939	NW_015927993.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold46615	Nicotiana tomentosiformis	4052	7458	97%	0.0	92.00%	68794	NW_008928725.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold517	Nicotiana tomentosiformis	4048	4713	82%	0.0	86.61%	290780	NW_008934374.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3253	Nicotiana tomentosiformis	4047	7936	99%	0.0	89.23%	73937	NW_008913075.1
Tnt1	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00009	Solanum tuberosum	4043	5584	91%	0.0	85.91%	3955149	NW_006238935.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13115	Nicotiana tomentosiformis	4041	9069	99%	0.0	90.41%	18953	NW_008863103.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold32727	Nicotiana tabacum	4024	4951	81%	0.0	88.37%	121303	NW_015890612.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold44444	Nicotiana tomentosiformis	4023	5848	80%	0.0	91.26%	15947	NW_008926308.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4019	7,47E+03	91%	0.0	91.93%	1499139	NW_017670270.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7459	Nicotiana tabacum	4010	7392	90%	0.0	89.65%	93177	NW_015934548.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold8648	Nicotiana sylvestris	4010	6078	88%	0.0	89.82%	209164	NW_009577697.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold10247	Nicotiana sylvestris	4002	9525	92%	0.0	96.10%	75202	NW_009341548.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	3999	6175	93%	0.0	91.91%	156741	NW_017672987.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold102983	Nicotiana tabacum	3989	1,02E+04	94%	0.0	94.30%	27688	NW_015789560.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 11, NIATT2	Nicotiana attenuata	3986	26012	92%	0.0	88.57%	61252062	NC_031999.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold3753	Nicotiana tabacum	3982	8989	100%	0.0	91.08%	92938	NW_015901228.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13764	Nicotiana tabacum	3973	14871	91%	0.0	91.25%	194850	NW_015814779.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold63722	Nicotiana tabacum	3962	7517	91%	0.0	90.88%	51917	NW_015925120.1
Tnt1	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00151	Solanum tuberosum	3949	4795	83%	0.0	82.97%	1427473	NW_006239077.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31534	Nicotiana tomentosiformis	3945	6835	89%	0.0	91.93%	24477	NW_008911969.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23487	Nicotiana tabacum	3940	6829	89%	0.0	91.89%	19909	NW_015868475.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23855	Nicotiana tabacum	3938	8094	97%	0.0	92.44%	108570	NW_015870190.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold22331	Nicotiana tabacum	3928	7647	90%	0.0	92.43%	58916	NW_015863021.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold23456	Nicotiana tomentosiformis	3928	8316	99%	0.0	89.85%	140848	NW_008902993.1
Tnt1	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3927	29765	90%	0.0	83.16%	215701946	NC_029980.1

Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold286	Nicotiana tabacum	3919	15893	90%	0.0	96.66%	51891	NW_015886725.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold35464	Nicotiana tomentosiformis	3912	7954	97%	0.0	89.73%	86820	NW_008916335.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3908	6194	88%	0.0	89.49%	149926	NW_017673130.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold29427	Nicotiana tabacum	3908	7512	84%	0.0	95.03%	130461	NW_015887504.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23709	Nicotiana tabacum	3904	7963	97%	0.0	89.69%	152180	NW_015869499.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold39359	Nicotiana sylvestris	3895	5137	84%	0.0	87.55%	48204	NW_009525341.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold26157	Nicotiana sylvestris	3892	8615	93%	0.0	94.76%	92458	NW_009510673.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold29316	Nicotiana tomentosiformis	3882	5866	87%	0.0	92.57%	37713	NW_008909504.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold77491	Nicotiana tabacum	3877	8627	90%	0.0	94.82%	22571	NW_015937021.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold20244	Nicotiana tomentosiformis	3873	8308	99%	0.0	92.22%	37044	NW_008899425.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8103	Nicotiana tabacum	3869	6645	93%	0.0	88.03%	277385	NW_015940038.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8398	Nicotiana tabacum	3869	8311	96%	0.0	92.36%	213188	NW_015942503.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold10951	Nicotiana tomentosiformis	3853	6547	93%	0.0	87.94%	203705	NW_008839060.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold64164	Nicotiana tomentosiformis	3818	7866	88%	0.0	91.35%	15545	NW_008948224.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4317	Nicotiana tabacum	3812	7436	91%	0.0	91.96%	58057	NW_015906465.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2641	Nicotiana tomentosiformis	3812	7458	91%	0.0	91.80%	54866	NW_008906275.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3807	7375	92%	0.0	91.46%	205250	NW_017672255.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold21950	Nicotiana tomentosiformis	3803	6723	85%	0.0	88.86%	121472	NW_008901320.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3797	10880	92%	0.0	89.74%	292058	NW_017671470.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold8208	Nicotiana sylvestris	3797	8612	91%	0.0	94.05%	36297	NW_009572809.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold74278	Nicotiana tabacum	3790	7972	89%	0.0	91.68%	40668	NW_015934269.1
Tnt1	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3775	28221	90%	0.0	85.21%	261510930	NC_029979.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3766	6528	88%	0.0	93.50%	180717	NW_017672605.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold63606	Nicotiana sylvestris	3762	5856	88%	0.0	89.21%	28214	NW_009552283.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold23420	Nicotiana tomentosiformis	3757	5,30E+03	83%	0.0	87.79%	75929	NW_008902954.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold14547	Nicotiana tabacum	3753	7344	89%	0.0	90.18%	206154	NW_01581994.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold30248	Nicotiana tabacum	3749	7897	99%	0.0	90.71%	24893	NW_015888281.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold100452	Nicotiana tabacum	3746	7397	92%	0.0	91.12%	47764	NW_015787590.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3744	8573	90%	0.0	92.49%	451606	NW_017670798.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8137	Nicotiana tabacum	3729	7432	89%	0.0	91.07%	163663	NW_015940333.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold10423	Nicotiana sylvestris	3714	10916	88%	0.0	91.45%	61192	NW_009343504.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold27644	Nicotiana tomentosiformis	3714	7334	92%	0.0	91.04%	21503	NW_008907646.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold39726	Nicotiana tabacum	3712	6433	88%	0.0	92.02%	58801	NW_015903284.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold123510	Nicotiana tabacum	3711	7,73E+03	87%	0.0	94.12%	20785	NW_015804850.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold6666	Nicotiana tomentosiformis	3709	7298	88%	0.0	91.54%	153770	NW_008950996.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold59226	Nicotiana tabacum	3698	7209	88%	0.0	91.44%	80388	NW_015921097.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3696	7483	92%	0.0	87.60%	722875	NW_017670432.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold14552	Nicotiana tomentosiformis	3696	6416	88%	0.0	91.91%	90377	NW_008879069.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold20166	Nicotiana tomentosiformis	3696	6416	89%	0.0	91.83%	24162	NW_008899338.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold60565	Nicotiana tabacum	3692	8556	88%	0.0	91.31%	8955	NW_015922290.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold180796	Nicotiana tabacum	3687	7299	93%	0.0	90.26%	25919	NW_015841257.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold9006	Nicotiana sylvestris	3685	7314	93%	0.0	90.26%	55494	NW_009581676.1
Tnt1	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3681	17164	90%	0.0	83.35%	222112641	NC_029983.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1027	Nicotiana tabacum	3672	5250	83%	0.0	87.39%	230627	NW_015789328.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3741	Nicotiana tomentosiformis	3661	5869	86%	0.0	90.09%	171772	NW_008918497.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3655	5789	86%	0.0	86.24%	471935	NW_017670738.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold17061	Nicotiana sylvestris	3653	8198	92%	0.0	93.65%	45223	NW_009417259.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold19110	Nicotiana tomentosiformis	3650	7529	88%	0.0	92.19%	154035	NW_008898165.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold10986	Nicotiana tomentosiformis	3646	16422	91%	0.0	90.96%	137028	NW_008839448.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold92772	Nicotiana tabacum	3640	10643	95%	0.0	95.68%	131555	NW_015949767.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold4727	Nicotiana sylvestris	3635	7133	92%	0.0	90.36%	88315	NW_009534131.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold24198	Nicotiana tomentosiformis	3633	7584	95%	0.0	90.32%	36790	NW_008903817.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold41161	Nicotiana tabacum	3631	6551	85%	0.0	90.30%	105525	NW_015904611.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold46265	Nicotiana tabacum	3631	7575	95%	0.0	90.32%	20876	NW_01590324.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold35759	Nicotiana tomentosiformis	3627	6322	95%	0.0	89.20%	6995	NW_008916662.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold111024	Nicotiana tabacum	3624	5709	82%	0.0	89.94%	59881	NW_015795712.1

Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold91771	Nicotiana tabacum	3624	7122	92%	0.0	90.30%	71387	NW_015948933.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9857	Nicotiana tabacum	3618	5819	83%	0.0	91.20%	32183	NW_015954336.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5077	Nicotiana tabacum	3616	7652	98%	0.0	89.13%	93465	NW_015913439.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold10378	Nicotiana tomentosiformis	3616	6045	90%	0.0	89.66%	63444	NW_008832693.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5170	Nicotiana tomentosiformis	3616	5389	85%	0.0	88.19%	107497	NW_008934375.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3611	5647	83%	0.0	90.13%	320405	NW_017671291.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3252	Nicotiana tomentosiformis	3609	6880	83%	0.0	92.03%	85200	NW_008913064.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold43177	Nicotiana tomentosiformis	3609	6943	95%	0.0	89.42%	91191	NW_008924905.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17050	Nicotiana tabacum	3602	5368	85%	0.0	88.10%	123534	NW_015835450.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold20780	Nicotiana tabacum	3598	6926	95%	0.0	89.35%	182166	NW_015855560.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold6933	Nicotiana tabacum	3592	6340	90%	0.0	88.59%	59409	NW_015930013.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold58993	Nicotiana tabacum	3581	5311	83%	0.0	89.42%	88922	NW_015920896.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold55979	Nicotiana sylvestris	3578	6608	87%	0.0	93.39%	58269	NW_009543807.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold24531	Nicotiana tomentosiformis	3578	6720	87%	0.0	91.14%	75806	NW_008904188.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold98105	Nicotiana tabacum	3570	7419	91%	0.0	90.89%	134467	NW_015953981.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold11260	Nicotiana sylvestris	3568	7241	95%	0.0	90.78%	187450	NW_00935204.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold35439	Nicotiana tomentosiformis	3568	7490	91%	0.0	90.89%	54089	NW_008916307.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold15893	Nicotiana tabacum	3565	7099	83%	0.0	91.75%	52900	NW_015828522.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold6443	Nicotiana tabacum	3555	6919	91%	0.0	90.17%	100864	NW_015925745.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold12901	Nicotiana tabacum	3552	7197	95%	0.0	90.66%	182667	NW_015808778.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold7251	Nicotiana sylvestris	3550	6913	91%	0.0	90.13%	101192	NW_009562176.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold24933	Nicotiana tomentosiformis	3544	6299	86%	0.0	92.59%	11497	NW_008904634.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23327	Nicotiana tabacum	3542	6885	91%	0.0	90.02%	146403	NW_015867717.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold23186	Nicotiana sylvestris	3542	6885	91%	0.0	90.02%	172860	NW_009485314.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold32673	Nicotiana tabacum	3539	5948	91%	0.0	90.35%	69498	NW_015890563.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold31202	Nicotiana tabacum	3537	6926	91%	0.0	91.70%	26297	NW_015889184.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold25103	Nicotiana tabacum	3535	8359	99%	0.0	91.99%	13510	NW_015875847.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold20522	Nicotiana tomentosiformis	3526	6,61E+03	91%	0.0	91.62%	68385	NW_008899734.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 12, NIATTR2	Nicotiana attenuata	3515	8713	93%	0.0	86.74%	53491057	NC_032000.1
Tnt1	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	3515	15169	89%	0.0	83.81%	68175699	NC_015444.3
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold11162	Nicotiana tomentosiformis	3511	6846	95%	0.0	91.51%	162051	NW_008841403.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold45022	Nicotiana tabacum	3500	5732	86%	0.0	89.04%	238682	NW_015908158.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold879	Nicotiana tabacum	3498	6881	90%	0.0	91.43%	68967	NW_015945741.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold27611	Nicotiana tabacum	3487	9714	91%	0.0	95.39%	62954	NW_015885787.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold56939	Nicotiana tabacum	3472	7786	84%	0.0	94.46%	65224	NW_015919069.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10459	Nicotiana tabacum	3463	7943	99%	0.0	91.21%	15429	NW_015790785.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold19678	Nicotiana sylvestris	3463	6823	90%	0.0	89.78%	83406	NW_009446335.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3461	7558	93%	0.0	90.41%	177713	NW_017672665.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3907	Nicotiana tomentosiformis	3459	7426	90%	0.0	89.60%	29681	NW_008920341.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5235	Nicotiana tomentosiformis	3445	7986	99%	0.0	89.88%	86109	NW_008935097.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold29425	Nicotiana tabacum	3437	7257	93%	0.0	91.82%	111475	NW_015887502.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold41737	Nicotiana tabacum	3435	7842	87%	0.0	94.76%	181014	NW_015905149.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold80492	Nicotiana tabacum	3426	6238	94%	0.0	88.44%	60977	NW_015939584.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1933	Nicotiana tabacum	3424	7499	89%	0.0	90.55%	91511	NW_015848101.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold60181	Nicotiana tabacum	3391	6321	80%	0.0	95.77%	7423	NW_015921951.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5078	Nicotiana tomentosiformis	3389	6951	97%	0.0	92.84%	90024	NW_008933352.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold43066	Nicotiana sylvestris	3384	13839	93%	0.0	96.77%	89833	NW_009529461.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7528	Nicotiana tabacum	3373	7846	92%	0.0	90.93%	191274	NW_015935130.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold122182	Nicotiana tabacum	3356	5175	87%	0.0	87.08%	33817	NW_015803903.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold15017	Nicotiana sylvestris	3356	7232	90%	0.0	91.97%	95267	NW_009394548.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13878	Nicotiana tomentosiformis	3356	9113	89%	0.0	92.97%	62040	NW_008871579.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold36865	Nicotiana sylvestris	3350	5148	87%	0.0	87.05%	36853	NW_009522570.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold81284	Nicotiana sylvestris	3345	9943	91%	0.0	94.37%	24576	NW_009571925.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold119706	Nicotiana tabacum	3339	7726	92%	0.0	91.10%	57114	NW_015802100.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold42528	Nicotiana tabacum	3336	5369	84%	0.0	87.01%	152515	NW_015905873.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold23607	Nicotiana tomentosiformis	3336	8164	96%	0.0	92.58%	180605	NW_008903161.1

Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16066	Nicotiana tomentosiformis	3330	5369	84%	0.0	86.98%	53501	NW_008894782.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold39391	Nicotiana attenuata	3330	5688	80%	0.0	89.15%	4408	NW_008920698.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana tomentosiformis	3326	5287	82%	0.0	88.52%	645505	NW_017670487.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2852	Nicotiana attenuata	3317	6912	92%	0.0	90.93%	203571	NW_008908619.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5342	Nicotiana tomentosiformis	3315	5939	84%	0.0	90.14%	279541	NW_008936286.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold98400	Nicotiana tabacum	3310	7863	90%	0.0	93.69%	19122	NW_015954207.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold48003	Nicotiana sylvestris	3306	6686	82%	0.0	93.25%	64775	NW_009534947.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17000	Nicotiana tabacum	3302	6824	91%	0.0	87.96%	114775	NW_015835169.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold73146	Nicotiana tomentosiformis	3301	6643	99%	0.0	89.72%	42694	NW_008958204.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold34503	Nicotiana tabacum	3295	6910	88%	0.0	90.74%	9343	NW_015892299.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold8924	Nicotiana tomentosiformis	3295	7592	98%	0.0	92.40%	99972	NW_008976085.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold39014	Nicotiana tabacum	3288	5935	84%	0.0	89.95%	85645	NW_015902622.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold4264	Nicotiana sylvestris	3288	8475	88%	0.0	94.25%	185422	NW_009528987.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold34927	Nicotiana tabacum	3284	6505	92%	0.0	91.27%	75068	NW_015896324.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold876	Nicotiana tomentosiformis	3282	7469	95%	0.0	90.89%	99473	NW_008974262.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4008	Nicotiana tomentosiformis	3278	11383	93%	0.0	91.23%	106059	NW_008921464.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold11904	Nicotiana tabacum	3275	7198	87%	0.0	90.16%	134089	NW_015801616.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold6132	Nicotiana tomentosiformis	3269	6670	94%	0.0	91.09%	160670	NW_008945064.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold52774	Nicotiana tabacum	3264	6659	94%	0.0	91.05%	59470	NW_015915283.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold15642	Nicotiana tomentosiformis	3264	7205	87%	0.0	90.08%	198956	NW_008891180.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold56075	Nicotiana tabacum	3262	7503	98%	0.0	92.14%	61093	NW_015918284.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold20979	Nicotiana tomentosiformis	3262	8244	97%	0.0	90.01%	34902	NW_008900240.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold92459	Nicotiana tabacum	3247	6961	93%	0.0	88.55%	51474	NW_015949511.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold12204	Nicotiana tomentosiformis	3243	7347	88%	0.0	90.33%	6935	NW_008852981.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold22119	Nicotiana sylvestris	3241	6956	93%	0.0	88.51%	138560	NW_009473459.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold72753	Nicotiana tomentosiformis	3238	5782	91%	0.0	87.84%	55730	NW_008957767.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold30619	Nicotiana tabacum	3236	7478	96%	0.0	92.13%	81030	NW_015888627.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold10437	Nicotiana tomentosiformis	3232	7262	89%	0.0	92.84%	33777	NW_008833349.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold74254	Nicotiana tabacum	3221	9440	85%	0.0	95.08%	180546	NW_015934251.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23928	Nicotiana tabacum	3219	6688	80%	0.0	89.84%	102261	NW_015870546.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold43911	Nicotiana sylvestris	3219	7829	82%	0.0	95.03%	50347	NW_009530400.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold5491	Nicotiana sylvestris	3210	8010	87%	0.0	94.35%	105891	NW_009542620.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1392	Nicotiana tomentosiformis	3208	6185	87%	0.0	89.64%	83327	NW_008872046.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold4755	Nicotiana sylvestris	3206	5003	80%	0.0	87.30%	72741	NW_009534442.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1010	Nicotiana sylvestris	3205	7327	80%	0.0	94.98%	25511	NW_009339915.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold16103	Nicotiana tabacum	3190	7826	98%	0.0	89.15%	52440	NW_015829832.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold26735	Nicotiana sylvestris	3182	8980	90%	0.0	95.57%	47503	NW_009511315.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold34860	Nicotiana tabacum	3175	6814	83%	0.0	90.58%	142117	NW_015895592.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1501	Nicotiana tomentosiformis	3171	7648	86%	0.0	92.55%	169954	NW_00884158.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold26750	Nicotiana tomentosiformis	3168	5460	88%	0.0	86.20%	95878	NW_008906653.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold107528	Nicotiana tabacum	3166	5475	88%	0.0	86.17%	92127	NW_015793041.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold15608	Nicotiana tomentosiformis	3142	5308	83%	0.0	88.01%	62917	NW_008890802.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1504	Nicotiana tabacum	3138	7081	88%	0.0	89.19%	166884	NW_015823133.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold69898	Nicotiana tomentosiformis	3138	5383	84%	0.0	87.46%	40158	NW_008954593.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold11291	Nicotiana tomentosiformis	3131	8099	97%	0.0	90.09%	48323	NW_008842836.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold62180	Nicotiana tabacum	3129	7019	90%	0.0	90.83%	107175	NW_015923736.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold10659	Nicotiana sylvestris	3123	7008	90%	0.0	90.79%	113562	NW_009346125.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5752	Nicotiana tabacum	3103	5647	90%	0.0	86.29%	201130	NW_015919603.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold11006	Nicotiana sylvestris	3103	5652	90%	0.0	86.29%	121736	NW_009349982.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1374	Nicotiana sylvestris	3101	8393	84%	0.0	96.34%	203238	NW_009380358.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold8512	Nicotiana sylvestris	3101	8604	91%	0.0	92.07%	98123	NW_009576187.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17486	Nicotiana tabacum	3099	7508	84%	0.0	91.79%	143640	NW_015837928.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold101469	Nicotiana tabacum	3092	6134	94%	0.0	88.07%	15174	NW_015788391.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 9, NIATT2	Nicotiana attenuata	3088	12200	93%	0.0	87.23%	50975570	NC_031997.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold69740	Nicotiana sylvestris	3085	10201	96%	0.0	95.15%	58209	NW_009559098.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold35139	Nicotiana tabacum	3079	8081	86%	0.0	90.14%	217011	NW_015898642.1

Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold26234	Nicotiana tomentosiformis	3072	6558	93%	0.0	89.13%	140437	NW_008906080.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold33670	Nicotiana tabacum	3070	10892	92%	0.0	89.88%	286809	NW_015891512.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1275	Nicotiana tomentosiformis	3068	7602	86%	0.0	90.06%	380063	NW_008859046.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold73607	Nicotiana tabacum	3066	6547	94%	0.0	89.06%	127436	NW_015933689.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold93771	Nicotiana tabacum	3057	9414	89%	0.0	93.02%	43316	NW_015950539.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold80715	Nicotiana tabacum	3053	6747	87%	0.0	87.59%	50554	NW_015939777.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold92714	Nicotiana tabacum	3048	6088	94%	0.0	87.76%	33110	NW_015949721.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold40668	Nicotiana tabacum	3046	7671	99%	0.0	89.81%	84363	NW_015904132.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold14849	Nicotiana tabacum	3035	8058	97%	0.0	89.51%	441724	NW_015821934.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold12852	Nicotiana tomentosiformis	3035	6442	85%	0.0	91.12%	112275	NW_008860180.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold24113	Nicotiana tomentosiformis	3029	7948	97%	0.0	92.74%	44503	NW_008903724.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold34201	Nicotiana tomentosiformis	3027	7340	90%	0.0	89.92%	88465	NW_008914933.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold26911	Nicotiana tomentosiformis	3020	7499	98%	0.0	91.61%	33862	NW_008906832.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold3177	Nicotiana tabacum	3014	7796	99%	0.0	90.35%	8934	NW_015889710.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold17029	Nicotiana sylvestris	3009	10451	93%	0.0	87.87%	143278	NW_009416903.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold38441	Nicotiana tabacum	3000	5986	82%	0.0	90.98%	28448	NW_015902092.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold54813	Nicotiana tabacum	2998	8699	91%	0.0	87.79%	73196	NW_015917151.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold15724	Nicotiana sylvestris	2996	8944	94%	0.0	94.77%	259450	NW_009402403.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold21037	Nicotiana sylvestris	2994	8896	82%	0.0	95.58%	66317	NW_009461437.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold59693	Nicotiana sylvestris	2992	7648	88%	0.0	95.43%	48294	NW_009547934.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9850	Nicotiana tomentosiformis	2992	12650	96%	0.0	89.98%	77036	NW_008986374.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold147526	Nicotiana tabacum	2983	7432	92%	0.0	91.85%	14952	NW_015821335.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold18076	Nicotiana sylvestris	2983	7428	92%	0.0	91.85%	78355	NW_009428536.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold116498	Nicotiana tabacum	2979	6308	88%	0.0	90.31%	85930	NW_015799768.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold59013	Nicotiana tabacum	2972	7805	89%	0.0	93.82%	43865	NW_015920918.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold8368	Nicotiana sylvestris	2968	6512	91%	0.0	86.18%	74886	NW_009574586.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold37240	Nicotiana tabacum	2955	7552	89%	0.0	92.85%	203128	NW_015900952.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold24683	Nicotiana tomentosiformis	2953	3208	90%	0.0	78.84%	153191	NW_008904356.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	2948	6578	87%	0.0	94.77%	322487	NW_017671282.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold51285	Nicotiana tabacum	2935	6988	85%	0.0	93.53%	21056	NW_015913910.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1472	Nicotiana tabacum	2933	6329	85%	0.0	90.21%	38084	NW_015821125.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold55250	Nicotiana tomentosiformis	2924	7728	99%	0.0	89.58%	48675	NW_008938320.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold2926	Nicotiana sylvestris	2911	11420	94%	0.0	95.89%	88958	NW_009514120.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold38135	Nicotiana tomentosiformis	2905	9344	91%	0.0	87.98%	14530	NW_008919303.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold20743	Nicotiana sylvestris	2892	7853	85%	0.0	95.20%	55320	NW_009458170.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold25200	Nicotiana tomentosiformis	2892	8208	100%	0.0	88.77%	5933	NW_008904932.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold210201	Nicotiana tabacum	2880	5263	88%	0.0	88.79%	38016	NW_015856775.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold86306	Nicotiana tomentosiformis	2876	6882	81%	0.0	92.19%	57737	NW_008972826.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold17063	Nicotiana sylvestris	2844	4440	80%	0.0	85.65%	134932	NW_009417281.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	2828	5526	82%	0.0	89.48%	130703	NW_017673533.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold28712	Nicotiana tabacum	2819	4967	80%	0.0	85.43%	92604	NW_015886830.1
Tnt1	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2819	28455	90%	0.0	83.83%	220335243	NC_029987.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold14842	Nicotiana sylvestris	2813	6549	82%	0.0	93.95%	65318	NW_009392603.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold16129	Nicotiana sylvestris	2798	10062	96%	0.0	95.26%	37172	NW_009406903.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold20597	Nicotiana sylvestris	2776	7446	86%	0.0	96.71%	47589	NW_009456547.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold401	Nicotiana tomentosiformis	2767	7,04E+03	98%	0.0	91.70%	14011	NW_008921486.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4922	Nicotiana tabacum	2761	5690	82%	0.0	89.20%	46881	NW_015912036.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold3398	Nicotiana sylvestris	2756	5272	82%	0.0	89.15%	44816	NW_009519364.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold27331	Nicotiana tabacum	2750	7531	89%	0.0	91.57%	55300	NW_015885520.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold10240	Nicotiana sylvestris	2750	9803	87%	0.0	95.50%	329378	NW_009341471.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold47393	Nicotiana tabacum	2717	7328	91%	0.0	92.93%	130654	NW_015910351.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold72564	Nicotiana tabacum	2717	6613	88%	0.0	90.54%	107847	NW_015932816.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold85240	Nicotiana tabacum	2715	5401	82%	0.0	87.27%	68646	NW_015943551.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold20574	Nicotiana tomentosiformis	2713	6804	90%	0.0	90.02%	461403	NW_008899791.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13058	Nicotiana tabacum	2702	7541	97%	0.0	90.17%	112303	NW_015809906.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold48757	Nicotiana tomentosiformis	2699	5368	82%	0.0	87.15%	73063	NW_008931104.1

Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold70444	Nicotiana tabacum	2673	6087	90%	0.0	87.37%	36701	NW_015930969.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13689	Nicotiana tomentosiformis	2671	7186	91%	0.0	92.51%	133199	NW_008869479.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	2662	5803	90%	0.0	89.17%	95677	NW_017674462.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold44807	Nicotiana tabacum	2654	7250	89%	0.0	92.23%	69939	NW_015907959.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold58773	Nicotiana tomentosiformis	2639	6674	85%	0.0	87.53%	36426	NW_008942233.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold26477	Nicotiana tabacum	2606	11137	90%	0.0	96.69%	88246	NW_015882268.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5387	Nicotiana tabacum	2606	6031	88%	0.0	88.12%	75258	NW_015916293.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold41327	Nicotiana tabacum	2591	8216	83%	0.0	95.66%	109349	NW_015904764.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold70805	Nicotiana tabacum	2590	7614	83%	0.0	95.66%	22930	NW_015931289.1
Tnt1	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2556	13701	90%	0.0	84.00%	238794889	NC_029985.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5745	Nicotiana tomentosiformis	2523	7105	87%	0.0	91.76%	24249	NW_008940763.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold68287	Nicotiana tabacum	2510	6861	96%	0.0	92.08%	67980	NW_015929099.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13953	Nicotiana tomentosiformis	2505	6866	96%	0.0	92.02%	24540	NW_008872413.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16273	Nicotiana tomentosiformis	2488	4922	81%	0.0	87.19%	137894	NW_008895012.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold19486	Nicotiana sylvestris	2475	5841	87%	0.0	87.38%	107802	NW_009444202.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4023	Nicotiana tabacum	2471	4854	81%	0.0	87.06%	129100	NW_015903740.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold22524	Nicotiana sylvestris	2431	7190	83%	0.0	86.58%	64465	NW_009477959.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold93472	Nicotiana tabacum	2425	5034	84%	0.0	85.66%	66433	NW_015950315.1
Tnt1	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2407	9057	89%	0.0	84.29%	153299543	NC_029984.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold11245	Nicotiana tabacum	2405	4748	81%	0.0	87.41%	69628	NW_015796773.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31677	Nicotiana tomentosiformis	2405	7445	89%	0.0	92.42%	79950	NW_008912127.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1319	Nicotiana sylvestris	2390	8074	86%	0.0	94.32%	147785	NW_009374247.1
Tnt1	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2381	16334	89%	0.0	83.02%	163962470	NC_029978.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold172870	Nicotiana tabacum	2374	8462	86%	0.0	94.27%	30121	NW_015836793.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold54283	Nicotiana tabacum	2361	6820	83%	0.0	91.54%	44199	NW_015916675.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold44259	Nicotiana tabacum	2335	8288	92%	0.0	93.77%	118404	NW_015907468.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23404	Nicotiana tabacum	2327	5895	86%	0.0	91.90%	159668	NW_015868073.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold26405	Nicotiana tomentosiformis	2320	7168	90%	0.0	91.26%	55726	NW_008906270.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold3296	Nicotiana tabacum	2314	7162	90%	0.0	91.20%	100345	NW_015890832.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold42555	Nicotiana tomentosiformis	2305	5856	86%	0.0	91.66%	61547	NW_008924214.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold23351	Nicotiana sylvestris	2292	6186	89%	0.0	87.48%	86726	NW_009487148.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold40885	Nicotiana tabacum	2287	5293	83%	0.0	87.43%	37627	NW_015904340.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold20836	Nicotiana tomentosiformis	2263	6242	86%	0.0	90.40%	219098	NW_00890082.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold35482	Nicotiana tabacum	2257	6270	86%	0.0	90.34%	213828	NW_015899311.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold927	Nicotiana tomentosiformis	2246	6026	85%	0.0	85.40%	76962	NW_008979929.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold7404	Nicotiana tomentosiformis	2237	6842	82%	0.0	91.20%	131825	NW_008959197.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold35181	Nicotiana tabacum	2233	6013	85%	0.0	85.28%	70263	NW_015899026.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold153621	Nicotiana tabacum	2220	5015	80%	0.0	88.55%	36013	NW_015825212.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold61282	Nicotiana tabacum	2213	5271	83%	0.0	88.73%	21861	NW_015922930.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold53	Nicotiana tomentosiformis	2174	5540	88%	0.0	88.56%	32298	NW_008935818.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold23160	Nicotiana tomentosiformis	2161	7264	91%	0.0	88.53%	180509	NW_008902665.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13876	Nicotiana tabacum	2156	7332	85%	0.0	92.46%	455350	NW_015815526.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold107939	Nicotiana tabacum	2150	6336	91%	0.0	88.42%	58463	NW_015793367.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2024	Nicotiana tomentosiformis	2145	7345	85%	0.0	92.33%	249441	NW_008899420.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1419	Nicotiana tabacum	2091	7982	82%	0.0	91.27%	482341	NW_015817660.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold11793	Nicotiana tomentosiformis	2063	6129	84%	0.0	90.04%	70473	NW_008848413.1
Tnt1	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold4971	Capsicum annuum	2037	4095	82%	0.0	83.20%	81603	NW_015960449.1
Tnt1	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold8548	Capsicum annuum	2026	4034	81%	0.0	83.08%	87797	NW_015961293.1
Tnt1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold31276	Nicotiana sylvestris	2026	5564	89%	0.0	89.96%	150718	NW_009516361.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold37360	Nicotiana tabacum	2021	5551	90%	0.0	89.55%	65958	NW_015901068.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	2017	6870	83%	0.0	86.45%	291638	NW_017671474.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31638	Nicotiana tomentosiformis	1999	7105	93%	0.0	91.79%	6946	NW_008912084.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold11874	Nicotiana tabacum	1949	7610	87%	0.0	92.74%	88948	NW_015801395.1
Tnt1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	1906	5098	87%	0.0	86.49%	120963	NW_017673766.1
Tnt1	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00160	Solanum tuberosum	1801	5602	89%	0.0	84.02%	1367927	NW_006239085.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold43392	Nicotiana tabacum	1792	6297	86%	0.0	93.73%	17993	NW_015906674.1

Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold20884	Nicotiana tomentosiformis	1777	3765	88%	0.0	79.34%	54029	NW_008900135.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold64336	Nicotiana tabacum	1768	6504	87%	0.0	91.02%	93867	NW_015925664.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold167272	Nicotiana tabacum	1729	3815	85%	0.0	79.38%	29890	NW_015833543.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold50715	Nicotiana tabacum	1711	8509	93%	0.0	90.92%	269829	NW_015913397.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold18882	Nicotiana tomentosiformis	1705	6915	87%	0.0	90.84%	79553	NW_008897910.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold50455	Nicotiana tabacum	1694	4557	86%	0.0	78.84%	111537	NW_015913157.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2217	Nicotiana tomentosiformis	1694	4535	86%	0.0	78.84%	179490	NW_008901564.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold3485	Nicotiana tabacum	1688	4319	87%	0.0	78.76%	194765	NW_015895481.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold12997	Nicotiana tomentosiformis	1685	6192	86%	0.0	78.84%	103562	NW_008861790.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold16974	Nicotiana tabacum	1679	3935	87%	0.0	78.69%	279842	NW_015835004.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold9564	Nicotiana tomentosiformis	1676	4289	87%	0.0	78.70%	153237	NW_008893196.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1235	Nicotiana tomentosiformis	1672	3476	85%	0.0	78.71%	126248	NW_008854602.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold75810	Nicotiana tabacum	1655	3443	84%	0.0	78.58%	64949	NW_015935585.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1710	Nicotiana tomentosiformis	1653	3818	84%	0.0	78.62%	301715	NW_008895931.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold18213	Nicotiana tabacum	1650	3800	84%	0.0	78.59%	106330	NW_015841996.1
Tnt1	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref.v1.0.scaffold2530	Capsicum annuum	1644	4050	89%	0.0	83.84%	86802	NW_015961129.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold22651	Nicotiana tabacum	1637	2982	80%	0.0	78.40%	63816	NW_015864550.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9847	Nicotiana tabacum	1626	6111	85%	0.0	78.41%	25355	NW_015954256.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold203	Nicotiana tabacum	1624	3709	87%	0.0	78.57%	178765	NW_015853131.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold67533	Nicotiana tabacum	1618	3380	86%	0.0	78.35%	55394	NW_015928450.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24108	Nicotiana tabacum	1583	3603	83%	0.0	78.08%	51010	NW_015871358.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold51855	Nicotiana tabacum	1583	3776	84%	0.0	78.25%	47500	NW_015914429.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold28420	Nicotiana tomentosiformis	1583	3804	85%	0.0	78.09%	25282	NW_008908509.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold23480	Nicotiana tomentosiformis	1576	3702	86%	0.0	78.13%	107486	NW_008903020.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3009	Nicotiana tomentosiformis	1555	3497	83%	0.0	78.44%	386545	NW_008910364.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold19145	Nicotiana tabacum	1554	3512	83%	0.0	78.45%	143789	NW_015847104.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold14090	Nicotiana tomentosiformis	1535	3735	85%	0.0	82.82%	71349	NW_008873936.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold58837	Nicotiana tomentosiformis	1533	3649	86%	0.0	77.84%	52170	NW_008942304.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold31971	Nicotiana tabacum	1528	3745	86%	0.0	77.73%	116111	NW_015889096.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5848	Nicotiana tabacum	1524	3599	87%	0.0	77.94%	217698	NW_015920454.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1728	Nicotiana tomentosiformis	1513	3537	88%	0.0	77.88%	194334	NW_008896130.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold46524	Nicotiana tomentosiformis	1513	2969	80%	0.0	82.58%	63442	NW_008928624.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold78966	Nicotiana tomentosiformis	1511	3357	81%	0.0	82.63%	13557	NW_008964669.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold36034	Nicotiana tomentosiformis	1493	3087	84%	0.0	77.67%	59025	NW_008916969.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold29904	Nicotiana tomentosiformis	1491	2969	80%	0.0	77.53%	30088	NW_008910157.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold47509	Nicotiana tabacum	1480	3304	85%	0.0	77.41%	157477	NW_015910454.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold32411	Nicotiana tabacum	1476	3274	83%	0.0	77.67%	68176	NW_015890322.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold17521	Nicotiana tomentosiformis	1472	3287	84%	0.0	77.41%	86450	NW_008896399.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31606	Nicotiana tomentosiformis	1469	2883	80%	0.0	77.30%	47854	NW_008912049.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold53397	Nicotiana tabacum	1467	2899	80%	0.0	82.17%	123130	NW_015915862.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold24563	Nicotiana tomentosiformis	1463	2908	80%	0.0	77.31%	85385	NW_008904223.1
Tnt1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1504	Nicotiana tomentosiformis	1395	3746	84%	0.0	81.61%	237830	NW_008884491.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold67586	Nicotiana tabacum	1384	3788	85%	0.0	81.43%	29372	NW_015928500.1
Tnt1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold38322	Nicotiana tabacum	1334	2828	82%	0.0	82.68%	133131	NW_015901980.1
Tto1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold6042	Nicotiana tabacum	9537	11552	100%	0.0	99.13%	158584	NW_015922161.1
Tto1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8326	Nicotiana tabacum	9535	11602	100%	0.0	99.08%	90194	NW_015941905.1
Tto1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold3934	Nicotiana tabacum	9529	11605	100%	0.0	99.06%	49271	NW_015902918.1
Tto1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold2835	Nicotiana sylvestris	9516	11580	100%	0.0	99.02%	250491	NW_009513109.1
Tto1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold23141	Nicotiana sylvestris	7312	9909	86%	0.0	99.38%	110081	NW_009484815.1
Tto1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold82885	Nicotiana tabacum	6876	9915	82%	0.0	99.14%	36495	NW_015941609.1
CopiaSL_17	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	14678	50030	100%	0.0	100.00%	66723567	NC_015442.3
CopiaSL_17	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	10813	51735	100%	0.0	92.63%	72906345	NC_015446.3
CopiaSL_17	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	9550	14840	100%	0.0	94.93%	56597135	NC_015448.3
CopiaSL_17	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	9284	39794	100%	0.0	94.26%	98455869	NC_015438.3
CopiaSL_17	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	8890	31717	100%	0.0	93.02%	68175699	NC_015444.3
CopiaSL_17	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	8468	25126	100%	0.0	91.81%	65987440	NC_015445.3

CopiaSL_17	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	8460	27784	100%	0.0	91.72%	49794276	NC_015443.3
CopiaSL_17	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	6617	22128	100%	0.0	86.47%	68126176	NC_015449.3
CopiaSL_17	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	5945	37669	100%	0.0	94.88%	65633393	NC_015447.3
CopiaSL_17	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	4217	19828	94%	0.0	94.89%	66557038	NC_015441.3
CopiaSL_17	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	10209	44510	100%	0.0	89.91%	109333515	NC_028637.1
CopiaSL_17	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	Solanum pennellii	9858	49003	100%	0.0	91.86%	84057508	NC_028645.1
CopiaSL_17	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	9696	83896	100%	0.0	95.35%	60730942	NC_028642.1
CopiaSL_17	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	Solanum pennellii	9666	37614	100%	0.0	95.29%	79292169	NC_028643.1
CopiaSL_17	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	9217	19818	100%	0.0	93.98%	75414019	NC_028639.1
CopiaSL_17	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum pennellii	9132	49987	100%	0.0	94.43%	77991103	NC_028641.1
CopiaSL_17	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum pennellii	8894	46157	100%	0.0	93.01%	82529941	NC_028646.1
CopiaSL_17	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	Solanum pennellii	8530	2,87E+04	100%	0.0	92.00%	66223686	NC_028647.1
CopiaSL_17	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	7607	24147	99%	0.0	96.31%	59803892	NC_028638.1
CopiaSL_17	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	Solanum pennellii	7308	24596	100%	0.0	88.39%	83305730	NC_028648.1
CopiaSL_17	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	6652	33802	100%	0.0	88.73%	77197300	NC_028640.1
CopiaSL_33	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	10211	1,01E+08	100%	0.0	100.0%	56597135	NC_015448.3
CopiaSL_33	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	9699	2,92E+08	100%	0.0	98.32%	109333515	NC_028637.1
CopiaSL_33	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	Solanum pennellii	9699	95478	100%	0.0	98.32%	66223686	NC_028647.1
CopiaSL_33	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	9697	1,03E+08	100%	0.0	98.32%	59803892	NC_028638.1
CopiaSL_33	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum pennellii	9696	3,06E+08	100%	0.0	98.32%	77991103	NC_028641.1
CopiaSL_33	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	Solanum pennellii	9696	2,31E+08	100%	0.0	98.32%	83305730	NC_028648.1
CopiaSL_33	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	Solanum pennellii	9692	1,74E+08	100%	0.0	98.30%	70546659	NC_028644.1
CopiaSL_33	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	9681	1,68E+08	100%	0.0	98.27%	75414019	NC_028639.1
CopiaSL_33	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum pennellii	9675	1,82E+08	100%	0.0	98.25%	82529941	NC_028646.1
CopiaSL_33	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	Solanum pennellii	9673	1,39E+08	100%	0.0	98.25%	84057508	NC_028645.1
CopiaSL_33	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	9662	1,96E+08	100%	0.0	98.21%	77197300	NC_028640.1
CopiaSL_33	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	9659	91978	100%	0.0	98.20%	60730942	NC_028642.1
CopiaSL_33	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	Solanum pennellii	9548	1,03E+08	100%	0.0	97.83%	79292169	NC_028643.1
CopiaSL_33	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	9548	1,72E+08	100%	0.0	97.82%	72906345	NC_015446.3
CopiaSL_33	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	9522	3,23E+08	100%	0.0	97.74%	98455869	NC_015438.3
CopiaSL_33	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	9454	1,16E+08	100%	0.0	97.49%	68175699	NC_015444.3
CopiaSL_33	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	9382	2,47E+08	100%	0.0	97.26%	65633393	NC_015447.3
CopiaSL_33	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	9321	1,35E+08	100%	0.0	97.18%	66557038	NC_015441.3
CopiaSL_33	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	9306	1,81E+08	100%	0.0	96.98%	66723567	NC_015442.3
CopiaSL_33	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	9306	2,02E+08	100%	0.0	97.04%	65987440	NC_015445.3
CopiaSL_33	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	9274	2,02E+08	100%	0.0	96.93%	68126176	NC_015449.3
CopiaSL_33	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	9153	1,44E+08	100%	0.0	96.55%	72290146	NC_015440.3
CopiaSL_33	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	8691	85080	100%	0.0	95.02%	55977580	NC_015439.3
CopiaSL_33	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	8045	83241	100%	0.0	96.55%	49794276	NC_015443.3
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	6543	3,94E+08	100%	0.0	88.13%	301019445	NC_029977.1
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	6338	2,11E+08	100%	0.0	87.57%	238794889	NC_029985.1
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	6237	3,11E+08	100%	0.0	87.23%	219521584	NC_029982.1
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	6220	2,43E+08	100%	0.0	87.10%	217274494	NC_029981.1
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1281	Capsicum annuum	6216	7186	100%	0.0	87.12%	233328	NW_015961028.1
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	6211	1,94E+08	100%	0.0	87.06%	215701946	NC_029980.1
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	6205	2,82E+08	100%	0.0	87.10%	261510930	NC_029979.1
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	6204	2,31E+08	100%	0.0	87.07%	220335243	NC_029987.1
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	6200	1,78E+08	100%	0.0	87.26%	153299543	NC_029984.1
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	6154	1,43E+08	100%	0.0	86.87%	205736368	NC_029986.1
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	6130	2,74E+08	100%	0.0	86.81%	229934170	NC_029988.1
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold592	Capsicum annuum	6120	1,25E+04	100%	0.0	86.75%	1905997	NW_015960988.1
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1037	Capsicum annuum	6106	7072	100%	0.0	86.77%	2186250	NW_015960681.1
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	6104	1,38E+08	100%	0.0	88.93%	163962470	NC_029978.1
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	6104	2,02E+08	100%	0.0	86.76%	222112641	NC_029983.1
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2348	Capsicum annuum	6063	6063	99%	0.0	86.63%	373378	NW_015960596.1
CopiaSL_33	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1422	Capsicum annuum	5927	22010	100%	0.0	86.70%	802096	NW_015960747.1

CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1633	Capisicum annum	5884	6234	97%	0.0	86.58%	1620172	NW_015961041.1
CopiaSL_33	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00029	Solanum tuberosum	5847	6286	90%	0.0	89.65%	2785282	NW_006238955.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1390	Capisicum annum	5681	5681	100%	0.0	85.39%	1730405	NW_015960783.1
CopiaSL_33	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00015	Solanum tuberosum	5677	5677	93%	0.0	86.60%	3327864	NW_006238941.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1341	Capisicum annum	5644	5644	99%	0.0	85.27%	1906221	NW_015960655.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold9	Capisicum annum	5546	1,02E+04	100%	0.0	84.93%	6567951	NW_015961815.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold688	Capisicum annum	5518	7,82E+03	96%	0.0	88.42%	1322070	NW_015960950.1
CopiaSL_33	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00022	Solanum tuberosum	5505	6241	100%	0.0	87.42%	3032639	NW_006238948.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3844	Capisicum annum	5485	5904	99%	0.0	85.27%	179318	NW_015961259.1
CopiaSL_33	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00069	Solanum tuberosum	5404	5633	87%	0.0	87.08%	2089258	NW_006238995.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold171	Capisicum annum	5358	5736	99%	0.0	85.08%	887051	NW_015960473.1
CopiaSL_33	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00066	Solanum tuberosum	5339	7639	84%	0.0	89.07%	2159599	NW_006238989.1
CopiaSL_33	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00120	Solanum tuberosum	5299	5299	87%	0.0	86.69%	1603502	NW_006239047.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold301	Capisicum annum	5273	5732	99%	0.0	84.30%	1226190	NW_015960697.1
CopiaSL_33	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00089	Solanum tuberosum	5251	13151	97%	0.0	86.53%	1856049	NW_006239012.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold148	Capisicum annum	5169	6820	100%	0.0	87.61%	1323090	NW_015961290.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1124	Capisicum annum	5169	7819	97%	0.0	84.31%	2110277	NW_015961370.1
CopiaSL_33	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00083	Solanum tuberosum	5162	5162	80%	0.0	87.91%	1836497	NW_006239015.1
CopiaSL_33	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00384	Solanum tuberosum	5156	5871	100%	0.0	87.53%	565346	NW_006239308.1
CopiaSL_33	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00589	Solanum tuberosum	5003	5069	85%	0.0	86.29%	277750	NW_006239511.1
CopiaSL_33	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00009	Solanum tuberosum	4955	31319	97%	0.0	89.31%	3955149	NW_006238935.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold372	Capisicum annum	4907	14284	100%	0.0	88.79%	2397553	NW_015961318.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold6868	Capisicum annum	4553	5032	90%	0.0	86.13%	22397	NW_015961342.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1973	Capisicum annum	4549	5100	90%	0.0	85.91%	1714326	NW_015961422.1
CopiaSL_33	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00038	Solanum tuberosum	4514	6576	90%	0.0	88.73%	2533956	NW_006238966.1
CopiaSL_33	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00295	Solanum tuberosum	4457	5582	81%	0.0	89.74%	825164	NW_006239214.1
CopiaSL_33	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13433	Nicotiana tomentosiformis	4346	4346	82%	0.0	84.03%	64717	NW_008866636.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1780	Capisicum annum	4331	4331	85%	0.0	83.38%	226878	NW_015961280.1
CopiaSL_33	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold15440	Nicotiana tabacum	4313	4313	82%	0.0	83.90%	59730	NW_015825707.1
CopiaSL_33	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold90048	Nicotiana tabacum	4301	4301	83%	0.0	83.74%	82605	NW_015947518.1
CopiaSL_33	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold24998	Nicotiana tomentosiformis	4292	4292	82%	0.0	83.71%	59222	NW_008904705.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold756	Capisicum annum	4226	8209	99%	0.0	86.38%	884456	NW_015961254.1
CopiaSL_33	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00086	Solanum tuberosum	4132	9369	94%	0.0	90.07%	1873685	NW_006239010.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold551	Capisicum annum	4130	5360	96%	0.0	84.38%	761547	NW_015961519.1
CopiaSL_33	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00606	Solanum tuberosum	4119	6251	89%	0.0	85.91%	265855	NW_006239521.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2239	Capisicum annum	3947	4893	99%	0.0	84.33%	615095	NW_015960816.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1378	Capisicum annum	3871	7155	98%	0.0	86.57%	1057095	NW_015961130.1
CopiaSL_33	refseq_genomes	8	Capisicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1688	Capisicum annum	3775	7134	100%	0.0	86.29%	693367	NW_015961726.1
CopiaSL_33	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00050	Solanum tuberosum	3705	5632	90%	0.0	87.37%	2328283	NW_006238975.1
CopiaSL_33	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold40802	Nicotiana tomentosiformis	3410	4201	81%	0.0	85.97%	67366	NW_008922267.1
CopiaSL_33	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1947	Nicotiana tabacum	3398	4184	81%	0.0	85.91%	135503	NW_015848868.1
Tork4/CopiaSL_37	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	8861	1,78E+08	100%	0.0	99.53%	55977580	NC_015439.3
Tork4/CopiaSL_37	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	8861	2,35E+08	100%	0.0	99.53%	68175699	NC_015444.3
Tork4/CopiaSL_37	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	8861	2,00E+08	100%	0.0	99.53%	65987440	NC_015445.3
Tork4/CopiaSL_37	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	8861	2,75E+08	100%	0.0	99.53%	65633393	NC_015447.3
Tork4/CopiaSL_37	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	8855	3,13E+08	100%	0.0	99.51%	98455869	NC_015438.3
Tork4/CopiaSL_37	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	8853	1,93E+08	100%	0.0	99.51%	66723567	NC_015442.3
Tork4/CopiaSL_37	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	8809	1,33E+08	100%	0.0	99.32%	56597135	NC_015448.3
Tork4/CopiaSL_37	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	8643	1,39E+08	100%	0.0	99.37%	49794276	NC_015443.3
Tork4/CopiaSL_37	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	8637	1,84E+08	100%	0.0	98.81%	68126176	NC_015449.3
Tork4/CopiaSL_37	refseq_genomes	8	Solanum pennellii chromosome 4, SPENNIV200	Solanum pennellii	8634	2,00E+08	100%	0.0	98.69%	77197300	NC_028640.1
Tork4/CopiaSL_37	refseq_genomes	8	Solanum pennellii chromosome 10, SPENNIV200	Solanum pennellii	8624	2,42E+08	100%	0.0	98.64%	82529941	NC_028646.1
Tork4/CopiaSL_37	refseq_genomes	8	Solanum pennellii chromosome 11, SPENNIV200	Solanum pennellii	8624	1,17E+08	100%	0.0	98.64%	66223686	NC_028647.1
Tork4/CopiaSL_37	refseq_genomes	8	Solanum pennellii chromosome 7, SPENNIV200	Solanum pennellii	8623	1,57E+08	100%	0.0	98.64%	79292169	NC_028643.1
Tork4/CopiaSL_37	refseq_genomes	8	Solanum pennellii chromosome 9, SPENNIV200	Solanum pennellii	8623	2,35E+08	100%	0.0	98.64%	84057508	NC_028645.1
Tork4/CopiaSL_37	refseq_genomes	8	Solanum pennellii chromosome 6, SPENNIV200	Solanum pennellii	8619	2,23E+08	100%	0.0	98.62%	60730942	NC_028642.1

Tork4/CopiaSL_37	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	8617	4,39E+08	100%	0.0	98.62%	109333515	NC_028637.1
Tork4/CopiaSL_37	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	8421	1,58E+08	100%	0.0	99.36%	72906345	NC_015446.3
Tork4/CopiaSL_37	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	8410	99664	100%	0.0	98.53%	75414019	NC_028639.1
Tork4/CopiaSL_37	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum pennellii	8351	1,32E+08	100%	0.0	98.11%	77991103	NC_028641.1
Tork4/CopiaSL_37	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	8312	53760	100%	0.0	98.76%	59803892	NC_028638.1
Tork4/CopiaSL_37	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	Solanum pennellii	8305	1,22E+08	100%	0.0	98.33%	70546659	NC_028644.1
Tork4/CopiaSL_37	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	8045	1,03E+08	100%	0.0	99.55%	72290146	NC_015440.3
Tork4/CopiaSL_37	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	Solanum pennellii	7854	1,31E+08	100%	0.0	98.69%	83305730	NC_028648.1
Tork4/CopiaSL_37	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	7851	1,66E+08	100%	0.0	99.33%	66557038	NC_015441.3
Tork4/CopiaSL_37	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC0000343	Solanum lycopersicum	6078	6635	86%	0.0	95.38%	5126	NW_020442824.1
Tork4/CopiaSL_37	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC0000003	Solanum lycopersicum	4021	7,25E+03	80%	0.0	94.89%	1066869	NW_020442484.1
CopiaSL_30	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	8468	3,54E+08	100%	0.0	100.00%	72906345	NC_015446.3
CopiaSL_30	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum pennellii	7411	4,43E+08	100%	0.0	95.84%	82529941	NC_028646.1
CopiaSL_30	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	7409	5,11E+08	100%	0.0	95.84%	109333515	NC_028637.1
CopiaSL_30	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	Solanum pennellii	7382	2,83E+08	100%	0.0	95.73%	79292169	NC_028643.1
CopiaSL_30	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	7376	2,17E+08	100%	0.0	95.71%	77197300	NC_028640.1
CopiaSL_30	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	7372	2,13E+08	100%	0.0	95.69%	75414019	NC_028639.1
CopiaSL_30	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	7369	1,58E+08	100%	0.0	95.65%	59803892	NC_028638.1
CopiaSL_30	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	Solanum pennellii	7345	1,63E+08	100%	0.0	95.58%	66223686	NC_028647.1
CopiaSL_30	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum pennellii	7323	2,56E+08	100%	0.0	95.49%	77991103	NC_028641.1
CopiaSL_30	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	7319	2,43E+08	100%	0.0	95.49%	60730942	NC_028642.1
CopiaSL_30	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	Solanum pennellii	7302	2,85E+08	100%	0.0	95.72%	83305730	NC_028648.1
CopiaSL_30	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	Solanum pennellii	7289	1,98E+08	100%	0.0	95.34%	70546659	NC_028644.1
CopiaSL_30	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	Solanum pennellii	7249	3,06E+08	100%	0.0	95.16%	84057508	NC_028645.1
CopiaSL_30	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	7143	3,29E+08	100%	0.0	94.81%	65633393	NC_015447.3
CopiaSL_30	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	7125	2,75E+08	100%	0.0	94.72%	72290146	NC_015440.3
CopiaSL_30	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	7038	2,29E+08	100%	0.0	94.39%	66723567	NC_015442.3
CopiaSL_30	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	7025	2,29E+08	100%	0.0	94.38%	49794276	NC_015443.3
CopiaSL_30	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	6966	4,46E+08	100%	0.0	94.09%	98455869	NC_015438.3
CopiaSL_30	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	6953	2,27E+08	100%	0.0	94.09%	65987440	NC_015445.3
CopiaSL_30	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	6905	1,26E+08	100%	0.0	94.14%	56597135	NC_015448.3
CopiaSL_30	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	6846	2,44E+08	100%	0.0	93.64%	66557038	NC_015441.3
CopiaSL_30	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	6841	2,81E+08	100%	0.0	93.61%	68126176	NC_015449.3
CopiaSL_30	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	6796	3,28E+08	100%	0.0	93.44%	68175699	NC_015444.3
CopiaSL_30	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	6721	1,72E+08	100%	0.0	93.25%	55977580	NC_015439.3
CopiaSL_30	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC000001	Solanum lycopersicum	6266	11811	98%	0.0	92.98%	2014460	NW_020442482.1
CopiaSL_30	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC000003	Solanum lycopersicum	5986	6560	100%	0.0	91.67%	1066869	NW_020442484.1
CopiaSL_30	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC000002	Solanum lycopersicum	5904	10556	100%	0.0	89.95%	177393	NW_020442483.1
CopiaSL_30	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC000009	Solanum lycopersicum	5816	6098	97%	0.0	90.12%	797759	NW_020442490.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3928	30145	91%	0.0	83.61%	219521584	NC_029982.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3849	52813	92%	0.0	83.26%	301019445	NC_029977.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3836	56884	93%	0.0	83.78%	217274494	NC_029981.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3786	35908	92%	0.0	83.50%	261510930	NC_029979.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3733	53418	92%	0.0	83.37%	215701946	NC_029980.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3731	50554	93%	0.0	82.77%	220335243	NC_029987.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3685	21023	91%	0.0	82.60%	163962470	NC_029978.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3639	35641	95%	0.0	83.02%	205736368	NC_029986.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3637	41555	93%	0.0	82.67%	229934170	NC_029988.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold920	Capsicum annuum	3500	3500	85%	0.0	82.95%	197492	NW_015961621.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold148	Capsicum annuum	3482	3482	85%	0.0	82.81%	1323090	NW_015961290.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1399	Capsicum annuum	3422	6393	88%	0.0	81.90%	1742971	NW_015961630.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2138	Capsicum annuum	3376	3376	91%	0.0	81.39%	681955	NW_015960370.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3373	48018	92%	0.0	81.29%	222112641	NC_029983.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold688	Capsicum annuum	3345	5329	92%	0.0	81.25%	1322070	NW_015960950.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3153	7700	88%	0.0	81.33%	238794889	NC_029985.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold258	Capsicum annuum	3094	3094	82%	0.0	81.54%	3098259	NW_015961180.1

CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2494	9408	89%	0.0	82.57%	153299543	NC_029984.1
CopiaSL_30	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1647	Capsicum annuum	2335	3244	91%	0.0	80.40%	1362880	NW_015960892.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	14837	58564	100%	0.0	100.00%	68126176	NC_015449.3
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	Solanum pennellii	9230	90913	95%	0.0	94.74%	84057508	NC_028645.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	9195	47180	100%	0.0	93.57%	55977580	NC_015439.3
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	Solanum pennellii	9092	90872	95%	0.0	94.11%	70546659	NC_028644.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	8997	62871	94%	0.0	93.99%	65987440	NC_015445.3
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	8977	74662	95%	0.0	93.97%	72290146	NC_015440.3
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum pennellii	8966	81011	96%	0.0	93.70%	82529941	NC_028646.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	8831	1,11E+08	98%	0.0	93.14%	109333515	NC_028637.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	8656	69748	95%	0.0	92.99%	49794276	NC_015443.3
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	8637	56042	95%	0.0	92.58%	66723567	NC_015442.3
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	Solanum pennellii	8253	56817	99%	0.0	90.84%	66223686	NC_028647.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00177	Solanum tuberosum	7827	12530	94%	0.0	89.98%	1275411	NW_006239101.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	7592	72522	95%	0.0	93.71%	72906345	NC_015446.3
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	7502	46944	95%	0.0	88.91%	56597135	NC_015448.3
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00405	Solanum tuberosum	7496	12848	91%	0.0	89.55%	533214	NW_006239323.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	7145	71161	94%	0.0	92.13%	98455869	NC_015438.3
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	6661	79660	94%	0.0	93.63%	68175699	NC_015444.3
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	Solanum pennellii	6523	93713	96%	0.0	87.94%	79292169	NC_028643.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00020	Solanum tuberosum	6274	20400	90%	0.0	91.88%	3188724	NW_006238946.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	Solanum pennellii	6054	1,35E+08	100%	0.0	96.13%	83305730	NC_028648.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC0000001	Solanum lycopersicum	5843	10790	90%	0.0	90.28%	2014460	NW_020442482.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	5819	67622	82%	0.0	95.59%	77197300	NC_028640.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum pennellii	5812	75023	95%	0.0	96.06%	77991103	NC_028641.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	5786	42972	89%	0.0	95.92%	59803892	NC_028638.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	5496	62803	94%	0.0	94.76%	65633393	NC_015447.3
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	5276	30826	94%	0.0	92.34%	66557038	NC_015441.3
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00333	Solanum tuberosum	5228	8357	84%	0.0	88.93%	657172	NW_006239270.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00532	Solanum tuberosum	5018	13874	90%	0.0	91.54%	325781	NW_006239461.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	3712	32261	92%	0.0	84.19%	60730942	NC_028642.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3046	34876	82%	0.0	80.81%	163962470	NC_029978.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2970	43045	80%	0.0	79.91%	301019445	NC_029977.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00558	Solanum tuberosum	2861	7368	84%	0.0	86.52%	310331	NW_006239478.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2617	30681	83%	0.0	77.51%	219521584	NC_029982.1
piaSL_monotypic Ch12_2s	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	1351	16504	88%	0.0	80.64%	215701946	NC_029980.1
CopiaSL_26	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	9799	56832	100%	0.0	100.00%	66723567	NC_015442.3
CopiaSL_26	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum pennellii	8626	62076	100%	0.0	96.00%	77991103	NC_028641.1
CopiaSL_26	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	8427	1,05E+08	100%	0.0	95.25%	109333515	NC_028637.1
CopiaSL_26	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	Solanum pennellii	8416	53457	100%	0.0	95.20%	84057508	NC_028645.1
CopiaSL_26	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	Solanum pennellii	8399	19373	100%	0.0	95.17%	70546659	NC_028644.1
CopiaSL_26	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	8181	35698	100%	0.0	94.44%	55977580	NC_015439.3
CopiaSL_26	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	8054	68627	100%	0.0	94.02%	65633393	NC_015447.3
CopiaSL_26	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	8000	26429	100%	0.0	93.82%	56597135	NC_015448.3
CopiaSL_26	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	8000	47777	100%	0.0	93.86%	68126176	NC_015449.3
CopiaSL_26	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	7989	41095	100%	0.0	93.82%	75414019	NC_028639.1
CopiaSL_26	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	Solanum pennellii	7974	32109	100%	0.0	93.73%	66223686	NC_028647.1
CopiaSL_26	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum pennellii	7899	30383	100%	0.0	93.48%	82529941	NC_028646.1
CopiaSL_26	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	7851	42304	100%	0.0	93.37%	72290146	NC_015440.3
CopiaSL_26	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	Solanum pennellii	7788	44419	100%	0.0	93.12%	79292169	NC_028643.1
CopiaSL_26	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	7618	23636	100%	0.0	92.77%	65987440	NC_015445.3
CopiaSL_26	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	7430	44726	100%	0.0	91.93%	77197300	NC_028640.1
CopiaSL_26	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00147	Solanum tuberosum	7136	7669	100%	0.0	90.91%	1440303	NW_006239070.1
CopiaSL_26	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	Solanum pennellii	7129	31770	100%	0.0	94.97%	83305730	NC_028648.1
CopiaSL_26	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00161	Solanum tuberosum	6828	7532	100%	0.0	89.87%	1344915	NW_006239086.1
CopiaSL_26	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00046	Solanum tuberosum	6663	8004	99%	0.0	91.07%	2411080	NW_006238971.1

CopiaSL_26	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00493	Solanum tuberosum	6582	7705	100%	0.0	90.81%	370066	NW_006239430.1
CopiaSL_26	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00281	Solanum tuberosum	6577	7655	100%	0.0	90.48%	861293	NW_006239206.1
CopiaSL_26	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	6453	39232	100%	0.0	89.31%	68175699	NC_015444.3
CopiaSL_26	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	6362	28795	100%	0.0	88.42%	72906345	NC_015446.3
CopiaSL_26	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	6215	38542	100%	0.0	92.08%	98455869	NC_015438.3
CopiaSL_26	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	6085	17848	100%	0.0	94.01%	59803892	NC_028638.1
CopiaSL_26	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00430	Solanum tuberosum	5891	7074	99%	0.0	88.95%	474478	NW_006239366.1
CopiaSL_26	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00115	Solanum tuberosum	5559	6273	100%	0.0	87.53%	1616611	NW_006239044.1
CopiaSL_26	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 contig00003	Solanum tuberosum	5044	7634	100%	0.0	90.12%	28034	NW_006239975.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	4857	61104	99%	0.0	84.97%	219521584	NC_029982.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	4848	89859	100%	0.0	84.91%	301019445	NC_029977.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	4820	65511	99%	0.0	84.79%	215701946	NC_029980.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	4780	36996	99%	0.0	84.63%	238794889	NC_029985.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	4745	57051	99%	0.0	84.66%	217274494	NC_029981.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	4743	87375	98%	0.0	84.51%	163962470	NC_029978.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	4706	80704	98%	0.0	84.60%	229934170	NC_029988.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	4698	76577	98%	0.0	84.52%	220335243	NC_029987.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	4673	96028	98%	0.0	84.28%	261510930	NC_029979.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold7059	Capsicum annuum	4673	5426	96%	0.0	84.41%	199068	NW_015961188.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2400	Capsicum annuum	4662	5044	96%	0.0	84.34%	1353284	NW_015960968.1
CopiaSL_26	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00001	Solanum tuberosum	4604	4833	81%	0.0	86.78%	7090834	NW_006238927.1
CopiaSL_26	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	4577	6228	98%	0.0	86.39%	60730942	NC_028642.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	4508	75761	98%	0.0	84.75%	205736368	NC_029986.1
CopiaSL_26	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00073	Solanum tuberosum	4490	7666	99%	0.0	90.86%	2002897	NW_006239002.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	4425	20392	98%	0.0	83.36%	153299543	NC_029984.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	4322	53132	100%	0.0	83.56%	222112641	NC_029983.1
CopiaSL_26	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	4185	39135	100%	0.0	87.54%	66557038	NC_015441.3
CopiaSL_26	refseq_genomes	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5971	Capsicum annum	4109	5261	96%	0.0	84.01%	156035	NW_015961177.1
CopiaSL_26	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00667	Solanum tuberosum	4089	7360	100%	0.0	90.69%	208703	NW_006239588.1
CopiaSL_26	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	3760	5944	99%	0.0	85.15%	49794276	NC_015443.3
CopiaSL_26	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00279	Solanum tuberosum	3714	7895	100%	0.0	89.89%	864930	NW_006239204.1
CopiaSL_26	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00060	Solanum tuberosum	3611	7272	98%	0.0	90.28%	2184951	NW_006238985.1
CopiaSL_26	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3116	4442	86%	0.0	83.70%	468561	NW_017670746.1
CopiaSL_26	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 12, NIATTR2	Nicotiana attenuata	3038	4125	84%	0.0	84.48%	53491057	NC_032000.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold5309	Capsicum annum	3007	5029	92%	0.0	84.06%	181449	NW_015960649.1
CopiaSL_26	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold62305	Nicotiana tabacum	2933	3951	86%	0.0	81.66%	97576	NW_015923851.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold561	Capsicum annum	2728	4835	99%	0.0	84.55%	1145873	NW_015961012.1
CopiaSL_26	refseq_genomes	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold3904	Capsicum annum	2686	5142	96%	0.0	85.48%	169740	NW_015960928.1
CopiaSL_26	refseq_genomes	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1894	Capsicum annum	2580	5056	98%	0.0	83.74%	2334823	NW_015961490.1
CopiaSL_26	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 11, NIATTR2	Nicotiana attenuata	2350	4278	83%	0.0	82.75%	61252062	NC_031999.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold572	Capsicum annum	2183	4788	92%	0.0	85.51%	590163	NW_015960560.1
CopiaSL_26	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold24535	Nicotiana tomentosiformis	2124	4510	85%	0.0	84.51%	79504	NW_008904192.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold170	Capsicum annum	2076	5813	82%	0.0	81.77%	2217302	NW_015960390.1
CopiaSL_26	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 5, NIATTR2	Nicotiana attenuata	2019	4386	86%	0.0	84.42%	57012777	NC_031993.1
CopiaSL_26	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold26753	Nicotiana sylvestris	1720	4246	86%	0.0	83.72%	138211	NW_009511335.1
CopiaSL_26	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	1681	4303	83%	0.0	84.41%	923158	NW_017670348.1
CopiaSL_26	refseq_genomes	8	Capsicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2138	Capsicum annum	1592	5889	98%	0.0	84.41%	681955	NW_015960370.1
CopiaSL_26	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold979	Capsicum annum	1471	4751	90%	0.0	84.63%	1172749	NW_015961157.1
CopiaSL_28	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	11500	4,23E+08	100%	0.0	100.00%	65633393	NC_015447.3
CopiaSL_28	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum pennellii Solanum	10436	4,65E+08	99%	0.0	96.95%	82529941	NC_028646.1
CopiaSL_28	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	pennellii Solanum pennellii	10405	2,65E+08	100%	0.0	96.91%	70546659	NC_028644.1
CopiaSL_28	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii Solanum	10390	2,05E+08	99%	0.0	96.82%	60730942	NC_028642.1
CopiaSL_28	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	lycopersicum Solanum	10312	2,20E+08	99%	0.0	96.65%	66223686	NC_028647.1
CopiaSL_28	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	lycopersicum Solanum	10179	5,94E+08	100%	0.0	96.31%	98455869	NC_015438.3
CopiaSL_28	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	pennellii	10135	1,82E+08	99%	0.0	96.13%	55977580	NC_015439.3
CopiaSL_28	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	10131	2,68E+08	99%	0.0	96.85%	84057508	NC_028645.1	

CopiaSL_28	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum pennellii	10122	3,26E+08	99%	0.0	96.83%	77991103	NC_028641.1
CopiaSL_28	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	Solanum pennellii	10078	2,28E+08	99%	0.0	96.77%	79292169	NC_028643.1
CopiaSL_28	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	10070	3,30E+08	99%	0.0	96.00%	66723567	NC_015442.3
CopiaSL_28	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	9614	3,21E+08	100%	0.0	96.85%	68126176	NC_015449.3
CopiaSL_28	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	9557	1,82E+08	99%	0.0	97.37%	75414019	NC_028639.1
CopiaSL_28	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	9369	3,32E+08	99%	0.0	96.21%	72906345	NC_015446.3
CopiaSL_28	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	9358	5,98E+08	100%	0.0	97.67%	109333515	NC_028637.1
CopiaSL_28	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	9354	4,43E+08	99%	0.0	97.39%	77197300	NC_028640.1
CopiaSL_28	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	9348	1,58E+08	99%	0.0	97.37%	59803892	NC_028638.1
CopiaSL_28	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	Solanum pennellii	9341	3,36E+08	99%	0.0	97.33%	83305730	NC_028648.1
CopiaSL_28	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	9136	3,84E+08	100%	0.0	96.67%	66557038	NC_015441.3
CopiaSL_28	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	8881	2,55E+08	100%	0.0	95.85%	65987440	NC_015445.3
CopiaSL_28	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	8841	1,68E+08	100%	0.0	95.86%	49794276	NC_015443.3
CopiaSL_28	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	8665	2,48E+08	100%	0.0	95.24%	56597135	NC_015448.3
CopiaSL_28	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	8510	2,13E+08	99%	0.0	95.49%	68175699	NC_015444.3
CopiaSL_28	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	8335	2,42E+08	100%	0.0	94.19%	72290146	NC_015440.3
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00001	Solanum tuberosum	6599	10413	91%	0.0	90.10%	7090834	NW_006238927.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00226	Solanum tuberosum	6462	15787	94%	0.0	88.45%	1057645	NW_006239155.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00058	Solanum tuberosum	6423	7605	87%	0.0	91.43%	2204211	NW_006238982.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00056	Solanum tuberosum	6412	16628	92%	0.0	90.21%	2177853	NW_006238986.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00764	Solanum tuberosum	6405	7347	85%	0.0	89.63%	123992	NW_006239690.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00809	Solanum tuberosum	6192	7048	91%	0.0	89.40%	97311	NW_006239738.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00145	Solanum tuberosum	6168	14112	91%	0.0	87.88%	1445052	NW_006239069.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00035	Solanum tuberosum	6154	10400	91%	0.0	89.37%	2572343	NW_006238963.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00249	Solanum tuberosum	6109	10961	85%	0.0	87.71%	921806	NW_006239187.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00271	Solanum tuberosum	6039	6144	86%	0.0	87.08%	902958	NW_006239192.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00691	Solanum tuberosum	5757	8154	85%	0.0	89.83%	195803	NW_006239605.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00316	Solanum tuberosum	5629	15490	92%	0.0	88.54%	755641	NW_006239240.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00647	Solanum tuberosum	5607	8817	84%	0.0	89.47%	219899	NW_006239575.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00225	Solanum tuberosum	5489	7214	92%	0.0	89.41%	1071188	NW_006239149.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00112	Solanum tuberosum	5339	8069	91%	0.0	89.53%	1674359	NW_006239035.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00013	Solanum tuberosum	5286	15523	92%	0.0	90.86%	3422333	NW_006238940.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00369	Solanum tuberosum	5068	6582	86%	0.0	89.02%	591576	NW_006239297.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00606	Solanum tuberosum	4868	7114	92%	0.0	88.51%	265855	NW_006239521.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00097	Solanum tuberosum	4745	9456	86%	0.0	87.00%	1770913	NW_006239022.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf01144	Solanum tuberosum	4658	6502	86%	0.0	88.02%	19337	NW_006240095.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00121	Solanum tuberosum	4599	7231	88%	0.0	90.27%	1607711	NW_006239046.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00320	Solanum tuberosum	4584	6265	80%	0.0	90.50%	738063	NW_006239242.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00786	Solanum tuberosum	4433	11300	84%	0.0	86.08%	123712	NW_006239691.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00037	Solanum tuberosum	4409	8643	80%	0.0	84.52%	2613480	NW_006238961.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00029	Solanum tuberosum	4324	8144	84%	0.0	85.68%	2785282	NW_006238955.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00403	Solanum tuberosum	4307	6293	84%	0.0	90.00%	537389	NW_006239321.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00028	Solanum tuberosum	4265	12014	91%	0.0	83.36%	2805095	NW_006238954.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00016	Solanum tuberosum	4261	21191	92%	0.0	88.44%	3295453	NW_006238942.1
CopiaSL_28	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.0 SCF00000350	Solanum lycopersicum	4095	7476	83%	0.0	94.14%	5101	NW_020442831.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00044	Solanum tuberosum	4061	8640	90%	0.0	89.50%	2434495	NW_006238970.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00204	Solanum tuberosum	4058	8236	89%	0.0	91.20%	1149434	NW_006239127.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00049	Solanum tuberosum	4013	11715	89%	0.0	90.44%	2333911	NW_006238974.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00213	Solanum tuberosum	3916	7218	85%	0.0	88.47%	1095126	NW_006239141.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00194	Solanum tuberosum	3895	6447	89%	0.0	89.20%	1175979	NW_006239120.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00420	Solanum tuberosum	3890	21160	93%	0.0	89.41%	512266	NW_006239340.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00344	Solanum tuberosum	3877	6581	82%	0.0	91.07%	652076	NW_006239273.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00254	Solanum tuberosum	3685	7292	86%	0.0	89.15%	968134	NW_006239174.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00151	Solanum tuberosum	3511	7886	84%	0.0	88.42%	1427473	NW_006239077.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00255	Solanum tuberosum	3498	7558	90%	0.0	89.66%	929374	NW_006239184.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00813	Solanum tuberosum	3413	6732	84%	0.0	90.41%	102418	NW_006239727.1

CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00206	Solanum tuberosum	3386	8567	81%	0.0	89.63%	1138255	NW_006239131.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00728	Solanum tuberosum	3374	7264	93%	0.0	88.37%	156345	NW_006239651.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00917	Solanum tuberosum	3343	6078	83%	0.0	87.80%	59371	NW_006239814.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00192	Solanum tuberosum	3251	9459	83%	0.0	89.45%	1185169	NW_006239151.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00464	Solanum tuberosum	3090	5847	84%	0.0	88.77%	411434	NW_006239400.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00027	Solanum tuberosum	3046	14132	94%	0.0	89.66%	2869105	NW_006238953.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00490	Solanum tuberosum	3022	6563	85%	0.0	91.15%	394222	NW_006239413.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00089	Solanum tuberosum	2988	7727	81%	0.0	87.76%	1856049	NW_006239012.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00011	Solanum tuberosum	2902	10122	85%	0.0	85.27%	3646191	NW_006238938.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00348	Solanum tuberosum	2891	6136	84%	0.0	90.70%	629130	NW_006239285.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00125	Solanum tuberosum	2826	11863	92%	0.0	89.63%	1553111	NW_006239052.1
CopiaSL_28	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00018	Solanum tuberosum	2817	14894	91%	0.0	90.52%	3280163	NW_006238944.1
CopiaSL_28	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2558	3,47E+08	80%	0.0	79.96%	215701946	NC_029980.1
CopiaSL_28	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2549	3,59E+08	81%	0.0	79.04%	205736368	NC_029986.1
CopiaSL_25	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	9808	92390	100%	0.0	100.00%	66557038	NC_015441.3
CopiaSL_25	refseq_genomes	8	Solanum pennellii chromosome 8, SPENNv200	Solanum pennellii	5553	79129	100%	0.0	96.26%	70546659	NC_028644.1
CopiaSL_25	refseq_genomes	8	Solanum pennellii chromosome 4, SPENNv200	Solanum pennellii	5522	1,33E+08	100%	0.0	96.11%	77197300	NC_028640.1
CopiaSL_25	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	5363	84239	100%	0.0	96.41%	65633393	NC_015447.3
CopiaSL_25	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	5275	53479	100%	0.0	95.92%	49794276	NC_015443.3
CopiaSL_25	refseq_genomes	8	Solanum pennellii chromosome 10, SPENNv200	Solanum pennellii	5129	1,34E+08	98%	0.0	95.72%	82529941	NC_028646.1
CopiaSL_25	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	5120	93616	100%	0.0	94.06%	68175699	NC_015444.3
CopiaSL_25	refseq_genomes	8	Solanum pennellii chromosome 7, SPENNv200	Solanum pennellii	5110	1,16E+08	99%	0.0	95.68%	79292169	NC_028643.1
CopiaSL_25	refseq_genomes	8	Solanum pennellii chromosome 6, SPENNv200	Solanum pennellii	5077	93603	100%	0.0	96.32%	60730942	NC_028642.1
CopiaSL_25	refseq_genomes	8	Solanum pennellii chromosome 9, SPENNv200	Solanum pennellii	5077	1,15E+08	98%	0.0	96.32%	84057508	NC_028645.1
CopiaSL_25	refseq_genomes	8	Solanum pennellii chromosome 1, SPENNv200	Solanum pennellii	5072	52617	100%	0.0	96.29%	109333515	NC_028637.1
CopiaSL_25	refseq_genomes	8	Solanum pennellii chromosome 2, SPENNv200	Solanum pennellii	5072	78327	100%	0.0	96.29%	59803892	NC_028638.1
CopiaSL_25	refseq_genomes	8	Solanum pennellii chromosome 11, SPENNv200	Solanum pennellii	5072	97647	96%	0.0	96.29%	66223686	NC_028647.1
CopiaSL_25	refseq_genomes	8	Solanum pennellii chromosome 12, SPENNv200	Solanum pennellii	5072	54844	94%	0.0	96.29%	83305730	NC_028648.1
CopiaSL_25	refseq_genomes	8	Solanum pennellii chromosome 3, SPENNv200	Solanum pennellii	5060	91536	91%	0.0	96.22%	75414019	NC_028639.1
CopiaSL_25	refseq_genomes	8	Solanum pennellii chromosome 5, SPENNv200	Solanum pennellii	5060	1,17E+08	99%	0.0	96.22%	77991103	NC_028641.1
CopiaSL_25	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	5003	48948	100%	0.0	95.17%	55977580	NC_015439.3
CopiaSL_25	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	4916	56342	98%	0.0	92.93%	68126176	NC_015449.3
CopiaSL_25	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	4822	51534	99%	0.0	94.90%	65987440	NC_015445.3
CopiaSL_25	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	4542	32698	93%	0.0	93.35%	72906345	NC_015446.3
CopiaSL_25	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	4516	64282	94%	0.0	93.20%	72290146	NC_015440.3
CopiaSL_25	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	4490	38497	91%	0.0	92.95%	66723567	NC_015442.3
CopiaSL_25	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	4490	50208	98%	0.0	93.05%	56597135	NC_015448.3
CopiaSL_25	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00145	Solanum tuberosum	4294	6381	85%	0.0	91.74%	1445052	NW_006239069.1
CopiaSL_25	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00237	Solanum tuberosum	4228	6554	89%	0.0	91.38%	1028729	NW_006239161.1
CopiaSL_25	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00070	Solanum tuberosum	4115	6216	90%	0.0	90.80%	2061072	NW_00623896.1
CopiaSL_25	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00085	Solanum tuberosum	4019	9442	90%	0.0	90.10%	1839851	NW_006239014.1
CopiaSL_25	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00134	Solanum tuberosum	3818	4684	85%	0.0	89.07%	1492581	NW_006239062.1
CopiaSL_25	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00019	Solanum tuberosum	3814	4751	90%	0.0	88.83%	3257788	NW_006238945.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold70723	Nicotiana tabacum	3472	3472	90%	0.0	80.02%	27986	NW_015931208.1
CopiaSL_25	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00047	Solanum tuberosum	3149	5399	84%	0.0	90.85%	2402127	NW_006238973.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	3116	3985	89%	0.0	84.94%	49050	NW_017676244.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold10379	Nicotiana tomentosiformis	3116	4288	89%	0.0	84.93%	196381	NW_008832704.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31284	Nicotiana tomentosiformis	3116	4304	89%	0.0	84.94%	143321	NW_008911691.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4404	Nicotiana tomentosiformis	3116	6899	89%	0.0	84.93%	98393	NW_008925864.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold88	Nicotiana tomentosiformis	3116	5044	89%	0.0	84.94%	243129	NW_008974706.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3309	Nicotiana tomentosiformis	3094	4299	89%	0.0	84.81%	150992	NW_008913697.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold106	Nicotiana tomentosiformis	3072	4253	89%	0.0	84.68%	142834	NW_008835159.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2342	Nicotiana tomentosiformis	3066	4147	88%	0.0	84.64%	164189	NW_008902953.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold48476	Nicotiana tabacum	3061	4142	89%	0.0	84.65%	49177	NW_015911352.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold11948	Nicotiana tomentosiformis	3061	4171	89%	0.0	84.62%	83338	NW_008850135.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	3055	4216	89%	0.0	84.60%	82204	NW_017674872.1

CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24146	Nicotiana tabacum	3055	4214	89%	0.0	84.62%	56338	NW_015871540.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold28586	Nicotiana tabacum	3055	5354	89%	0.0	84.62%	57113	NW_015886712.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold42487	Nicotiana tabacum	3049	4214	89%	0.0	84.58%	67139	NW_015905834.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold64849	Nicotiana tabacum	3049	4236	89%	0.0	84.58%	55060	NW_015926109.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold33870	Nicotiana sylvestris	3049	6225	89%	0.0	84.58%	118539	NW_009519243.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold72853	Nicotiana tabacum	3038	4164	89%	0.0	84.52%	52624	NW_015933057.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1516	Nicotiana sylvestris	3038	4192	89%	0.0	84.52%	73449	NW_009396136.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold22194	Nicotiana sylvestris	3027	4208	89%	0.0	84.45%	76251	NW_009474292.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 2, NIATTr2	Nicotiana attenuata	3011	51770	89%	0.0	84.34%	96359640	NC_031990.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold9935	Nicotiana sylvestris	3011	4153	89%	0.0	84.36%	395863	NW_009519971.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3009	4168	89%	0.0	84.36%	227599	NW_017671979.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold14908	Nicotiana tabacum	2996	4138	89%	0.0	84.27%	35405	NW_015822307.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7775	Nicotiana tabacum	2985	4802	89%	0.0	84.21%	233187	NW_015937251.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 11, NIATTr2	Nicotiana attenuata	2953	31716	89%	0.0	84.01%	61252062	NC_031999.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 7, NIATTr2	Nicotiana attenuata	2950	72973	90%	0.0	84.01%	114765206	NC_031995.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold10746	Nicotiana sylvestris	2950	4040	89%	0.0	84.02%	130135	NW_009347092.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 10, NIATTr2	Nicotiana attenuata	2944	27903	89%	0.0	83.99%	51082206	NC_031998.1
CopiaSL_25	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2944	87035	91%	0.0	83.91%	217274494	NC_029981.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	2924	4050	89%	0.0	83.82%	285848	NW_017671514.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold11906	Nicotiana tomentosiformis	2913	3915	89%	0.0	83.76%	115774	NW_008849669.1
CopiaSL_25	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2909	80238	89%	0.0	83.49%	205736368	NC_029986.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold73374	Nicotiana tabacum	2907	3992	89%	0.0	83.75%	32556	NW_015933483.1
CopiaSL_25	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2907	99692	91%	0.0	83.51%	261510930	NC_029979.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9941	Nicotiana tabacum	2902	3752	85%	0.0	83.69%	152282	NW_015955006.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13868	Nicotiana tabacum	2889	3948	89%	0.0	83.63%	134467	NW_015815476.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 3, NIATTr2	Nicotiana attenuata	2885	45894	90%	0.0	83.63%	72893213	NC_031991.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	2885	3942	89%	0.0	83.66%	200864	NW_017672311.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold65735	Nicotiana tabacum	2883	4002	89%	0.0	83.61%	96160	NW_015926868.1
CopiaSL_25	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2883	1,03E+08	89%	0.0	83.36%	219521584	NC_029982.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold60862	Nicotiana sylvestris	2880	3889	87%	0.0	83.59%	58890	NW_009549234.1
CopiaSL_25	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2874	65562	89%	0.0	83.54%	163962470	NC_029978.1
CopiaSL_25	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2874	45252	89%	0.0	83.52%	153299543	NC_029984.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold55444	Nicotiana tabacum	2861	3944	89%	0.0	83.47%	179457	NW_015917721.1
CopiaSL_25	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 S16 R44 unplaced genomic scaffold, SolTub_3.0 scf0043	Solanum tuberosum	2850	15388	90%	0.0	88.87%	2504196	NW_006238969.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold11111	Nicotiana tabacum	2846	3872	89%	0.0	83.40%	110871	NW_015795774.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 12, NIATTr2	Nicotiana attenuata	2837	41061	89%	0.0	83.34%	53491057	NC_032000.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold106431	Nicotiana tabacum	2837	3882	89%	0.0	83.31%	17027	NW_015792222.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 9, NIATTr2	Nicotiana attenuata	2835	33688	87%	0.0	83.82%	50975570	NC_031997.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23012	Nicotiana tabacum	2835	4294	89%	0.0	83.31%	95574	NW_015866247.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold24539	Nicotiana sylvestris	2828	10002	89%	0.0	83.31%	26440	NW_009500347.1
CopiaSL_25	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 S16 R44 unplaced genomic scaffold, SolTub_3.0 scf00464	Solanum tuberosum	2819	5673	81%	0.0	90.05%	411434	NW_006239400.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold64820	Nicotiana tabacum	2806	3887	89%	0.0	83.18%	48718	NW_015926085.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	2800	3819	89%	0.0	83.16%	582787	NW_017670557.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold25062	Nicotiana tabacum	2800	3800	89%	0.0	83.15%	239346	NW_015875667.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold46775	Nicotiana tabacum	2798	3893	89%	0.0	83.38%	57263	NW_015909796.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold20712	Nicotiana tabacum	2796	3848	89%	0.0	83.15%	48515	NW_015855243.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold29311	Nicotiana tabacum	2795	3789	89%	0.0	83.10%	65512	NW_015887397.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold48899	Nicotiana tomentosiformis	2795	3518	83%	0.0	83.11%	18964	NW_008931261.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold65832	Nicotiana tabacum	2793	3786	89%	0.0	83.17%	15390	NW_015926954.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold50892	Nicotiana tabacum	2789	3800	89%	0.0	83.05%	134077	NW_015913547.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold25636	Nicotiana tabacum	2785	3835	89%	0.0	83.04%	68990	NW_015878252.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold66941	Nicotiana tabacum	2785	7149	88%	0.0	83.02%	139198	NW_015927940.1
CopiaSL_25	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2782	51870	89%	0.0	81.25%	222112641	NC_029983.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5002	Nicotiana tomentosiformis	2782	3892	89%	0.0	83.07%	165988	NW_008932509.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold65241	Nicotiana tabacum	2769	3840	89%	0.0	82.95%	16166	NW_015926443.1
CopiaSL_25	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2767	71280	89%	0.0	82.64%	220335243	NC_029987.1

CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold32528	Nicotiana tabacum	2763	3730	89%	0.0	82.99%	28847	NW_015890430.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2482	Nicotiana tomentosiformis	2761	3501	83%	0.0	82.90%	83457	NW_008904508.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold57969	Nicotiana tabacum	2760	3797	89%	0.0	82.93%	109786	NW_015920005.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsy Nsy_scaffold5473	Nicotiana sylvestris	2760	3721	89%	0.0	82.97%	95057	NW_009542420.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold20975	Nicotiana tabacum	2758	3780	89%	0.0	82.87%	236010	NW_015856563.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold54415	Nicotiana tabacum	2758	3802	89%	0.0	82.89%	129201	NW_015916796.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsy Nsy_scaffold2829	Nicotiana sylvestris	2758	3686	89%	0.0	82.93%	139875	NW_009513042.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold7530	Nicotiana tomentosiformis	2752	3708	89%	0.0	82.84%	42054	NW_008960597.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold33339	Nicotiana tabacum	2750	3735	88%	0.0	82.81%	92052	NW_015891198.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9134	Nicotiana tabacum	2748	3710	89%	0.0	82.85%	36348	NW_015948588.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4569	Nicotiana tabacum	2747	5935	88%	0.0	82.82%	31641	NW_015908778.1
CopiaSL_25	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2747	94168	89%	0.0	82.58%	229934170	NC_029988.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2948	Nicotiana tomentosiformis	2745	3757	89%	0.0	82.81%	109912	NW_008909685.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold103698	Nicotiana tabacum	2741	3762	89%	0.0	82.80%	48625	NW_015790111.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold29400	Nicotiana tabacum	2741	3721	88%	0.0	82.81%	57393	NW_015887481.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold44197	Nicotiana tabacum	2741	3763	83%	0.0	82.54%	67946	NW_015907412.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4038	Nicotiana tomentosiformis	2741	3756	89%	0.0	82.78%	81983	NW_008921797.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold2523	Nicotiana tabacum	2739	3800	89%	0.0	82.75%	77170	NW_015876407.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8171	Nicotiana tabacum	2739	6227	89%	0.0	82.80%	37086	NW_015940618.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold12490	Nicotiana tomentosiformis	2739	3789	89%	0.0	82.75%	290929	NW_008856158.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5562	Nicotiana tabacum	2737	3749	89%	0.0	82.97%	72894	NW_015917874.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5911	Nicotiana tomentosiformis	2737	3683	87%	0.0	82.81%	517654	NW_008942608.1
CopiaSL_25	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2736	1,10E+08	91%	0.0	82.50%	301019445	NC_029977.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4300	Nicotiana tomentosiformis	2736	3780	89%	0.0	82.79%	344126	NW_008924709.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold15486	Nicotiana tabacum	2734	3810	89%	0.0	82.75%	134618	NW_015825994.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17203	Nicotiana tabacum	2734	4128	89%	0.0	82.81%	234505	NW_015836299.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold28928	Nicotiana tabacum	2734	3706	89%	0.0	82.74%	76998	NW_015887028.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsy Nsy_scaffold53930	Nicotiana sylvestris	2734	3719	89%	0.0	82.73%	148314	NW_009541532.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsy Nsy_scaffold1062	Nicotiana sylvestris	2732	3719	89%	0.0	82.76%	11693	NW_009345692.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold131970	Nicotiana tabacum	2728	3752	89%	0.0	82.73%	33367	NW_015810872.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold19802	Nicotiana tabacum	2728	3708	89%	0.0	82.68%	95352	NW_015850591.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2r	Nicotiana attenuata	2726	5966	90%	0.0	82.70%	1243377	NW_017670288.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold33105	Nicotiana tabacum	2726	3677	89%	0.0	82.71%	20275	NW_015890971.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold15762	Nicotiana tomentosiformis	2726	3704	89%	0.0	82.79%	189671	NW_008892513.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold38522	Nicotiana tomentosiformis	2726	5418	86%	0.0	83.20%	50504	NW_008919733.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5574	Nicotiana tomentosiformis	2724	6233	89%	0.0	82.73%	158996	NW_008938863.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold14556	Nicotiana tabacum	2723	3767	89%	0.0	82.68%	60873	NW_015820059.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold32955	Nicotiana tabacum	2723	3743	89%	0.0	82.66%	22770	NW_015890829.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold50145	Nicotiana tabacum	2723	3695	87%	0.0	82.68%	14795	NW_015912877.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold484	Nicotiana tomentosiformis	2723	5175	89%	0.0	82.66%	191032	NW_008930707.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5681	Nicotiana tomentosiformis	2723	3488	83%	0.0	82.73%	47892	NW_008940052.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold67005	Nicotiana tabacum	2721	7024	89%	0.0	82.75%	83939	NW_015927993.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold15158	Nicotiana tomentosiformis	2721	3686	89%	0.0	82.70%	127001	NW_008885802.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold31865	Nicotiana tabacum	2719	3815	89%	0.0	82.74%	67832	NW_015889803.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold96322	Nicotiana tabacum	2719	3544	83%	0.0	82.66%	62139	NW_015952568.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold97198	Nicotiana tabacum	2719	4839	89%	0.0	82.65%	13841	NW_015953256.1
CopiaSL_25	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2719	53374	89%	0.0	82.67%	215701946	NC_029980.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold36321	Nicotiana tomentosiformis	2719	3752	89%	0.0	82.68%	58602	NW_008917288.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold11177	Nicotiana tomentosiformis	2717	4821	84%	0.0	82.65%	229492	NW_008841569.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold41874	Nicotiana tabacum	2713	3656	89%	0.0	82.72%	72296	NW_015905273.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold39287	Nicotiana tabacum	2712	3756	89%	0.0	82.63%	156027	NW_015902877.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold30867	Nicotiana tomentosiformis	2706	3781	87%	0.0	82.58%	49852	NW_008911227.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold34992	Nicotiana tabacum	2704	3749	89%	0.0	82.59%	14880	NW_015897035.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold46735	Nicotiana tabacum	2704	3750	89%	0.0	82.65%	49813	NW_015909760.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4179	Nicotiana tomentosiformis	2704	4693	89%	0.0	82.72%	227617	NW_008923363.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23443	Nicotiana tabacum	2702	3507	83%	0.0	82.63%	55046	NW_015868252.1

CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold28405	Nicotiana tomentosiformis	2702	3739	89%	0.0	82.58%	71370	NW_008908492.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold88976	Nicotiana tabacum	2700	3738	89%	0.0	82.57%	65370	NW_015946636.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold14605	Nicotiana tomentosiformis	2700	4920	89%	0.0	82.56%	23341	NW_008879658.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4056	Nicotiana tabacum	2699	8499	89%	0.0	82.54%	201862	NW_015904038.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold32585	Nicotiana tabacum	2697	3776	89%	0.0	82.53%	78462	NW_015890481.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold17112	Nicotiana tomentosiformis	2697	3472	84%	0.0	82.57%	79869	NW_008895945.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 5, NIATTr2	Nicotiana attenuata	2695	22883	90%	0.0	82.66%	57012777	NC_031993.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1098	Nicotiana tabacum	2691	3664	89%	0.0	82.65%	341244	NW_015794785.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold26885	Nicotiana tomentosiformis	2691	3760	89%	0.0	82.49%	73620	NW_008906802.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold63326	Nicotiana tabacum	2689	3807	85%	0.0	82.50%	48267	NW_015924759.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1641	Nicotiana tomentosiformis	2689	3695	89%	0.0	82.52%	133998	NW_008895164.1
CopiaSL_25	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold181	Capsicum annuum	2687	8606	80%	0.0	82.30%	2703691	NW_015961388.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold32027	Nicotiana tabacum	2686	3695	89%	0.0	82.53%	17155	NW_015889961.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold49451	Nicotiana tabacum	2686	3645	88%	0.0	82.46%	49723	NW_015912231.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold27863	Nicotiana tabacum	2684	3429	87%	0.0	82.59%	163106	NW_015886036.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17044	Nicotiana tabacum	2682	3693	89%	0.0	82.53%	35129	NW_015835420.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold39463	Nicotiana tabacum	2680	3634	89%	0.0	82.44%	71785	NW_015903036.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9508	Nicotiana tabacum	2680	5450	89%	0.0	82.47%	212694	NW_015951581.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold8429	Nicotiana tomentosiformis	2680	3601	89%	0.0	82.52%	221969	NW_008970585.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold57213	Nicotiana tabacum	2678	3721	89%	0.0	82.43%	116352	NW_015919314.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold21398	Nicotiana tomentosiformis	2678	3634	89%	0.0	82.59%	219202	NW_008900706.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold45287	Nicotiana tabacum	2675	3593	86%	0.0	82.46%	93217	NW_015908404.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold14755	Nicotiana tomentosiformis	2675	3632	89%	0.0	82.41%	39171	NW_008881324.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold16975	Nicotiana tomentosiformis	2675	4648	86%	0.0	82.43%	55487	NW_008895791.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold37122	Nicotiana tomentosiformis	2675	5960	83%	0.0	82.49%	106680	NW_008918178.1
CopiaSL_25	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00662	Solanum tuberosum	2671	5393	83%	0.0	88.82%	215247	NW_006239580.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24231	Nicotiana tabacum	2669	4656	89%	0.0	82.38%	59941	NW_015871924.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2586	Nicotiana tomentosiformis	2669	3683	87%	0.0	82.56%	125635	NW_008905663.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold6496	Nicotiana tomentosiformis	2667	3746	87%	0.0	82.89%	62680	NW_008949107.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold51911	Nicotiana tabacum	2665	3697	89%	0.0	82.36%	89518	NW_015914483.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4845	Nicotiana tabacum	2662	4656	89%	0.0	83.44%	139016	NW_015911334.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold47903	Nicotiana tabacum	2660	4230	82%	0.0	82.39%	74209	NW_015910829.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold12231	Nicotiana tabacum	2658	3918	89%	0.0	82.31%	135288	NW_015804005.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1496	Nicotiana tabacum	2658	3597	89%	0.0	82.35%	99225	NW_015822620.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1874	Nicotiana tabacum	2656	6635	89%	0.0	82.55%	191802	NW_015844914.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold26138	Nicotiana tabacum	2656	7122	89%	0.0	82.99%	354171	NW_015880641.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold48323	Nicotiana tabacum	2656	3705	86%	0.0	82.97%	44308	NW_015911222.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsy1 Nsy1_scaffold1902	Nicotiana sylvestris	2656	3529	89%	0.0	82.33%	146119	NW_009439025.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24394	Nicotiana tabacum	2654	6,21E+03	89%	0.0	82.36%	12611	NW_015872679.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold39882	Nicotiana tabacum	2654	3787	88%	0.0	82.52%	119430	NW_015903426.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold86782	Nicotiana tabacum	2652	3689	89%	0.0	82.42%	29332	NW_015944837.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold20354	Nicotiana tomentosiformis	2652	3693	89%	0.0	82.35%	55596	NW_008899547.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold87324	Nicotiana tabacum	2651	3,69E+03	89%	0.0	82.27%	145340	NW_015945281.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsy1 Nsy1_scaffold27623	Nicotiana sylvestris	2651	3731	89%	0.0	82.95%	79250	NW_009512302.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold21747	Nicotiana tomentosiformis	2651	3697	89%	0.0	82.26%	95370	NW_008910944.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2994	Nicotiana tomentosiformis	2651	3512	88%	0.0	82.26%	171488	NW_008910196.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold22424	Nicotiana tabacum	2649	3440	89%	0.0	82.26%	53421	NW_015863452.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold36028	Nicotiana tabacum	2645	3263	81%	0.0	82.27%	42750	NW_015899827.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold10746	Nicotiana tomentosiformis	2645	3608	88%	0.0	82.52%	283500	NW_008836782.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5092	Nicotiana tomentosiformis	2641	3886	89%	0.0	82.22%	124493	NW_008933508.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold8583	Nicotiana tomentosiformis	2639	3692	86%	0.0	82.23%	262523	NW_008972296.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold20575	Nicotiana tabacum	2638	3651	89%	0.0	82.27%	139348	NW_015854552.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold30929	Nicotiana tabacum	2638	4145	89%	0.0	83.28%	81297	NW_015888927.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5598	Nicotiana tabacum	2638	3695	89%	0.0	82.20%	135222	NW_015918199.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2844	Nicotiana tomentosiformis	2638	3629	87%	0.0	82.66%	243778	NW_008908530.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold36195	Nicotiana tabacum	2636	3656	89%	0.0	82.23%	20416	NW_015899981.1

CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold49421	Nicotiana tabacum	2636	3602	89%	0.0	82.32%	54256	NW_015912204.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3153	Nicotiana tomentosiformis	2636	3547	89%	0.0	82.24%	147072	NW_008911964.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold12780	Nicotiana tabacum	2628	3549	88%	0.0	82.43%	438606	NW_015807950.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold48631	Nicotiana tabacum	2628	3664	89%	0.0	82.11%	119175	NW_015911498.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold10274	Nicotiana sylvestris	2628	3676	80%	0.0	84.63%	151572	NW_009341848.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold7652	Nicotiana tomentosiformis	2628	3584	89%	0.0	82.18%	60547	NW_008961952.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold38670	Nicotiana tabacum	2625	3705	89%	0.0	82.19%	102374	NW_015902302.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold47106	Nicotiana tabacum	2625	3614	89%	0.0	82.12%	25338	NW_015910094.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold84869	Nicotiana tabacum	2625	3573	89%	0.0	82.30%	35756	NW_015943245.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2062	Nicotiana tomentosiformis	2625	5324	89%	0.0	82.08%	131621	NW_008899842.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1014	Nicotiana tabacum	2621	3610	89%	0.0	82.15%	57512	NW_015788330.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold14714	Nicotiana tomentosiformis	2621	3462	80%	0.0	82.84%	103166	NW_008880869.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold24921	Nicotiana tomentosiformis	2621	3697	89%	0.0	82.26%	19168	NW_008904621.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold64876	Nicotiana tabacum	2619	3701	89%	0.0	82.10%	124117	NW_015926130.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold20260	Nicotiana tomentosiformis	2619	3654	89%	0.0	82.16%	219114	NW_008899443.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5848	Nicotiana tabacum	2617	3672	87%	0.0	82.06%	217698	NW_015920454.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold65858	Nicotiana tabacum	2617	3665	89%	0.0	82.08%	84720	NW_015926977.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3240	Nicotiana tomentosiformis	2617	3736	82%	0.0	84.15%	51029	NW_008912931.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold7892	Nicotiana tomentosiformis	2612	3643	87%	0.0	82.03%	233327	NW_008964618.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23022	Nicotiana tabacum	2608	3595	89%	0.0	82.03%	66609	NW_015866296.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold11315	Nicotiana tomentosiformis	2608	3534	89%	0.0	82.09%	15646	NW_008843103.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4368	Nicotiana tomentosiformis	2608	3560	90%	0.0	81.91%	783164	NW_008925463.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold51939	Nicotiana tomentosiformis	2608	3625	88%	0.0	82.41%	47166	NW_008934640.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold91527	Nicotiana tabacum	2606	3697	89%	0.0	82.11%	51903	NW_015948725.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold102930	Nicotiana tabacum	2604	3588	87%	0.0	82.08%	31078	NW_015789523.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold14719	Nicotiana tabacum	2604	3488	84%	0.0	82.01%	144850	NW_015821120.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold63916	Nicotiana tabacum	2604	3566	89%	0.0	82.03%	10631	NW_015925297.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold49621	Nicotiana tabacum	2603	4184	89%	0.0	82.06%	143343	NW_015912393.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold29013	Nicotiana tomentosiformis	2603	3612	89%	0.0	81.99%	22434	NW_008909168.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 1, NIATT2r	Nicotiana attenuata	2601	50793	89%	0.0	82.01%	110538649	NC_031989.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold47586	Nicotiana tabacum	2601	4933	89%	0.0	82.19%	94921	NW_015910523.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold2857	Nicotiana sylvestris	2601	3568	88%	0.0	82.44%	87988	NW_009513353.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold9045	Nicotiana sylvestris	2601	3663	83%	0.0	82.17%	56057	NW_009582109.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5884	Nicotiana tomentosiformis	2599	3562	89%	0.0	81.95%	167785	NW_008942307.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold30389	Nicotiana tomentosiformis	2595	3644	86%	0.0	82.02%	30775	NW_008910696.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold35709	Nicotiana tomentosiformis	2593	3619	88%	0.0	82.26%	27468	NW_008916607.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold19916	Nicotiana tabacum	2591	3717	88%	0.0	82.22%	54742	NW_015851185.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold54310	Nicotiana tabacum	2591	7153	89%	0.0	82.05%	160905	NW_015916701.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold55149	Nicotiana sylvestris	2591	3485	84%	0.0	82.04%	33352	NW_009542886.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold2501	Nicotiana sylvestris	2588	3680	83%	0.0	83.80%	203304	NW_009505581.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold14989	Nicotiana tabacum	2586	3614	88%	0.0	82.28%	259993	NW_015822805.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 6, NIATT2r	Nicotiana attenuata	2584	51110	90%	0.0	81.93%	97571839	NC_031994.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold11265	Nicotiana tabacum	2584	3629	89%	0.0	81.86%	37943	NW_015796191.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold202736	Nicotiana tabacum	2584	3599	90%	0.0	81.92%	53152	NW_015852991.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold14506	Nicotiana tomentosiformis	2582	3562	89%	0.0	81.87%	76300	NW_008878558.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2r	Nicotiana attenuata	2579	3592	89%	0.0	81.97%	459022	NW_017670777.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold33845	Nicotiana tomentosiformis	2579	3621	89%	0.0	81.87%	55973	NW_008914536.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold77263	Nicotiana tabacum	2575	3510	87%	0.0	81.82%	17301	NW_015936817.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold22938	Nicotiana tomentosiformis	2575	6575	89%	0.0	81.95%	45536	NW_008902417.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold7276	Nicotiana sylvestris	2573	3538	89%	0.0	81.92%	73420	NW_009562453.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold60944	Nicotiana tabacum	2567	3577	89%	0.0	81.79%	61145	NW_015922636.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold845	Nicotiana tabacum	2566	3529	89%	0.0	81.85%	63114	NW_015942931.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold25717	Nicotiana tomentosiformis	2566	3459	84%	0.0	82.75%	53334	NW_008905505.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold387	Nicotiana sylvestris	2562	3363	86%	0.0	81.84%	302458	NW_009524608.1
CopiaSL_25	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2r	Nicotiana attenuata	2560	3472	89%	0.0	81.79%	722875	NW_017670432.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold17033	Nicotiana tomentosiformis	2558	3603	87%	0.0	82.19%	63442	NW_008895857.1

CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9194	Nicotiana tabacum	2556	5498	89%	0.0	81.74%	179722	NW_015949073.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold10916	Nicotiana sylvestris	2556	3762	85%	0.0	81.88%	56894	NW_009348981.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold36722	Nicotiana sylvestris	2556	3660	84%	0.0	83.19%	81588	NW_009522412.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5389	Nicotiana tomentosiformis	2556	3632	89%	0.0	82.19%	257796	NW_008936807.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4570	Nicotiana tomentosiformis	2555	3438	84%	0.0	81.77%	113084	NW_008927708.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold119191	Nicotiana tabacum	2553	3529	89%	0.0	81.76%	77192	NW_015801726.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold54080	Nicotiana tabacum	2551	5202	89%	0.0	81.87%	199814	NW_015916480.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold21337	Nicotiana tomentosiformis	2551	3601	89%	0.0	81.72%	61156	NW_008900639.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold95079	Nicotiana tabacum	2549	3545	89%	0.0	81.68%	14382	NW_015951580.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13337	Nicotiana tabacum	2545	3355	86%	0.0	81.69%	59273	NW_015811818.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1928	Nicotiana tabacum	2545	3575	89%	0.0	81.71%	100516	NW_015847830.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24339	Nicotiana tabacum	2543	3503	89%	0.0	81.68%	36413	NW_015872426.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold69274	Nicotiana tabacum	2540	3582	89%	0.0	81.64%	39023	NW_015929967.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold71517	Nicotiana tabacum	2540	3725	85%	0.0	81.79%	49985	NW_015931892.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1920	Nicotiana tomentosiformis	2538	4861	89%	0.0	81.72%	177443	NW_008898264.1
CopiaSL_25	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00025	Solanum tuberosum	2534	4074	84%	0.0	83.24%	2980670	NW_006238951.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold31511	Nicotiana tabacum	2521	5924	86%	0.0	82.44%	183861	NW_015889474.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold92590	Nicotiana tabacum	2521	3523	89%	0.0	81.68%	144875	NW_015949622.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold19584	Nicotiana tomentosiformis	2521	3540	86%	0.0	82.33%	110401	NW_008898690.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold35934	Nicotiana tomentosiformis	2519	3345	86%	0.0	81.55%	169841	NW_008916857.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold47975	Nicotiana tabacum	2512	3448	83%	0.0	83.27%	153674	NW_015910898.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold53745	Nicotiana tabacum	2512	3485	89%	0.0	81.51%	127682	NW_015916178.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold10035	Nicotiana tomentosiformis	2510	3414	89%	0.0	81.47%	58683	NW_008828883.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold15985	Nicotiana tomentosiformis	2507	3226	84%	0.0	81.51%	134424	NW_008894691.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2670	Nicotiana tomentosiformis	2501	3357	89%	0.0	81.51%	18949	NW_008906597.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold27039	Nicotiana tabacum	2497	3326	83%	0.0	82.11%	107677	NW_015885248.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold14399	Nicotiana tabacum	2494	3569	82%	0.0	83.46%	130517	NW_015819031.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold90693	Nicotiana tabacum	2490	3468	87%	0.0	82.01%	16272	NW_015948044.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold22647	Nicotiana tomentosiformis	2488	3237	83%	0.0	81.41%	72783	NW_008902094.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold40449	Nicotiana tabacum	2483	3390	89%	0.0	81.52%	23928	NW_015903932.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold41268	Nicotiana tabacum	2479	4590	85%	0.0	82.51%	52027	NW_015904709.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold126524	Nicotiana tabacum	2468	3516	89%	0.0	81.38%	83098	NW_015807042.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold7362	Nicotiana tabacum	2462	3411	81%	0.0	82.92%	72050	NW_015933702.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold3764	Nicotiana tabacum	2460	3377	88%	0.0	81.32%	77686	NW_015901333.1
CopiaSL_25	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold51044	Nicotiana sylvestris	2460	3388	88%	0.0	81.32%	50961	NW_009538326.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold35876	Nicotiana tabacum	2457	3439	89%	0.0	81.15%	316815	NW_015899681.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold42458	Nicotiana tabacum	2451	3737	89%	0.0	82.50%	197412	NW_015905802.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold27621	Nicotiana tabacum	2446	3918	89%	0.0	81.13%	98018	NW_015885795.1
CopiaSL_25	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2834	Nicotiana tomentosiformis	2444	3335	87%	0.0	81.18%	366940	NW_008908419.1
CopiaSL_25	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold153438	Nicotiana tabacum	2440	3562	82%	0.0	83.15%	15468	NW_015825086.1
CopiaSL_38	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	10504	82556	100%	0.0	100.00%	55977580	NC_015439.3
CopiaSL_38	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	9546	87841	100%	0.0	96.96%	60730942	NC_028642.1
CopiaSL_38	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	9526	1,23E+08	100%	0.0	96.89%	109333515	NC_028637.1
CopiaSL_38	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum pennellii	9520	1,04E+08	100%	0.0	96.88%	77991103	NC_028641.1
CopiaSL_38	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	Solanum pennellii	9520	1,01E+08	100%	0.0	96.88%	84057508	NC_028645.1
CopiaSL_38	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	Solanum pennellii	9520	85135	100%	0.0	96.88%	66223686	NC_028647.1
CopiaSL_38	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	Solanum pennellii	9518	84029	100%	0.0	96.88%	83305730	NC_028648.1
CopiaSL_38	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	9446	1,02E+08	100%	0.0	96.68%	75414019	NC_028639.1
CopiaSL_38	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	9435	77612	100%	0.0	96.65%	77197300	NC_028640.1
CopiaSL_38	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	9367	1,09E+08	100%	0.0	96.39%	72906345	NC_015446.3
CopiaSL_38	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	9282	90944	100%	0.0	96.19%	68175699	NC_015444.3
CopiaSL_38	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	Solanum pennellii	9073	1,41E+08	100%	0.0	96.02%	70546659	NC_028644.1
CopiaSL_38	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	8914	84014	100%	0.0	95.09%	65633393	NC_015447.3
CopiaSL_38	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	8672	72985	100%	0.0	94.37%	66557038	NC_015441.3
CopiaSL_38	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	8578	76363	100%	0.0	94.01%	66723567	NC_015442.3
CopiaSL_38	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	8421	93913	100%	0.0	93.54%	65987440	NC_015445.3

CopiaSL_38	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	8183	1,25E+08	100%	0.0	92.74%	56597135	NC_015448.3
CopiaSL_38	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	8109	51662	100%	0.0	95.17%	49794276	NC_015443.3
CopiaSL_38	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	Solanum pennellii	7947	76656	100%	0.0	95.99%	79292169	NC_028643.1
CopiaSL_38	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	7533	73998	100%	0.0	94.98%	72290146	NC_015440.3
CopiaSL_38	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	6870	83873	100%	0.0	96.28%	59803892	NC_028638.1
CopiaSL_38	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum pennellii	6370	66239	100%	0.0	94.20%	82529941	NC_028646.1
CopiaSL_38	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	6292	79262	100%	0.0	93.82%	98455869	NC_015438.3
CopiaSL_38	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	6107	53729	100%	0.0	93.06%	68126176	NC_015449.3
CopiaSL_38	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00364	Solanum tuberosum	5600	8697	99%	0.0	90.72%	609531	NW_006239291.1
CopiaSL_38	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00084	Solanum tuberosum	5249	6447	93%	0.0	89.22%	1918041	NW_006239099.1
CopiaSL_38	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00190	Solanum tuberosum	3951	4620	80%	0.0	85.83%	1188304	NW_006239114.1
CopiaSL_38	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00186	Solanum tuberosum	3917	12069	80%	0.0	85.69%	1196459	NW_006239112.1
CopiaSL_38	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 8, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3570	1,07E+08	89%	0.0	84.53%	153299543	NC_029984.1
CopiaSL_38	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 6, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3552	2,17E+08	89%	0.0	84.31%	219521584	NC_029982.1
CopiaSL_38	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 1, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3535	2,81E+08	93%	0.0	84.21%	301019445	NC_029977.1
CopiaSL_38	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 2, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3533	1,98E+08	89%	0.0	84.27%	163962470	NC_029978.1
CopiaSL_38	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3526	2,13E+08	92%	0.0	84.24%	220335243	NC_029987.1
CopiaSL_38	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3517	1,81E+08	88%	0.0	84.55%	215701946	NC_029980.1
CopiaSL_38	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3491	1,74E+08	89%	0.0	83.72%	222112641	NC_029983.1
CopiaSL_38	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3485	3,04E+08	89%	0.0	84.01%	261510930	NC_029979.1
CopiaSL_38	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3480	1,98E+08	89%	0.0	83.96%	205736368	NC_029986.1
CopiaSL_38	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3461	2,18E+08	88%	0.0	84.19%	229934170	NC_029988.1
CopiaSL_38	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3456	1,51E+08	93%	0.0	84.26%	217274494	NC_029981.1
CopiaSL_38	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	3434	2,05E+08	91%	0.0	84.16%	238794889	NC_029985.1
CopiaSL_38	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2719	Capsicum annuum	3378	4654	85%	0.0	83.81%	277434	NW_015960900.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 6, NIATTR2	Nicotiana attenuata	3343	60756	81%	0.0	83.00%	97571839	NC_031994.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 7, NIATTR2	Nicotiana attenuata	3306	88649	87%	0.0	82.82%	114765206	NC_031995.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 12, NIATTR2	Nicotiana attenuata	3301	51885	88%	0.0	82.79%	53491057	NC_032000.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 11, NIATTR2	Nicotiana attenuata	3295	56181	87%	0.0	82.78%	61252062	NC_031999.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3293	3701	86%	0.0	82.72%	440631	NW_017670830.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3282	3697	81%	0.0	82.73%	357002	NW_017671104.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 9, NIATTR2	Nicotiana attenuata	3273	50911	88%	0.0	82.66%	50975570	NC_031997.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3271	3660	86%	0.0	82.67%	142057	NW_017673286.1
CopiaSL_38	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold2267	Capsicum annuum	3267	12338	88%	0.0	82.97%	3493078	NW_015960517.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3264	3689	86%	0.0	82.74%	69780	NW_017675337.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3254	3643	80%	0.0	82.57%	234944	NW_017671928.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 1, NIATTR2	Nicotiana attenuata	3249	80759	88%	0.0	82.56%	110538649	NC_031989.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 3, NIATTR2	Nicotiana attenuata	3238	60574	82%	0.0	82.53%	72893213	NC_031991.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3238	3592	80%	0.0	82.48%	87827	NW_017674699.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 2, NIATTR2	Nicotiana attenuata	3236	68498	88%	0.0	82.49%	96359640	NC_031990.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 8, NIATTR2	Nicotiana attenuata	3227	50028	87%	0.0	82.45%	47755514	NC_031996.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3219	3718	86%	0.0	82.48%	170758	NW_017672766.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3219	3671	86%	0.0	82.43%	119157	NW_017673810.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3214	3595	85%	0.0	82.42%	361967	NW_017671081.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3201	3610	86%	0.0	82.31%	109842	NW_017674065.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3175	3571	80%	0.0	82.21%	210106	NW_017672208.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3173	3650	81%	0.0	82.19%	1570162	NW_017670262.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3169	3685	86%	0.0	82.22%	187560	NW_017672500.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3157	3579	86%	0.0	82.21%	45636	NW_017676394.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3155	3732	86%	0.0	82.13%	1243377	NW_017670288.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3136	3571	86%	0.0	82.05%	207692	NW_017672229.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3118	3514	86%	0.0	81.91%	81235	NW_017674905.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3096	3561	84%	0.0	82.34%	954013	NW_017670339.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 10, NIATTR2	Nicotiana attenuata	3092	37130	82%	0.0	81.75%	51082206	NC_031998.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3085	3455	81%	0.0	81.82%	233323	NW_017671942.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTR2	Nicotiana attenuata	3083	3448	86%	0.0	81.77%	270994	NW_017671615.1

CopiaSL_38	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold5313	Nicotiana sylvestris	3053	3488	86%	0.0	81.60%	160185	NW_009540643.1
CopiaSL_38	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold968	Nicotiana sylvestris	3040	3462	86%	0.0	81.54%	129181	NW_009589163.1
CopiaSL_38	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 4, NIATTR2	Nicotiana attenuata	2970	10759	88%	0.0	81.44%	12333206	NC_031992.1
CopiaSL_38	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold20987	Nicotiana tomentosiformis	2802	5336	82%	0.0	79.58%	173145	NW_008900249.1
GypsySL_04	refseq_genomes	8	Capicum annum cultivar Zunla-1 unplaced genomic scaffold, Pepper Zunla 1 Ref_v1.0 scaffold1165	Capicum annum	2763	4156	84%	0.0	83.86%	67805	NW_015960689.1
GypsySL_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	25994	3,08E+09	100%	0.0	100.00%	98455869	NC_015438.3
GypsySL_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	13173	1,71E+09	100%	0.0	94.62%	49794276	NC_015443.3
GypsySL_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	13101	2,23E+09	100%	0.0	94.49%	56597135	NC_015448.3
GypsySL_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	12992	2,49E+09	100%	0.0	87.12%	72290146	NC_015440.3
GypsySL_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	12936	3,01E+09	100%	0.0	94.27%	68126176	NC_015449.3
GypsySL_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	12562	2,43E+09	100%	0.0	93.32%	66557038	NC_015441.3
GypsySL_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	12305	2,51E+09	100%	0.0	96.27%	65987440	NC_015445.3
GypsySL_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	12209	3,01E+09	100%	0.0	92.79%	65633393	NC_015447.3
GypsySL_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	12098	3,21E+09	100%	0.0	92.32%	66723567	NC_015442.3
GypsySL_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	12046	3,31E+09	100%	0.0	95.27%	68175699	NC_015444.3
GypsySL_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	11972	1,44E+09	100%	0.0	92.09%	55977580	NC_015439.3
GypsySL_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	11707	3,00E+09	100%	0.0	91.58%	72906345	NC_015446.3
GypsySL_04	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	Solanum pennellii	11703	2,78E+09	100%	0.0	91.49%	70546659	NC_028644.1
GypsySL_04	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	11418	2,85E+09	100%	0.0	92.05%	75414019	NC_028639.1
GypsySL_04	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	Solanum pennellii	11295	2,68E+09	100%	0.0	92.81%	66223686	NC_028647.1
GypsySL_04	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum pennellii	11215	3,87E+09	100%	0.0	92.49%	77991103	NC_028641.1
GypsySL_04	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum pennellii	11188	3,33E+09	100%	0.0	92.60%	82529941	NC_028646.1
GypsySL_04	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	Solanum pennellii	11031	2,88E+09	100%	0.0	90.32%	84057508	NC_028645.1
GypsySL_04	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	Solanum pennellii	10473	3,30E+09	100%	0.0	94.37%	79292169	NC_028643.1
GypsySL_04	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	Solanum pennellii	10469	3,51E+09	100%	0.0	91.55%	83305730	NC_028648.1
GypsySL_04	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	10434	2,22E+09	100%	0.0	92.78%	60730942	NC_028642.1
GypsySL_04	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	10286	1,70E+09	100%	0.0	93.26%	59803892	NC_028638.1
GypsySL_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC00000008	Solanum lycopersicum	10220	94345	97%	0.0	89.45%	1577018	NW_020442489.1
GypsySL_04	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	9310	3,03E+09	100%	0.0	87.58%	77197300	NC_028640.1
GypsySL_04	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	8410	2,95E+09	100%	0.0	87.25%	109333515	NC_028637.1
GypsySL_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC00000001	Solanum lycopersicum	5869	1,34E+08	92%	0.0	88.16%	2014460	NW_020442482.1
GypsySL_04	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC00000002	Solanum lycopersicum	4034	25789	80%	0.0	82.48%	177393	NW_020442483.1
GypsySL_06	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	36044	4,03E+08	100%	0.0	99.94%	65633393	NC_015447.3
GypsySL_06	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	22864	3,80E+08	95%	0.0	97.87%	68175699	NC_015444.3
GypsySL_06	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	21213	5,04E+08	85%	0.0	95.92%	98455869	NC_015438.3
GypsySL_06	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	19765	4,40E+08	82%	0.0	96.90%	72906345	NC_015446.3
GypsySL_06	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	19734	3,70E+08	80%	0.0	96.70%	65987440	NC_015445.3
GypsySL_06	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	15272	3,40E+08	84%	0.0	97.16%	72290146	NC_015440.3
GypsySL_06	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	14840	1,42E+08	81%	0.0	90.11%	49794276	NC_015443.3
GypsySL_06	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	14218	2,13E+08	83%	0.0	86.40%	60730942	NC_028642.1
GypsySL_06	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	13923	3,11E+08	81%	0.0	86.49%	75414019	NC_028639.1
GypsySL_06	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	13324	2,68E+08	84%	0.0	85.97%	66557038	NC_015441.3
GypsySL_06	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	13282	2,70E+08	81%	0.0	86.57%	77197300	NC_028640.1
GypsySL_06	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	12957	1,39E+08	90%	0.0	85.73%	55977580	NC_015439.3
GypsySL_06	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	12748	3,99E+08	81%	0.0	86.57%	109333515	NC_028637.1
GypsySL_06	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	11932	2,70E+08	82%	0.0	84.04%	68126176	NC_015449.3
GypsySL_06	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	10733	3,05E+08	83%	0.0	85.84%	66723567	NC_015442.3
GypsySL_06	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	10534	2,19E+08	82%	0.0	85.75%	59803892	NC_028638.1
GypsySL_06	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	10081	2,39E+08	81%	0.0	85.79%	56597135	NC_015448.3
GypsySL_12	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	22005	1,95E+08	100%	0.0	100.00%	56597135	NC_015448.3
GypsySL_12	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	13376	4,49E+08	100%	0.0	95.25%	60730942	NC_028642.1
GypsySL_12	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum pennellii	12172	2,46E+08	100%	0.0	97.45%	77991103	NC_028641.1
GypsySL_12	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	12028	1,65E+08	100%	0.0	97.09%	75414019	NC_028639.1
GypsySL_12	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	11995	3,58E+08	100%	0.0	97.01%	77197300	NC_028640.1
GypsySL_12	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum pennellii	11936	4,38E+08	100%	0.0	96.85%	82529941	NC_028646.1
GypsySL_12	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	11756	4,17E+08	100%	0.0	96.42%	109333515	NC_028637.1

GypsySL_12	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	Solanum pennellii	11622	1,54E+08	100%	0.0	96.05%	79292169	NC_028643.1
GypsySL_12	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	11609	1,44E+08	100%	0.0	96.04%	66557038	NC_015441.3
GypsySL_12	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	Solanum pennellii	11339	3,58E+08	100%	0.0	95.36%	84057508	NC_028645.1
GypsySL_12	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	11337	3,00E+08	100%	0.0	95.35%	66723567	NC_015442.3
GypsySL_12	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	11308	1,49E+08	100%	0.0	95.34%	68126176	NC_015449.3
GypsySL_12	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	11169	4,26E+08	100%	0.0	94.89%	59803892	NC_028638.1
GypsySL_12	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	11051	1,94E+08	100%	0.0	94.96%	72290146	NC_015440.3
GypsySL_12	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	10918	1,77E+08	100%	0.0	94.33%	72906345	NC_015446.3
GypsySL_12	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	10909	1,57E+08	100%	0.0	94.35%	65987440	NC_015445.3
GypsySL_12	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	Solanum pennellii	10562	4,46E+08	100%	0.0	93.39%	70546659	NC_028644.1
GypsySL_12	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	Solanum pennellii	10532	1,70E+08	100%	0.0	93.40%	66223686	NC_028647.1
GypsySL_12	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	10414	3,16E+08	100%	0.0	95.50%	65633393	NC_015447.3
GypsySL_12	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	10146	2,03E+08	100%	0.0	94.55%	98455869	NC_015438.3
GypsySL_12	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	10050	1,40E+08	100%	0.0	95.26%	68175699	NC_015444.3
GypsySL_12	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	Solanum pennellii	9845	3,53E+08	100%	0.0	94.77%	83305730	NC_028648.1
GypsySL_12	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00368	Solanum tuberosum	8087	13167	81%	0.0	87.19%	608325	NW_006239292.1
GypsySL_12	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	7849	94725	100%	0.0	92.66%	49794276	NC_015443.3
GypsySL_12	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	7731	1,34E+08	100%	0.0	95.15%	55977580	NC_015439.3
GypsySL_12	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC0000001	Solanum lycopersicum	6698	18378	92%	0.0	91.94%	2014460	NW_020442482.1
GypsySL_12	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00085	Solanum tuberosum	6466	13100	80%	0.0	87.41%	1839851	NW_006239014.1
GypsySL_12	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00006	Solanum tuberosum	5112	25564	89%	0.0	82.81%	4150974	NW_006238932.1
GypsySL_12	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 3, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2061	1,30E+08	82%	0.0	78.19%	261510930	NC_029979.1
GypsySL_12	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2026	68486	81%	0.0	78.39%	215701946	NC_02980.1
GypsySL_12	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	1941	83439	80%	0.0	78.04%	222112641	NC_029983.1
psySL_monotypic Ch03_1s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	11049	14322	100%	0.0	100.00%	72290146	NC_015440.3
psySL_monotypic Ch03_1s	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	6693	12649	100%	0.0	97.27%	60730942	NC_028642.1
psySL_monotypic Ch03_1s	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	Solanum pennellii	5834	10193	84%	0.0	93.52%	84057508	NC_028645.1
psySL_monotypic Ch03_1s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	3308	10686	89%	0.0	92.71%	72906345	NC_015446.3
psySL_monotypic Ch03_1s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	1633	8885	93%	0.0	88.81%	55977580	NC_015439.3
psySL_monotypic Ch12_1s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	10115	11653	100%	0.0	100.00%	68126176	NC_015449.3
psySL_monotypic Ch12_1s	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	8780	10405	100%	0.0	95.62%	75414019	NC_028639.1
psySL_monotypic Ch12_1s	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	Solanum pennellii	8227	18797	99%	0.0	93.95%	66223686	NC_028647.1
psySL_monotypic Ch12_1s	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum pennellii	7738	8828	99%	0.0	92.29%	82529941	NC_028646.1
psySL_monotypic Ch12_1s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	7505	12134	99%	0.0	95.11%	98455869	NC_015438.3
psySL_monotypic Ch12_1s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	7265	7791	99%	0.0	90.80%	65633393	NC_015447.3
psySL_monotypic Ch12_1s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	6756	9634	98%	0.0	94.62%	68175699	NC_015444.3
psySL_monotypic Ch12_1s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	5958	13716	99%	0.0	94.36%	56597135	NC_015448.3
psySL_monotypic Ch12_1s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	5483	15568	99%	0.0	93.90%	49794276	NC_015443.3
psySL_monotypic Ch12_1s	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	5317	8342	98%	0.0	91.94%	77197300	NC_028640.1
psySL_monotypic Ch12_1s	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	Solanum pennellii	4065	4065	82%	0.0	82.90%	70546659	NC_028644.1
psySL_monotypic Ch12_1s	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	4058	4058	82%	0.0	82.89%	109333515	NC_028637.1
psySL_monotypic Ch12_1s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	3862	3862	82%	0.0	82.09%	65987440	NC_015445.3
psySL_monotypic Ch12_1s	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	3179	9558	87%	0.0	79.51%	60730942	NC_028642.1
psySL_monotypic Ch12_1s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	2617	6211	82%	0.0	79.07%	72290146	NC_015440.3
psySL_monotypic Ch12_1s	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	2364	4711	82%	0.0	83.84%	66723567	NC_015442.3
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold22471	Nicotiana tomentosiformis	5609	7379	96%	0.0	87.41%	52716	NW_008901899.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2282	Nicotiana tomentosiformis	5603	7615	96%	0.0	87.39%	237567	NW_008902286.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold2755	Nicotiana tomentosiformis	5600	7617	96%	0.0	87.37%	146752	NW_008907541.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold34143	Nicotiana tomentosiformis	5598	7613	96%	0.0	87.36%	121277	NW_008914868.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold24902	Nicotiana tomentosiformis	5583	14019	96%	0.0	87.31%	54776	NW_008904600.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31687	Nicotiana tomentosiformis	5581	13647	96%	0.0	87.31%	77527	NW_008912138.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold61699	Nicotiana tomentosiformis	5581	7891	99%	0.0	87.31%	32015	NW_008945484.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold11955	Nicotiana tomentosiformis	5559	6540	94%	0.0	86.64%	349838	NW_008850213.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold27563	Nicotiana tabacum	5485	6640	99%	0.0	87.01%	127279	NW_015885740.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold32900	Nicotiana tabacum	5465	6674	96%	0.0	88.26%	53101	NW_015890776.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold7623	Nicotiana tomentosiformis	5432	6707	97%	0.0	88.19%	66778	NW_008961630.1

Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13868	Nicotiana tabacum	5402	7110	94%	0.0	87.97%	134467	NW_015815476.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17909	Nicotiana tabacum	5402	6840	93%	0.0	88.05%	287030	NW_015840282.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold13433	Nicotiana tomentosiformis	5400	8297	99%	0.0	87.09%	64717	NW_008866636.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold36367	Nicotiana tabacum	5385	5905	83%	0.0	87.90%	27763	NW_015900142.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold39652	Nicotiana tabacum	5385	6922	99%	0.0	86.65%	96441	NW_015903218.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold64876	Nicotiana tabacum	5380	6606	92%	0.0	86.88%	124117	NW_015926130.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold57928	Nicotiana tabacum	5376	6768	97%	0.0	87.97%	63354	NW_015919969.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold43655	Nicotiana tabacum	5369	6126	87%	0.0	87.84%	96415	NW_015906914.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3471	Nicotiana tomentosiformis	5352	6117	88%	0.0	87.78%	184701	NW_008915497.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13367	Nicotiana tabacum	5349	6486	99%	0.0	86.48%	65428	NW_015812023.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold81533	Nicotiana tabacum	5339	9013	96%	0.0	87.24%	73683	NW_015940479.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold28867	Nicotiana tabacum	5336	6783	98%	0.0	87.72%	118211	NW_015886974.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold246479	Nicotiana tabacum	5319	7744	98%	0.0	87.72%	10289	NW_015873837.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold28097	Nicotiana tabacum	5319	7153	97%	0.0	87.67%	40676	NW_015886258.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1251	Nicotiana tomentosiformis	5317	7331	94%	0.0	87.67%	155978	NW_008856380.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold7430	Nicotiana tomentosiformis	5313	6417	94%	0.0	87.64%	53933	NW_008959486.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5057	Nicotiana tomentosiformis	5312	7395	94%	0.0	87.66%	139171	NW_008933119.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold29244	Nicotiana tabacum	5301	6195	86%	0.0	87.60%	98940	NW_015887333.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold102161	Nicotiana tabacum	5299	6392	85%	0.0	89.36%	5356	NW_015788916.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold50219	Nicotiana tomentosiformis	5297	7115	97%	0.0	87.58%	45028	NW_008932730.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold21492	Nicotiana tabacum	5295	6601	99%	0.0	86.28%	214967	NW_015859021.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10122	Nicotiana tabacum	5289	6578	94%	0.0	87.54%	150655	NW_015788196.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold15436	Nicotiana tomentosiformis	5289	6471	99%	0.0	86.27%	403882	NW_008888891.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold47540	Nicotiana tomentosiformis	5286	7758	98%	0.0	87.53%	46274	NW_008929753.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold173	Nicotiana tomentosiformis	5276	6824	92%	0.0	86.24%	122173	NW_008896152.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold84221	Nicotiana tabacum	5258	7342	99%	0.0	86.17%	61925	NW_015942700.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5408	Nicotiana tomentosiformis	5256	6313	91%	0.0	87.38%	119859	NW_008937019.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold21789	Nicotiana tabacum	5240	7029	94%	0.0	87.45%	116944	NW_015860458.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold44711	Nicotiana tomentosiformis	5240	7639	98%	0.0	87.39%	76181	NW_008926610.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold35404	Nicotiana tabacum	5238	6256	89%	0.0	87.92%	39824	NW_015899239.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold41507	Nicotiana tomentosiformis	5238	7069	94%	0.0	87.45%	141329	NW_008923050.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold107834	Nicotiana tabacum	5230	7008	94%	0.0	87.45%	50278	NW_015793285.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold21025	Nicotiana tomentosiformis	5228	6799	89%	0.0	87.35%	11743	NW_008900293.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold12157	Nicotiana tomentosiformis	5225	7014	94%	0.0	87.43%	208333	NW_008852458.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold107774	Nicotiana tabacum	5221	7309	95%	0.0	87.36%	18353	NW_015793235.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold10844	Nicotiana tomentosiformis	5221	12148	98%	0.0	87.35%	317981	NW_008837871.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9612	Nicotiana tabacum	5219	6256	89%	0.0	87.42%	360222	NW_015952413.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold14682	Nicotiana tomentosiformis	5216	7072	95%	0.0	87.35%	149872	NW_008880513.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4876	Nicotiana tabacum	5210	7328	98%	0.0	87.33%	313082	NW_015911612.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold55355	Nicotiana tabacum	5210	6200	89%	0.0	87.24%	66341	NW_015917636.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold7971	Nicotiana tomentosiformis	5203	6245	89%	0.0	87.28%	124454	NW_008965496.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold24021	Nicotiana tomentosiformis	5199	7750	98%	0.0	87.18%	34549	NW_008903622.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold412	Nicotiana tomentosiformis	5192	5999	85%	0.0	87.02%	131134	NW_008922708.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold2428	Nicotiana tabacum	5190	6354	93%	0.0	87.15%	115131	NW_015872154.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold977	Nicotiana tomentosiformis	5179	12829	99%	0.0	85.86%	190588	NW_008985484.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold100771	Nicotiana tabacum	5177	6814	98%	0.0	87.09%	15965	NW_015787838.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold49889	Nicotiana tabacum	5177	6284	88%	0.0	88.51%	128702	NW_015912645.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold93949	Nicotiana tabacum	5160	12052	98%	0.0	87.11%	110237	NW_015950675.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold8263	Nicotiana tomentosiformis	5149	6928	98%	0.0	86.98%	75359	NW_008968741.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold72476	Nicotiana tabacum	5136	6739	99%	0.0	86.92%	57868	NW_015932742.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold38706	Nicotiana tomentosiformis	5131	5651	89%	0.0	85.70%	7567	NW_008919937.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold57100	Nicotiana tabacum	5121	5752	84%	0.0	87.58%	68372	NW_015919215.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3829	Nicotiana tomentosiformis	5116	9830	98%	0.0	88.41%	81699	NW_008919474.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold41437	Nicotiana tabacum	5103	6064	89%	0.0	87.60%	37387	NW_015904868.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold21451	Nicotiana tomentosiformis	5075	6715	89%	0.0	87.87%	49133	NW_008900766.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold94114	Nicotiana tabacum	5068	5949	88%	0.0	86.78%	35560	NW_015950817.1

Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold45661	Nicotiana tabacum	5059	7057	93%	0.0	88.13%	44993	NW_015908753.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold12044	Nicotiana tabacum	5020	6506	94%	0.0	88.00%	140177	NW_015802636.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold29654	Nicotiana tabacum	4990	6622	94%	0.0	87.89%	48458	NW_015887720.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold23853	Nicotiana tomentosiformis	4987	6273	91%	0.0	86.46%	59679	NW_008903434.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold98280	Nicotiana tabacum	4975	6059	90%	0.0	86.41%	41720	NW_015954119.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold3209	Nicotiana tomentosiformis	4974	6645	94%	0.0	86.34%	61053	NW_008912586.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold12554	Nicotiana tomentosiformis	4918	5718	82%	0.0	87.80%	227451	NW_008856869.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 1, NIATTr2	Nicotiana attenuata	4911	1,10E+08	98%	0.0	86.09%	110538649	NC_031989.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold131700	Nicotiana tabacum	4881	6569	92%	0.0	86.58%	27395	NW_015810690.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold41550	Nicotiana tabacum	4881	5381	81%	0.0	86.61%	160711	NW_015904978.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold38003	Nicotiana tabacum	4878	5683	82%	0.0	87.64%	54461	NW_015901679.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 4, NIATTr2	Nicotiana attenuata	4876	38269	98%	0.0	86.11%	12333206	NC_031992.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold8249	Nicotiana tomentosiformis	4876	6004	92%	0.0	86.58%	260191	NW_008968585.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold5528	Nicotiana sylvestris	4868	11591	95%	0.0	85.95%	113997	NW_009543031.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 3, NIATTr2	Nicotiana attenuata	4863	81257	98%	0.0	85.96%	72893213	NC_031991.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 10, NIATTr2	Nicotiana attenuata	4863	38904	98%	0.0	85.94%	5108206	NC_031998.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 6, NIATTr2	Nicotiana attenuata	4850	89661	99%	0.0	85.88%	97571839	NC_031994.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4839	6260	94%	0.0	85.83%	432375	NW_017670855.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4839	6684	97%	0.0	85.85%	249786	NW_017671800.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4837	6055	98%	0.0	85.83%	99899	NW_017674356.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 5, NIATTr2	Nicotiana attenuata	4835	46131	98%	0.0	85.86%	57012777	NC_031993.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold175476	Nicotiana tabacum	4835	6120	94%	0.0	85.79%	42841	NW_015838272.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4824	6234	97%	0.0	85.77%	674142	NW_017670458.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4824	5931	89%	0.0	85.77%	236676	NW_017671917.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold15881	Nicotiana tabacum	4824	11057	95%	0.0	85.75%	50792	NW_015828447.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold57240	Nicotiana tabacum	4824	5647	92%	0.0	85.75%	128756	NW_015919338.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4815	6035	89%	0.0	85.74%	663437	NW_017670473.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold37781	Nicotiana tabacum	4813	5670	94%	0.0	85.70%	119546	NW_0319901468.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4807	6397	90%	0.0	85.71%	150848	NW_017673111.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold80	Nicotiana sylvestris	4800	6023	87%	0.0	85.72%	226580	NW_009570497.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4798	6379	98%	0.0	85.78%	185398	NW_017672537.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 2, NIATTr2	Nicotiana attenuata	4796	68346	98%	0.0	85.66%	96359640	NC_031990.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4796	6650	97%	0.0	85.63%	195897	NW_017672385.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4795	13008	98%	0.0	85.66%	321437	NW_017671287.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1311	Nicotiana tomentosiformis	4789	10250	98%	0.0	86.80%	48416	NW_008863047.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4782	6231	95%	0.0	85.59%	219144	NW_017672091.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4780	6235	97%	0.0	85.57%	526680	NW_017670637.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4778	7089	97%	0.0	85.60%	1026265	NW_017670317.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4776	6561	98%	0.0	85.61%	435747	NW_017670841.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4763	6134	97%	0.0	85.58%	347664	NW_017671149.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4761	5780	94%	0.0	85.56%	360021	NW_017671088.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4759	6217	97%	0.0	85.51%	28186	NW_017677689.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold31152	Nicotiana sylvestris	4758	5993	94%	0.0	85.48%	199439	NW_009516224.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4756	5577	88%	0.0	85.54%	287943	NW_017671495.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4739	63412	98%	0.0	85.43%	61252062	NC_031999.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold23275	Nicotiana sylvestris	4737	6840	95%	0.0	85.41%	52891	NW_009486303.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4735	6366	97%	0.0	85.39%	85001	NW_017674777.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4726	6498	98%	0.0	85.38%	153820	NW_017673045.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4719	6158	97%	0.0	85.36%	74228	NW_017675158.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4717	8438	97%	0.0	85.36%	46410	NW_017676365.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4713	7558	94%	0.0	85.38%	192976	NW_017672423.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold23423	Nicotiana sylvestris	4713	5330	85%	0.0	85.31%	70177	NW_009487948.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4711	5519	86%	0.0	85.35%	336284	NW_017671202.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4708	5653	87%	0.0	85.34%	373821	NW_017671042.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4706	6021	90%	0.0	85.28%	650487	NW_017670484.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4704	5843	89%	0.0	85.32%	446766	NW_017670814.1

Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4700	7223	97%	0.0	85.32%	1612275	NW_017670256.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4700	6416	97%	0.0	86.17%	459799	NW_017670773.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4697	5582	88%	0.0	85.29%	192746	NW_017672426.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold49054	Nicotiana sylvestris	4697	5260	85%	0.0	85.27%	76173	NW_009536114.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4695	6209	97%	0.0	85.26%	443537	NW_017670821.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4693	6108	97%	0.0	85.21%	149826	NW_017673132.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4689	12189	98%	0.0	85.25%	500387	NW_017670684.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold27364	Nicotiana sylvestris	4686	5657	86%	0.0	85.22%	59137	NW_009512014.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 7, NIATTr2	Nicotiana attenuata	4684	80380	98%	0.0	85.26%	114765206	NC_031995.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24853	Nicotiana tabacum	4678	5225	87%	0.0	85.20%	234295	NW_015874717.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4673	5564	91%	0.0	85.13%	490964	NW_017670699.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4669	6009	93%	0.0	85.13%	75273	NW_017675128.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4658	6437	98%	0.0	85.12%	166205	NW_017672832.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4656	5294	89%	0.0	85.11%	10111	NW_017682319.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold7606	Nicotiana tomentosiformis	4652	5716	85%	0.0	87.61%	276920	NW_008961441.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 12, NIATTr2	Nicotiana attenuata	4650	41583	98%	0.0	85.09%	53491057	NC_032000.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4650	5446	88%	0.0	85.58%	141412	NW_017673299.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold99684	Nicotiana tabacum	4650	5220	85%	0.0	85.09%	43248	NW_015955213.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4649	5784	97%	0.0	85.10%	1065697	NW_017670311.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4649	5278	85%	0.0	85.14%	85934	NW_017674756.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4638	5629	91%	0.0	85.04%	168467	NW_017672793.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold33253	Nicotiana sylvestris	4632	5453	87%	0.0	84.98%	88439	NW_009518558.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4630	5544	87%	0.0	85.03%	457135	NW_017670785.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold22454	Nicotiana sylvestris	4626	5691	99%	0.0	83.81%	141987	NW_009477181.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold19211	Nicotiana sylvestris	4625	5615	95%	0.0	85.00%	206434	NW_009441148.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold58775	Nicotiana tabacum	4617	5982	94%	0.0	85.00%	80956	NW_015920707.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold39059	Nicotiana tabacum	4615	6016	94%	0.0	84.98%	165586	NW_015902664.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 9, NIATTr2	Nicotiana attenuata	4614	31101	98%	0.0	85.00%	50975570	NC_031997.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8174	Nicotiana tabacum	4614	5521	93%	0.0	84.96%	19877	NW_015940639.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold90771	Nicotiana tabacum	4610	5669	99%	0.0	83.83%	60358	NW_015948114.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 8, NIATTr2	Nicotiana attenuata	4606	37785	98%	0.0	84.95%	47755514	NC_031996.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4602	5560	87%	0.0	85.63%	247665	NW_017671821.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1472	Nicotiana sylvestris	4599	5538	94%	0.0	84.92%	226595	NW_009391247.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4597	4978	83%	0.0	84.86%	251695	NW_017671782.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold12996	Nicotiana sylvestris	4595	5991	94%	0.0	84.92%	214613	NW_009372091.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold5512	Nicotiana sylvestris	4595	5847	94%	0.0	84.88%	119718	NW_009542854.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4591	5448	86%	0.0	84.91%	197841	NW_017672358.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold123387	Nicotiana sylvestris	4578	5439	87%	0.0	84.87%	12970	NW_009364789.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold57680	Nicotiana tomentosiformis	4575	5682	81%	0.0	88.00%	16840	NW_008941019.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold106011	Nicotiana tabacum	4573	5701	91%	0.0	84.76%	36765	NW_015791899.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold59123	Nicotiana tabacum	4573	5498	93%	0.0	84.79%	18644	NW_015921013.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4564	5734	97%	0.0	84.83%	265973	NW_017671664.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4553	5698	94%	0.0	84.73%	29083	NW_017677581.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold49463	Nicotiana sylvestris	4549	5045	86%	0.0	84.76%	96659	NW_00951568.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4547	10195	98%	0.0	84.76%	365158	NW_017671070.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold8326	Nicotiana tabacum	4545	4930	86%	0.0	84.70%	90194	NW_015941905.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold114830	Nicotiana tabacum	4543	5489	94%	0.0	84.71%	31234	NW_015798555.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4542	5456	94%	0.0	84.76%	394233	NW_017670965.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold84038	Nicotiana tabacum	4538	5549	94%	0.0	84.67%	26168	NW_015942552.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold17547	Nicotiana sylvestris	4538	5477	94%	0.0	84.69%	37565	NW_009422658.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold84013	Nicotiana sylvestris	4538	5544	94%	0.0	84.67%	44145	NW_009574958.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	4534	5978	84%	0.0	85.90%	294851	NW_017671447.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold100757	Nicotiana tabacum	4532	4930	84%	0.0	84.70%	69808	NW_015787832.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold3818	Nicotiana sylvestris	4532	6026	94%	0.0	84.62%	292384	NW_009524031.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold6229	Nicotiana sylvestris	4519	4962	86%	0.0	84.57%	131481	NW_009550820.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold30997	Nicotiana sylvestris	4516	5492	87%	0.0	84.57%	176597	NW_009516050.1

Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold21402	Nicotiana tabacum	4510	5335	84%	0.0	84.54%	194144	NW_015858587.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold2242	Nicotiana sylvestris	4508	5212	86%	0.0	84.59%	76533	NW_009476803.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4505	5744	92%	0.0	84.63%	258279	NW_017671719.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold4055	Nicotiana tabacum	4495	5490	87%	0.0	84.50%	318295	NW_015904028.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold32799	Nicotiana tabacum	4494	5965	94%	0.0	84.47%	283883	NW_015890680.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold2938	Nicotiana sylvestris	4494	5272	94%	0.0	84.44%	107624	NW_009514253.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold149308	Nicotiana tabacum	4488	6540	94%	0.0	87.12%	7509	NW_015822458.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold96186	Nicotiana tabacum	4486	5413	98%	0.0	83.40%	79153	NW_015952462.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold47678	Nicotiana tabacum	4484	4927	86%	0.0	84.44%	178459	NW_015910615.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4479	5461	86%	0.0	84.69%	125342	NW_017673660.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10980	Nicotiana tabacum	4475	5298	85%	0.0	84.42%	42835	NW_015794786.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold26004	Nicotiana sylvestris	4468	5524	87%	0.0	84.36%	136912	NW_009510504.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold56588	Nicotiana tabacum	4464	5623	93%	0.0	84.39%	155146	NW_015918745.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold61563	Nicotiana sylvestris	4464	5383	85%	0.0	84.38%	61634	NW_009550013.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold20881	Nicotiana tabacum	4462	7536	98%	0.0	88.15%	31722	NW_015856081.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold76247	Nicotiana tabacum	4460	5520	96%	0.0	83.32%	30427	NW_015935970.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold86760	Nicotiana tabacum	4460	5485	87%	0.0	84.36%	48685	NW_015944816.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold2332	Nicotiana tabacum	4458	6082	98%	0.0	84.31%	68855	NW_015867677.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1306	Nicotiana sylvestris	4458	6347	98%	0.0	84.31%	78631	NW_009372803.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4455	4799	83%	0.0	84.46%	329677	NW_017671242.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold520	Nicotiana tabacum	4449	5955	94%	0.0	84.32%	94142	NW_015914563.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold17173	Nicotiana tabacum	4447	5923	95%	0.0	84.40%	78275	NW_015836138.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold32380	Nicotiana sylvestris	4440	5052	84%	0.0	84.62%	37948	NW_009517588.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold6487	Nicotiana tabacum	4436	4836	86%	0.0	84.34%	62693	NW_015926126.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold11905	Nicotiana sylvestris	4436	5736	93%	0.0	84.29%	272020	NW_009359970.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4427	5802	94%	0.0	84.39%	1699785	NW_017670251.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold12348	Nicotiana sylvestris	4427	5839	89%	0.0	84.29%	72803	NW_009364892.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold17790	Nicotiana sylvestris	4420	5480	96%	0.0	83.20%	61339	NW_009425358.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold2835	Nicotiana sylvestris	4418	5196	86%	0.0	84.27%	250491	NW_009513109.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold122484	Nicotiana tabacum	4410	4901	81%	0.0	85.33%	58090	NW_015804127.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4409	6215	94%	0.0	84.96%	331877	NW_017671228.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold2103	Nicotiana tabacum	4407	5716	94%	0.0	84.18%	130931	NW_015856817.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold15836	Nicotiana sylvestris	4407	5732	94%	0.0	84.14%	63583	NW_009403647.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold33667	Nicotiana tabacum	4405	4824	82%	0.0	84.09%	14294	NW_015891508.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold3005	Nicotiana tabacum	4403	5294	87%	0.0	85.72%	197213	NW_015888088.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold35954	Nicotiana sylvestris	4403	5282	94%	0.0	84.15%	86977	NW_009521558.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold139625	Nicotiana tabacum	4397	5244	82%	0.0	85.08%	23219	NW_015816109.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold15027	Nicotiana sylvestris	4397	4906	87%	0.0	84.19%	57778	NW_009394659.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold53156	Nicotiana tabacum	4394	4983	84%	0.0	84.45%	74971	NW_015915632.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold26669	Nicotiana sylvestris	4392	4902	85%	0.0	84.15%	108786	NW_009511241.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold85842	Nicotiana tabacum	4386	5589	97%	0.0	84.15%	39521	NW_015944056.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold10873	Nicotiana sylvestris	4385	4898	85%	0.0	84.04%	99268	NW_009348503.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold54087	Nicotiana tabacum	4383	5130	94%	0.0	84.05%	63645	NW_015916486.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold57387	Nicotiana tabacum	4383	5750	89%	0.0	84.11%	74046	NW_015919484.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold6806	Nicotiana sylvestris	4383	5835	97%	0.0	84.13%	57418	NW_009557231.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold14575	Nicotiana tabacum	4379	5694	94%	0.0	84.04%	80269	NW_015820186.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold15474	Nicotiana tabacum	4379	4995	86%	0.0	84.02%	25850	NW_015825927.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold54984	Nicotiana tabacum	4379	4882	85%	0.0	84.02%	97540	NW_015917301.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold96016	Nicotiana sylvestris	4375	6212	98%	0.0	84.03%	52607	NW_009588294.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold8714	Nicotiana tomentosiformis	4372	5245	98%	0.0	84.02%	77865	NW_008973752.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4370	5474	90%	0.0	85.72%	26282	NW_017677889.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold14210	Nicotiana sylvestris	4366	5295	87%	0.0	84.03%	62761	NW_009385582.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold650	Nicotiana sylvestris	4362	5264	87%	0.0	85.56%	210552	NW_009553831.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold1786	Nicotiana sylvestris	4353	5658	94%	0.0	84.00%	351612	NW_009426135.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold18825	Nicotiana sylvestris	4353	4792	86%	0.0	84.02%	151572	NW_009436858.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9972	Nicotiana tabacum	4349	5234	87%	0.0	85.40%	114725	NW_015955243.1

Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold16107	Nicotiana tabacum	4342	4781	86%	0.0	83.98%	241788	NW_015829859.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold020930	Nicotiana sylvestris	4340	5705	91%	0.0	83.84%	107699	NW_009460248.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold21666	Nicotiana tabacum	4338	4711	84%	0.0	83.90%	105360	NW_015859855.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold18589	Nicotiana tabacum	4333	5181	98%	0.0	83.87%	81994	NW_015844059.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold9696	Nicotiana sylvestris	4331	4810	85%	0.0	83.84%	43153	NW_009589341.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4325	10187	98%	0.0	85.27%	227622	NW_017671996.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold397	Nicotiana sylvestris	4318	5195	89%	0.0	84.43%	113007	NW_009525719.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold24174	Nicotiana tabacum	4311	5470	94%	0.0	83.77%	54897	NW_015871679.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold10961	Nicotiana sylvestris	4311	5470	94%	0.0	83.77%	53229	NW_009349481.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold100015	Nicotiana tabacum	4296	5329	90%	0.0	84.45%	35908	NW_015787245.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold51229	Nicotiana sylvestris	4296	5329	90%	0.0	84.45%	35245	NW_009538531.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold27714	Nicotiana tabacum	4274	5016	89%	0.0	83.98%	81215	NW_015885890.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9746	Nicotiana tabacum	4274	5383	93%	0.0	83.69%	81695	NW_015953467.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold2459	Nicotiana sylvestris	4270	9365	83%	0.0	84.20%	174103	NW_009500913.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold17046	Nicotiana sylvestris	4263	5459	93%	0.0	83.65%	123686	NW_009417092.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold96428	Nicotiana tabacum	4253	4803	80%	0.0	84.66%	51920	NW_015952654.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold68100	Nicotiana tomentosiformis	4246	6431	93%	0.0	87.62%	22746	NW_008952598.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold9389	Nicotiana tabacum	4242	9326	83%	0.0	84.09%	98476	NW_015950629.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold637	Nicotiana sylvestris	4233	5077	85%	0.0	85.42%	476905	NW_009552386.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold43443	Nicotiana tabacum	4231	4758	82%	0.0	84.20%	90978	NW_015906720.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold29577	Nicotiana sylvestris	4231	5182	83%	0.0	84.18%	179029	NW_009514472.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold208	Nicotiana tabacum	4229	5203	92%	0.0	83.47%	140747	NW_015855661.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold101805	Nicotiana tabacum	4220	5633	95%	0.0	83.47%	55105	NW_015788649.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold224928	Nicotiana tabacum	4220	5111	85%	0.0	85.38%	20642	NW_015863778.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold23458	Nicotiana sylvestris	4220	5118	85%	0.0	85.43%	42416	NW_009488336.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold31891	Nicotiana sylvestris	4220	5633	95%	0.0	83.47%	186659	NW_009517044.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold188269	Nicotiana tabacum	4213	4776	81%	0.0	84.78%	10992	NW_015845379.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold15784	Nicotiana tabacum	4191	5533	82%	0.0	85.91%	37561	NW_015827850.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold37560	Nicotiana sylvestris	4180	5281	94%	0.0	84.36%	113441	NW_009523343.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold34345	Nicotiana tabacum	4176	4519	83%	0.0	83.36%	73502	NW_015892146.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold12672	Nicotiana sylvestris	4174	5581	87%	0.0	85.14%	236673	NW_009368492.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold44969	Nicotiana tomentosiformis	4169	5576	82%	0.0	85.81%	42854	NW_008926895.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4156	5071	88%	0.0	83.31%	61312	NW_017675714.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold45200	Nicotiana tabacum	4156	4977	89%	0.0	84.31%	112687	NW_015908326.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold17927	Nicotiana sylvestris	4141	4972	93%	0.0	83.14%	235894	NW_009426880.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold147251	Nicotiana tabacum	4139	4970	93%	0.0	83.15%	35449	NW_015821159.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4126	4986	80%	0.0	85.41%	328155	NW_017671251.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4106	5689	85%	0.0	85.90%	1677488	NW_017670254.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold25401	Nicotiana tabacum	4091	5075	87%	0.0	84.54%	26321	NW_015877164.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold53061	Nicotiana sylvestris	4089	4997	87%	0.0	83.69%	53828	NW_009540567.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4082	5712	88%	0.0	85.68%	303324	NW_017671393.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold16228	Nicotiana sylvestris	4061	4830	90%	0.0	82.97%	216634	NW_009408003.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4047	6530	84%	0.0	83.08%	81668	NW_017674894.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold13135	Nicotiana tabacum	4047	4796	85%	0.0	83.89%	63105	NW_015810438.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold119939	Nicotiana tabacum	4041	4808	81%	0.0	84.93%	61396	NW_015802274.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold8834	Nicotiana sylvestris	4041	4796	85%	0.0	83.86%	73830	NW_009579764.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold6101	Nicotiana tabacum	4037	12573	99%	0.0	86.91%	266780	NW_015922693.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold47887	Nicotiana sylvestris	4032	4855	85%	0.0	83.63%	27912	NW_009534816.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold51518	Nicotiana sylvestris	4017	4841	86%	0.0	84.39%	27608	NW_009538852.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	4013	4933	87%	0.0	82.88%	375176	NW_017671038.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold5272	Nicotiana tabacum	3999	6060	89%	0.0	87.97%	167441	NW_015915232.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	3988	4612	81%	0.0	83.70%	597674	NW_017670542.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	3958	5331	80%	0.0	86.64%	140783	NW_017673321.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold5750	Nicotiana tomentosiformis	3945	4872	89%	0.0	83.45%	256512	NW_008940819.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATT2	Nicotiana attenuata	3930	6175	80%	0.0	86.87%	474251	NW_017670736.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold33753	Nicotiana tabacum	3917	4850	89%	0.0	83.34%	102642	NW_015891589.1

Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3904	5814	80%	0.0	85.97%	232428	NW_017671953.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold25914	Nicotiana sylvestris	3904	5875	94%	0.0	82.42%	214657	NW_009510403.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold114081	Nicotiana tabacum	3897	5883	80%	0.0	87.36%	46885	NW_015798000.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold25823	Nicotiana tomentosiformis	3882	6838	89%	0.0	86.37%	49722	NW_008905623.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1069	Nicotiana tomentosiformis	3871	6871	96%	0.0	87.53%	170958	NW_008836159.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold1886	Nicotiana tabacum	3864	5996	82%	0.0	87.97%	13594	NW_015845550.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold15292	Nicotiana tomentosiformis	3858	7219	99%	0.0	85.79%	53616	NW_008887291.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold28562	Nicotiana sylvestris	3856	4995	84%	0.0	84.07%	32337	NW_009513345.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold105904	Nicotiana tabacum	3847	5770	94%	0.0	82.22%	107011	NW_015791814.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3823	4999	83%	0.0	85.22%	73401	NW_017675197.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold24479	Nicotiana sylvestris	3818	8059	91%	0.0	85.12%	88132	NW_009499680.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold26204	Nicotiana tabacum	3814	4704	86%	0.0	81.96%	48440	NW_015880953.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold7591	Nicotiana sylvestris	3808	5252	83%	0.0	84.47%	49741	NW_009565953.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold2415	Nicotiana tabacum	3807	7972	91%	0.0	85.07%	116635	NW_015871561.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold58463	Nicotiana tabacum	3807	5312	87%	0.0	85.57%	7176	NW_015920441.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold24327	Nicotiana sylvestris	3790	4757	81%	0.0	84.48%	95529	NW_009497992.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold21102	Nicotiana tabacum	3775	7024	95%	0.0	86.73%	173521	NW_015857153.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3773	7548	82%	0.0	86.38%	583767	NW_017670553.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4461	Nicotiana tomentosiformis	3771	4667	86%	0.0	81.80%	261601	NW_008926497.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3718	8573	97%	0.0	86.33%	458155	NW_017670783.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold91891	Nicotiana tabacum	3709	8989	96%	0.0	93.10%	69752	NW_015949033.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold10172	Nicotiana tomentosiformis	3705	7201	92%	0.0	86.93%	217121	NW_008830405.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold10728	Nicotiana tomentosiformis	3674	5153	98%	0.0	81.34%	57149	NW_008836582.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold20222	Nicotiana sylvestris	3672	4786	82%	0.0	85.06%	64345	NW_009452382.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold31826	Nicotiana tabacum	3659	5842	80%	0.0	86.81%	142986	NW_015889764.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold20963	Nicotiana tomentosiformis	3653	5038	94%	0.0	81.29%	277129	NW_008900223.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold26479	Nicotiana tabacum	3637	6310	90%	0.0	87.36%	91350	NW_015882277.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold33376	Nicotiana tabacum	3626	4915	94%	0.0	81.19%	329493	NW_015891234.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold36476	Nicotiana tabacum	3624	3976	84%	0.0	81.23%	68568	NW_015900242.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold267955	Nicotiana tabacum	3620	4245	82%	0.0	81.93%	11805	NW_015884172.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold37198	Nicotiana tomentosiformis	3618	4131	85%	0.0	81.15%	74187	NW_008918261.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold6869	Nicotiana tomentosiformis	3611	3968	84%	0.0	81.18%	120927	NW_008953251.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3609	6262	92%	0.0	86.80%	370296	NW_017671058.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3603	4746	80%	0.0	85.19%	212399	NW_017672182.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold12504	Nicotiana tomentosiformis	3598	6523	91%	0.0	87.15%	157552	NW_008856314.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold34924	Nicotiana tabacum	3581	4465	93%	0.0	81.01%	191410	NW_015896292.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3576	4019	84%	0.0	81.05%	348009	NW_017671147.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold32313	Nicotiana tomentosiformis	3576	4819	84%	0.0	84.31%	127009	NW_008912835.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold1608	Nicotiana tomentosiformis	3572	4099	81%	0.0	82.17%	309886	NW_008894797.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3555	5758	94%	0.0	86.19%	62422	NW_017675668.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3548	5507	88%	0.0	84.98%	149014	NW_017673143.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold321	Nicotiana tomentosiformis	3533	6689	86%	0.0	88.77%	225289	NW_008912597.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold21235	Nicotiana tabacum	3515	6653	94%	0.0	86.70%	115538	NW_015857775.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold50139	Nicotiana tabacum	3511	3592	83%	0.0	79.64%	44274	NW_015912870.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold23732	Nicotiana tabacum	3506	5606	83%	0.0	87.61%	42917	NW_015869612.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold2875	Nicotiana sylvestris	3506	3586	83%	0.0	79.62%	70400	NW_009513553.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3493	6533	95%	0.0	87.15%	378987	NW_017671024.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold4962	Nicotiana tomentosiformis	3482	5112	81%	0.0	85.93%	139036	NW_008932063.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold50627	Nicotiana tabacum	3474	5299	81%	0.0	85.89%	139157	NW_015913322.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold109268	Nicotiana tabacum	3465	5783	94%	0.0	85.02%	26584	NW_015794382.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold53732	Nicotiana sylvestris	3439	3675	80%	0.0	80.57%	59672	NW_009541312.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold48159	Nicotiana sylvestris	3432	5515	87%	0.0	84.86%	48782	NW_009535119.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold4884	Nicotiana sylvestris	3395	4117	88%	0.0	81.23%	208661	NW_009535875.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold8863	Nicotiana sylvestris	3393	5687	89%	0.0	86.11%	117359	NW_009580086.1
Tntom1	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold31810	Nicotiana tomentosiformis	3393	9515	96%	0.0	87.23%	139349	NW_008912276.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold61937	Nicotiana sylvestris	3391	9176	92%	0.0	85.11%	41847	NW_009550428.1

Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold35864	Nicotiana tabacum	3384	4084	88%	0.0	81.19%	130654	NW_015899669.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10658	Nicotiana tabacum	3360	7609	96%	0.0	87.79%	193269	NW_015792333.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold25055	Nicotiana tabacum	3328	6017	86%	0.0	87.62%	143344	NW_015875632.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3323	6219	97%	0.0	86.67%	397494	NW_017670957.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3308	5934	93%	0.0	87.29%	227578	NW_017671999.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3265	5793	96%	0.0	84.98%	150068	NW_017673125.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3253	6563	97%	0.0	85.58%	334748	NW_017671211.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3251	6117	97%	0.0	86.91%	232909	NW_017671950.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, NsyL_NsyL_scaffold32469	Nicotiana sylvestris	3206	5864	97%	0.0	84.20%	62904	NW_009517686.1
Tntom1	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, NsyL_NsyL_scaffold28367	Nicotiana sylvestris	3173	5357	97%	0.0	83.66%	70372	NW_009513128.1
Tntom1	refseq_genomes	8	Nicotiana attenuata strain UT unplaced genomic scaffold, NIATTr2	Nicotiana attenuata	3171	7821	94%	0.0	85.04%	504629	NW_017670677.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold96044	Nicotiana tabacum	3168	5346	97%	0.0	83.63%	17783	NW_015952350.1
Tntom1	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold10784	Nicotiana tabacum	3157	4970	87%	0.0	84.51%	58392	NW_015793290.1
GypsySL_11	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	11073	69437	100%	0.0	100.00%	98455869	NC_015438.3
GypsySL_11	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	8641	1,03E+08	100%	0.0	92.64%	56597135	NC_015448.3
GypsySL_11	refseq_genomes	8	Solanum pennellii chromosome 4, SPENNv200	Solanum pennellii	8567	33451	100%	0.0	92.39%	77197300	NC_028640.1
GypsySL_11	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	8272	67128	100%	0.0	91.58%	55977580	NC_015439.3
GypsySL_11	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	8117	72538	100%	0.0	91.06%	72906345	NC_015446.3
GypsySL_11	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	8082	36471	100%	0.0	91.00%	68126176	NC_015449.3
GypsySL_11	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	7771	26039	100%	0.0	90.18%	72290146	NC_015440.3
GypsySL_11	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	7744	35581	100%	0.0	90.02%	65987440	NC_015445.3
GypsySL_11	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	7662	42164	100%	0.0	90.38%	49794276	NC_015443.3
GypsySL_11	refseq_genomes	8	Solanum pennellii chromosome 12, SPENNv200	Solanum pennellii	7439	27066	100%	0.0	92.38%	83305730	NC_028648.1
GypsySL_11	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	7396	31289	100%	0.0	91.97%	65633393	NC_015447.3
GypsySL_11	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC0000008	Solanum lycopersicum	7260	9232	97%	0.0	91.53%	1577018	NW_020442489.1
GypsySL_11	refseq_genomes	8	Solanum pennellii chromosome 1, SPENNv200	Solanum pennellii	7167	27423	99%	0.0	91.44%	109333515	NC_028637.1
GypsySL_11	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	7105	42164	100%	0.0	91.25%	66557038	NC_015441.3
GypsySL_11	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC0000032	Solanum lycopersicum	7083	8988	97%	0.0	90.92%	62247	NW_020442513.1
GypsySL_11	refseq_genomes	8	Solanum pennellii chromosome 7, SPENNv200	Solanum pennellii	6390	49932	100%	0.0	93.15%	79292169	NC_028643.1
GypsySL_11	refseq_genomes	8	Solanum pennellii chromosome 10, SPENNv200	Solanum pennellii	5980	27144	98%	0.0	89.73%	82529941	NC_028646.1
GypsySL_11	refseq_genomes	8	Solanum pennellii chromosome 6, SPENNv200	Solanum pennellii	5452	7762	96%	0.0	87.84%	60730942	NC_028642.1
GypsySL_11	refseq_genomes	8	Solanum pennellii chromosome 5, SPENNv200	Solanum pennellii	5411	19097	90%	0.0	87.60%	77991103	NC_028641.1
GypsySL_11	refseq_genomes	8	Solanum pennellii chromosome 9, SPENNv200	Solanum pennellii	5094	18022	100%	0.0	92.39%	84057508	NC_028645.1
GypsySL_11	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	3934	23080	100%	0.0	90.88%	66723567	NC_015442.3
GypsySL_11	refseq_genomes	8	Solanum pennellii chromosome 2, SPENNv200	Solanum pennellii	3286	9084	88%	0.0	80.75%	59803892	NC_028638.1
GypsySL_11	refseq_genomes	8	Solanum pennellii chromosome 3, SPENNv200	Solanum pennellii	2662	8449	93%	0.0	77.69%	75414019	NC_028639.1
GypsySL_11	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC0000162	Solanum lycopersicum	2564	6292	82%	0.0	87.20%	7677	NW_020442643.1
GypsySL_11	refseq_genomes	8	Solanum pennellii chromosome 8, SPENNv200	Solanum pennellii	2470	7766	86%	0.0	76.91%	70546659	NC_028644.1
GypsySL_11	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	2451	11700	92%	0.0	77.11%	68175699	NC_015444.3
GypsySL_11	refseq_genomes	8	Solanum pennellii chromosome 11, SPENNv200	Solanum pennellii	2255	5737	80%	0.0	75.79%	66223686	NC_028647.1
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	16281	3,51E+09	100%	0.0	100.00%	72290146	NC_015440.3
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	11029	2,19E+09	100%	0.0	89.29%	66557038	NC_015441.3
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	10892	3,61E+09	100%	0.0	89.05%	72906345	NC_015446.3
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	10829	3,81E+09	100%	0.0	88.89%	65633393	NC_015447.3
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	10384	3,67E+09	100%	0.0	87.98%	68126176	NC_015449.3
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	10294	3,15E+09	100%	0.0	87.90%	56597135	NC_015448.3
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	10275	1,61E+09	100%	0.0	87.74%	49794276	NC_015443.3
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	10198	3,55E+09	100%	0.0	90.04%	68175699	NC_015444.3
GypsySL_01	refseq_genomes	8	Solanum pennellii chromosome 12, SPENNv200	Solanum pennellii	10137	1,63E+09	100%	0.0	89.38%	83305730	NC_028648.1
GypsySL_01	refseq_genomes	8	Solanum pennellii chromosome 5, SPENNv200	Solanum pennellii	10131	1,38E+09	100%	0.0	90.06%	77991103	NC_028641.1
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	10028	3,68E+09	100%	0.0	91.89%	65987440	NC_015445.3
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	10024	3,19E+09	100%	0.0	91.92%	66723567	NC_015442.3
GypsySL_01	refseq_genomes	8	Solanum pennellii chromosome 10, SPENNv200	Solanum pennellii	9963	1,57E+09	100%	0.0	89.64%	82529941	NC_028646.1
GypsySL_01	refseq_genomes	8	Solanum pennellii chromosome 1, SPENNv200	Solanum pennellii	9958	2,45E+09	100%	0.0	91.81%	109333515	NC_028637.1
GypsySL_01	refseq_genomes	8	Solanum pennellii chromosome 4, SPENNv200	Solanum pennellii	9721	1,43E+09	100%	0.0	91.13%	77197300	NC_028640.1
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	9686	1,48E+09	100%	0.0	89.03%	55977580	NC_015439.3

GypsySL_01	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	9603	9,45E+08	100%	0.0	91.11%	60730942	NC_028642.1
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	9570	4,76E+09	100%	0.0	90.77%	98455869	NC_015438.3
GypsySL_01	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	Solanum pennellii	9520	1,13E+09	100%	0.0	90.87%	70546659	NC_028644.1
GypsySL_01	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	Solanum pennellii	9295	1,72E+09	100%	0.0	90.57%	84057508	NC_028645.1
GypsySL_01	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	Solanum pennellii	9282	1,66E+09	100%	0.0	90.34%	79292169	NC_028643.1
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC0000008	Solanum lycopersicum	8708	52247	100%	0.0	90.89%	1577018	NW_020442489.1
GypsySL_01	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	8586	9,58E+08	100%	0.0	91.97%	75414019	NC_028639.1
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC0000003	Solanum lycopersicum	8486	1,02E+08	98%	0.0	91.44%	1066869	NW_020442484.1
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC0000005	Solanum lycopersicum	8144	53201	100%	0.0	90.51%	86546	NW_020442486.1
GypsySL_01	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	Solanum pennellii	8137	1,32E+09	100%	0.0	88.33%	66223686	NC_028647.1
GypsySL_01	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	7520	4,76E+08	100%	0.0	89.94%	59803892	NC_028638.1
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC0000004	Solanum lycopersicum	6955	20010	97%	0.0	90.11%	27484	NW_020442485.1
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC0000001	Solanum lycopersicum	6499	60770	97%	0.0	86.39%	2014460	NW_020442482.1
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC0000009	Solanum lycopersicum	5694	26130	96%	0.0	91.05%	797759	NW_020442490.1
GypsySL_01	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00164	Solanum tuberosum	5267	6527	81%	0.0	84.45%	1312146	NW_006239092.1
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC0000023	Solanum lycopersicum	5195	12928	97%	0.0	88.85%	89524	NW_020442504.1
GypsySL_01	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC0000006	Solanum lycopersicum	3620	54027	99%	0.0	86.22%	784488	NW_020442487.1
GypsySL_03	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	18912	6,22E+08	100%	0.0	100.00%	98455869	NC_015438.3
GypsySL_03	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	12168	2,03E+08	100%	0.0	94.06%	77197300	NC_028640.1
GypsySL_03	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	Solanum pennellii	11284	2,63E+08	89%	0.0	95.19%	79292169	NC_028643.1
GypsySL_03	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum pennellii	11143	2,90E+08	100%	0.0	94.71%	77991103	NC_028641.1
GypsySL_03	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	10104	4,32E+08	100%	0.0	94.92%	10933351	NC_028637.1
GypsySL_03	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	Solanum pennellii	10986	2,65E+08	92%	0.0	94.36%	83305730	NC_028648.1
GypsySL_03	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	10608	2,79E+08	91%	0.0	93.58%	65633393	NC_015447.3
GypsySL_03	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	10418	2,57E+08	90%	0.0	93.41%	65987440	NC_015445.3
GypsySL_03	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	10390	2,58E+08	91%	0.0	92.81%	72906345	NC_015446.3
GypsySL_03	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	10379	2,67E+08	90%	0.0	93.29%	68175699	NC_015444.3
GypsySL_03	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	10368	3,11E+08	92%	0.0	93.23%	56597135	NC_015448.3
GypsySL_03	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	10364	1,82E+08	92%	0.0	93.34%	55977580	NC_015439.3
GypsySL_03	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum pennellii	10353	2,39E+08	92%	0.0	96.30%	82529941	NC_028646.1
GypsySL_03	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	10353	2,60E+08	100%	0.0	92.27%	66723567	NC_015442.3
GypsySL_03	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	Solanum pennellii	10342	2,92E+08	92%	0.0	93.19%	84057508	NC_028645.1
GypsySL_03	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	10202	2,66E+08	91%	0.0	92.77%	72290146	NC_015440.3
GypsySL_03	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	10129	2,03E+08	92%	0.0	95.41%	75414019	NC_028639.1
GypsySL_03	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	10098	92117	88%	0.0	92.27%	49794276	NC_015443.3
GypsySL_03	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	Solanum pennellii	10094	2,17E+08	92%	0.0	92.03%	66223686	NC_028647.1
GypsySL_03	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	9821	2,55E+08	92%	0.0	90.96%	68126176	NC_015449.3
GypsySL_03	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	Solanum pennellii	9710	92233	89%	0.0	91.37%	70546659	NC_028644.1
GypsySL_03	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	9683	1,76E+08	88%	0.0	91.55%	66557038	NC_015441.3
GypsySL_03	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	8665	1,32E+08	89%	0.0	91.34%	60730942	NC_028642.1
GypsySL_03	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	8395	1,24E+08	91%	0.0	96.69%	59803892	NC_028638.1
GypsySL_03	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00030	Solanum tuberosum	5517	10655	80%	0.0	88.54%	2778153	NW_006238956.1
GypsySL_03	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC0000001	Solanum lycopersicum	4787	16509	85%	0.0	85.88%	2014460	NW_020442482.1
GypsySL_03	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00057	Solanum tuberosum	4405	13569	86%	0.0	85.99%	2187528	NW_006238984.1
GypsySL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	21420	1,24E+09	100%	0.0	100.00%	72290146	NC_015440.3
GypsySL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	20831	1,15E+09	100%	0.0	99.04%	55977580	NC_015439.3
GypsySL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	14763	6,56E+08	100%	0.0	98.65%	49794276	NC_015443.3
GypsySL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	14307	1,39E+09	100%	0.0	99.07%	98455869	NC_015438.3
GypsySL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	14235	9,28E+08	100%	0.0	97.35%	68175699	NC_015444.3
GypsySL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	13782	9,44E+08	100%	0.0	97.86%	65987440	NC_015445.3
GypsySL_05	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	13415	1,49E+09	100%	0.0	96.95%	77197300	NC_028640.1
GypsySL_05	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	13332	1,11E+09	100%	0.0	97.01%	60730942	NC_028642.1
GypsySL_05	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	Solanum pennellii	13309	1,83E+09	100%	0.0	96.96%	82529941	NC_028646.1
GypsySL_05	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum pennellii	12920	1,19E+09	100%	0.0	96.66%	59803892	NC_028638.1
GypsySL_05	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum lycopersicum	11723	9,74E+08	100%	0.0	92.16%	68126176	NC_015449.3

GypsySL_05	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	11105	2,06E+09	100%	0.0	93.21%	75414019	NC_028639.1
GypsySL_05	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	Solanum pennellii	10929	1,65E+09	99%	0.0	93.02%	70546659	NC_028644.1
GypsySL_05	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	Solanum pennellii	10656	1,27E+09	99%	0.0	94.07%	66223686	NC_028647.1
GypsySL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	9660	9,80E+08	100%	0.0	90.64%	66557038	NC_015441.3
GypsySL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.0SC00000003	Solanum lycopersicum	9361	25180	97%	0.0	89.32%	1066869	NW_020442484.1
GypsySL_05	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	8772	1,93E+09	100%	0.0	90.31%	109333515	NC_028637.1
GypsySL_05	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	Solanum pennellii	8167	1,66E+09	99%	0.0	87.92%	79292169	NC_028643.1
GypsySL_05	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	Solanum pennellii	7393	1,57E+09	99%	0.0	86.58%	83305730	NC_028648.1
GypsySL_05	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum pennellii	7184	1,60E+09	100%	0.0	89.34%	77991103	NC_028641.1
GypsySL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	6281	9,75E+08	100%	0.0	86.68%	65633393	NC_015447.3
GypsySL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	6209	8,79E+08	100%	0.0	86.37%	66723567	NC_015442.3
GypsySL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	6067	1,11E+09	100%	0.0	85.92%	72906345	NC_015446.3
GypsySL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	6013	6,20E+08	100%	0.0	85.96%	56597135	NC_015448.3
GypsySL_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	20247	63174	100%	0.0	100.00%	56597135	NC_015448.3
GypsySL_07	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	17692	2,15E+08	100%	0.0	95.83%	109333515	NC_028637.1
GypsySL_07	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	17680	1,90E+08	100%	0.0	95.81%	75414019	NC_028639.1
GypsySL_07	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	17680	1,11E+08	100%	0.0	95.81%	77197300	NC_028640.1
GypsySL_07	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum pennellii	17680	1,13E+08	100%	0.0	95.81%	77991103	NC_028641.1
GypsySL_07	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	Solanum pennellii	17680	73938	100%	0.0	95.81%	70546659	NC_028644.1
GypsySL_07	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum pennellii	17680	72647	100%	0.0	95.81%	82529941	NC_028646.1
GypsySL_07	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	Solanum pennellii	17675	1,54E+08	100%	0.0	95.80%	66223686	NC_028647.1
GypsySL_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	16235	1,50E+08	100%	0.0	95.31%	65633393	NC_015447.3
GypsySL_07	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	16211	1,45E+08	100%	0.0	95.70%	59803892	NC_028638.1
GypsySL_07	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	Solanum pennellii	15605	85510	100%	0.0	95.77%	84057508	NC_028645.1
GypsySL_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	12981	94041	100%	0.0	96.05%	72906345	NC_015446.3
GypsySL_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	12977	1,56E+08	100%	0.0	96.00%	98455869	NC_015438.3
GypsySL_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	11995	50746	100%	0.0	93.68%	66723567	NC_015442.3
GypsySL_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	11985	65880	100%	0.0	93.69%	49794276	NC_015443.3
GypsySL_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	11736	80852	100%	0.0	95.83%	68126176	NC_015449.3
GypsySL_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	11559	86829	100%	0.0	92.70%	66557038	NC_015441.3
GypsySL_07	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	Solanum pennellii	9642	57595	100%	0.0	88.45%	79292169	NC_028643.1
GypsySL_07	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	9461	66115	100%	0.0	93.70%	60730942	NC_028642.1
GypsySL_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	8340	38110	100%	0.0	85.56%	68175699	NC_015444.3
GypsySL_07	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	Solanum pennellii	8287	1,01E+08	100%	0.0	89.12%	83305730	NC_028648.1
GypsySL_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	7801	78861	100%	0.0	88.22%	72290146	NC_015440.3
GypsySL_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	6152	47779	100%	0.0	92.04%	65987440	NC_015445.3
GypsySL_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00218	Solanum tuberosum	5576	25175	92%	0.0	81.98%	1093294	NW_006239144.1
GypsySL_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00440	Solanum tuberosum	5094	42806	90%	0.0	83.26%	479509	NW_006239363.1
GypsySL_07	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	4855	1,32E+08	99%	0.0	96.50%	55977580	NC_015439.3
GypsySL_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00479	Solanum tuberosum	4682	11021	93%	0.0	87.34%	406342	NW_006239404.1
GypsySL_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00378	Solanum tuberosum	4397	11664	89%	0.0	85.30%	585542	NW_006239300.1
GypsySL_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00113	Solanum tuberosum	4010	9996	92%	0.0	80.53%	1648986	NW_006239039.1
GypsySL_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00208	Solanum tuberosum	3940	9757	86%	0.0	87.91%	1133327	NW_006239132.1
GypsySL_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00157	Solanum tuberosum	3319	8178	89%	0.0	82.08%	1381298	NW_006239084.1
GypsySL_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00009	Solanum tuberosum	2952	21898	90%	0.0	85.17%	3955149	NW_006238935.1
GypsySL_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00585	Solanum tuberosum	2841	28056	89%	0.0	86.83%	263234	NW_006239524.1
GypsySL_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00085	Solanum tuberosum	2808	12037	88%	0.0	85.08%	1839851	NW_006239014.1
GypsySL_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00163	Solanum tuberosum	2479	19962	98%	0.0	85.53%	1340955	NW_006239087.1
GypsySL_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00063	Solanum tuberosum	2433	20394	96%	0.0	87.84%	2166112	NW_006238988.1
GypsySL_07	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00146	Solanum tuberosum	2174	10066	86%	0.0	83.18%	1428744	NW_006239076.1
ToRTL1/CopiaSL_19	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	9502	1,38E+08	95%	0.0	99.54%	55977580	NC_015439.3
ToRTL1/CopiaSL_19	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	8696	1,53E+08	99%	0.0	95.30%	59803892	NC_028638.1
CopiaSL_21	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	8434	1,04E+08	100%	0.0	100.00%	72906345	NC_015446.3
CopiaSL_21	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum pennellii	7456	1,13E+08	100%	0.0	96.13%	77991103	NC_028641.1
CopiaSL_21	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum pennellii	7404	2,19E+08	100%	0.0	95.93%	82529941	NC_028646.1
CopiaSL_21	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	Solanum pennellii	7332	1,61E+08	100%	0.0	95.67%	79292169	NC_028643.1

CopiaSL_21	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	7330	2,79E+08	100%	0.0	95.65%	109333515	NC_028637.1
CopiaSL_21	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	Solanum pennellii	7304	1,53E+08	100%	0.0	95.54%	83305730	NC_028648.1
CopiaSL_21	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	7169	95886	100%	0.0	95.10%	75414019	NC_028639.1
CopiaSL_21	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	Solanum pennellii	7142	91957	100%	0.0	94.91%	66223686	NC_028647.1
CopiaSL_21	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	7123	57594	100%	0.0	94.91%	59803892	NC_028638.1
CopiaSL_21	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	7105	46477	100%	0.0	94.75%	60730942	NC_028642.1
CopiaSL_21	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	7062	1,21E+08	100%	0.0	94.60%	68175699	NC_015444.3
CopiaSL_21	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	Solanum pennellii	7022	59274	100%	0.0	94.68%	70546659	NC_028644.1
CopiaSL_21	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	Solanum pennellii	6988	1,35E+08	100%	0.0	94.32%	84057508	NC_028645.1
CopiaSL_21	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	6922	64242	100%	0.0	94.11%	56597135	NC_015448.3
CopiaSL_21	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	6892	1,70E+08	100%	0.0	93.92%	98455869	NC_015438.3
CopiaSL_21	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	6892	45782	100%	0.0	93.96%	55977580	NC_015439.3
CopiaSL_21	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	6807	1,75E+08	100%	0.0	93.62%	77197300	NC_028640.1
CopiaSL_21	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	6774	2,19E+08	100%	0.0	93.48%	65633393	NC_015447.3
CopiaSL_21	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	6663	92189	100%	0.0	93.06%	68126176	NC_015449.3
CopiaSL_21	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	Solanum lycopersicum	6503	1,38E+08	100%	0.0	92.41%	66557038	NC_015441.3
CopiaSL_21	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	6416	40623	100%	0.0	92.85%	49794276	NC_015443.3
CopiaSL_21	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	6220	49772	100%	0.0	91.30%	65987440	NC_015445.3
CopiaSL_21	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	6133	64900	100%	0.0	91.04%	72290146	NC_015440.3
CopiaSL_21	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00211	Solanum tuberosum	5864	6344	100%	0.0	90.48%	1109151	NW_006239137.1
CopiaSL_21	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	5677	65153	100%	0.0	93.49%	66723567	NC_015442.3
CopiaSL_21	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC00000003	Solanum lycopersicum	5332	5956	99%	0.0	87.92%	1066869	NW_020442484.1
CopiaSL_21	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 unplaced genomic scaffold, SL3.0 SL3.00SC00000008	Solanum lycopersicum	5313	5313	99%	0.0	87.76%	1577018	NW_020442489.1
CopiaSL_21	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00038	Solanum tuberosum	5278	5278	100%	0.0	87.66%	2533956	NW_006238966.1
CopiaSL_21	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00142	Solanum tuberosum	4710	5545	98%	0.0	89.17%	1452299	NW_006239067.1
CopiaSL_21	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00492	Solanum tuberosum	4650	6635	100%	0.0	90.41%	395864	NW_006239410.1
CopiaSL_21	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00031	Solanum tuberosum	4108	5686	96%	0.0	90.37%	2735125	NW_006238957.1
CopiaSL_21	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf01051	Solanum tuberosum	3760	5603	100%	0.0	87.92%	20269	NW_006240081.1
CopiaSL_21	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00003	Solanum tuberosum	2418	4571	94%	0.0	85.59%	4590222	NW_006238929.1
CopiaSL_21	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00236	Solanum tuberosum	2359	9870	82%	0.0	89.01%	1037213	NW_006239160.1
CopiaSL_21	refseq_genomes	8	Nicotiana attenuata strain Ut chromosome 10, NIATT2	Nicotiana attenuata	1347	2830	90%	0.0	81.59%	51082206	NC_031998.1
CopiaSL_23	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	8841	26677	100%	0.0	100.00%	55977580	NC_015439.3
CopiaSL_23	refseq_genomes	8	Solanum pennellii chromosome 1, SPENN200	Solanum pennellii	8536	48827	100%	0.0	98.87%	109333515	NC_028637.1
CopiaSL_23	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum pennellii	8530	31527	100%	0.0	98.85%	77991103	NC_028641.1
CopiaSL_23	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	Solanum pennellii	8527	11977	100%	0.0	98.83%	60730942	NC_028642.1
CopiaSL_23	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 5, SL3.0	Solanum lycopersicum	8167	24191	100%	0.0	97.47%	66723567	NC_015442.3
CopiaSL_23	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	Solanum lycopersicum	8133	25705	100%	0.0	97.37%	56597135	NC_015448.3
CopiaSL_23	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	8052	43899	100%	0.0	97.32%	98455869	NC_015438.3
CopiaSL_23	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	Solanum pennellii	7830	17918	100%	0.0	96.22%	66223686	NC_028647.1
CopiaSL_23	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	Solanum pennellii	7598	26905	100%	0.0	95.36%	83305730	NC_028648.1
CopiaSL_23	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 9, SL3.0	Solanum lycopersicum	7500	22384	100%	0.0	94.97%	72906345	NC_015446.3
CopiaSL_23	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	7393	27493	100%	0.0	97.66%	65987440	NC_015445.3
CopiaSL_23	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	7391	16177	100%	0.0	94.54%	49794276	NC_015443.3
CopiaSL_23	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum lycopersicum	7138	36495	100%	0.0	93.66%	68126176	NC_015449.3
CopiaSL_23	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	6303	18257	100%	0.0	98.15%	68175699	NC_015444.3
CopiaSL_23	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	5987	27011	100%	0.0	95.55%	65633393	NC_015447.3
CopiaSL_23	refseq_genomes	8	Solanum pennellii chromosome 2, SPENN200	Solanum pennellii	4366	12982	98%	0.0	96.82%	59803892	NC_028638.1
CopiaSL_23	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum pennellii	3637	25874	97%	0.0	82.67%	82529941	NC_028646.1
CopiaSL_23	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00091	Solanum tuberosum	3269	3269	90%	0.0	80.56%	1834259	NW_006239016.1
CopiaSL_23	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00708	Solanum tuberosum	3079	3079	87%	0.0	80.12%	177089	NW_006239630.1
CopiaSL_23	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00329	Solanum tuberosum	3048	3048	86%	0.0	80.18%	703406	NW_006239250.1
CopiaSL_23	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00143	Solanum tuberosum	3035	3035	96%	0.0	78.87%	1446147	NW_006239068.1
CopiaSL_23	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 10, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2988	11832	86%	0.0	80.00%	205736368	NC_029986.1
CopiaSL_23	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii	2830	23790	100%	0.0	95.44%	75414019	NC_028639.1
CopiaSL_23	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 9, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2769	5221	86%	0.0	78.99%	238794889	NC_029985.1
CopiaSL_23	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00289	Solanum tuberosum	2724	5087	88%	0.0	87.47%	799419	NW_006239225.1

CopiaSL_23	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 4, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2706	6307	86%	0.0	78.80%	215701946	NC_029980.1
CopiaSL_23	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 12, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2619	2672	85%	0.0	78.34%	229934170	NC_029988.1
CopiaSL_23	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00009	Solanum tuberosum	2579	4831	87%	0.0	86.64%	3955149	NW_006238935.1
CopiaSL_23	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00220	Solanum tuberosum	2545	4821	88%	0.0	86.51%	1072888	NW_006239148.1
CopiaSL_23	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 5, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	2499	7068	89%	0.0	77.95%	217274494	NC_029981.1
CopiaSL_23	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	Solanum pennellii	2495	25774	100%	0.0	83.98%	84057508	NC_028645.1
CopiaSL_23	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00123	Solanum tuberosum	2300	9504	89%	0.0	84.44%	1602102	NW_006239048.1
CopiaSL_23	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	Solanum pennellii	2152	8755	98%	0.0	95.36%	79292169	NC_028643.1
CopiaSL_23	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii	2128	18657	86%	0.0	83.48%	77197300	NC_028640.1
CopiaSL_23	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	Solanum pennellii	2030	11284	85%	0.0	82.47%	70546659	NC_028644.1
CopiaSL_23	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 11, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	1683	8112	84%	0.0	76.42%	220335243	NC_029987.1
CopiaSL_23	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00069	Solanum tuberosum	1668	4970	90%	0.0	82.01%	2089258	NW_006238995.1
CopiaSL_23	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold20247	Nicotiana tabacum	1664	2262	83%	0.0	79.67%	62189	NW_015852854.1
CopiaSL_23	refseq_genomes	8	Nicotiana sylvestris unplaced genomic scaffold, Nsyl Nsyl_scaffold25512	Nicotiana sylvestris	1664	2247	83%	0.0	79.66%	20976	NW_009509957.1
CopiaSL_23	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00001	Solanum tuberosum	1646	4456	82%	0.0	88.29%	7090834	NW_006238927.1
CopiaSL_23	refseq_genomes	8	Nicotiana attenuata strain UT chromosome 1, NIATT2	Nicotiana attenuata	1618	3953	85%	0.0	83.66%	110538649	NC_031989.1
CopiaSL_23	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00033	Solanum tuberosum	1609	4139	86%	0.0	75.69%	2695005	NW_006238959.1
CopiaSL_23	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00187	Solanum tuberosum	1557	6376	88%	0.0	80.65%	1214092	NW_006239111.1
CopiaSL_23	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00296	Solanum tuberosum	1519	3309	88%	0.0	82.31%	820454	NW_006239217.1
CopiaSL_23	refseq_genomes	8	Nicotiana tomentosiformis unplaced genomic scaffold, Ntom_v01 Ntom_scaffold54762	Nicotiana tomentosiformis	1413	1891	80%	0.0	77.95%	36099	NW_008937777.1
CopiaSL_23	refseq_genomes	8	Nicotiana tabacum cultivar TN90 unplaced genomic scaffold, Ntab-TN90 Ntab-TN90_scaffold36594	Nicotiana tabacum	1402	1874	80%	0.0	77.86%	64966	NW_015900355.1
CopiaSL_23	refseq_genomes	8	Capsicum annuum cultivar Zunla-1 chromosome 7, Pepper Zunla 1 Ref_v1.0	Capsicum annuum	1048	4213	84%	0.0	83.67%	222112641	NC_029983.1
CopiaSL_15	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 1, SL3.0	Solanum lycopersicum	10030	35643	100%	0.0	100.00%	98455869	NC_015438.3
CopiaSL_15	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 2, SL3.0	Solanum lycopersicum	8850	11061	100%	0.0	96.12%	55977580	NC_015439.3
CopiaSL_15	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	Solanum lycopersicum	7812	23512	100%	0.0	92.73%	72290146	NC_015440.3
CopiaSL_15	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	Solanum lycopersicum	7810	26612	100%	0.0	92.77%	68175699	NC_015444.3
CopiaSL_15	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 10, SL3.0	Solanum lycopersicum	7755	8331	100%	0.0	92.59%	65633393	NC_015447.3
CopiaSL_15	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	Solanum lycopersicum	7513	14654	98%	0.0	92.05%	49794276	NC_015443.3
CopiaSL_15	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	Solanum lycopersicum	7406	15920	100%	0.0	91.45%	65987440	NC_015445.3
CopiaSL_15	refseq_genomes	8	Solanum pennellii chromosome 4, SPENN200	Solanum pennellii Solanum	7090	19069	99%	0.0	90.61%	77197300	NC_028640.1
CopiaSL_15	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	pennellii Solanum	7005	13764	98%	0.0	91.61%	60730942	NC_028642.1
CopiaSL_15	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	lycopersicum Solanum	5786	6569	93%	0.0	87.40%	56597135	NC_015448.3
CopiaSL_15	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	pennellii Solanum	5631	17970	82%	0.0	94.41%	84057508	NC_028649.3
CopiaSL_15	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	lycopersicum Solanum	5469	15492	93%	0.0	93.16%	68126176	NC_015449.3
CopiaSL_15	refseq_genomes	8	Solanum pennellii chromosome 7, SPENN200	pennellii Solanum pennellii	5299	11854	87%	0.0	86.87%	79292169	NC_028643.1
CopiaSL_15	refseq_genomes	8	Solanum pennellii chromosome 3, SPENN200	Solanum pennellii Solanum	5234	11825	98%	0.0	93.54%	75414019	NC_028639.1
CopiaSL_15	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	lycopersicum Solanum	4977	5787	83%	0.0	86.87%	66223686	NC_028647.1
CopiaSL_15	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 4, SL3.0	pennellii Solanum	3450	9017	96%	0.0	87.42%	66557038	NC_015441.3
CopiaSL_15	refseq_genomes	8	Solanum pennellii chromosome 12, SPENN200	tuberosum Solanum	2617	4849	85%	0.0	83.03%	83305730	NC_028648.1
CopiaSL_15	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00165	tuberosum Solanum	1435	3689	82%	0.0	81.73%	1320774	NW_006239090.1
CopiaSL_15	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00116	tuberosum Solanum	1380	3799	81%	0.0	82.42%	1639857	NW_006239040.1
CopiaSL_15	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00022	lycopersicum Solanum	1182	3157	81%	0.0	79.21%	3032639	NW_006238948.1
CopiaSL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 11, SL3.0	pennellii Solanum	9806	16013	100%	0.0	100.00%	56597135	NC_015448.3
CopiaSL_05	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	lycopersicum Solanum	8545	9409	100%	0.0	95.67%	60730942	NC_028642.1
CopiaSL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	pennellii Solanum pennellii	8482	14964	100%	0.0	95.49%	49794276	NC_015443.3
CopiaSL_05	refseq_genomes	8	Solanum pennellii chromosome 5, SPENN200	Solanum lycopersicum	7930	20443	100%	0.0	93.59%	77991103	NC_028641.1
CopiaSL_05	refseq_genomes	8	Solanum pennellii chromosome 10, SPENN200	Solanum lycopersicum	7723	8921	99%	0.0	93.00%	82529941	NC_028646.1
CopiaSL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 12, SL3.0	Solanum pennellii Solanum	7099	17630	100%	0.0	90.95%	68126176	NC_015449.3
CopiaSL_05	refseq_genomes	8	Solanum pennellii chromosome 6, SPENN200	pennellii Solanum	7083	7083	100%	0.0	90.80%	65633393	NC_015447.3
CopiaSL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 6, SL3.0	lycopersicum Solanum	6999	14795	100%	0.0	90.49%	70546659	NC_028644.1
CopiaSL_05	refseq_genomes	8	Solanum pennellii chromosome 8, SPENN200	lycopersicum Solanum	6894	7369	97%	0.0	90.64%	75414019	NC_028639.1
CopiaSL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 8, SL3.0	pennellii Solanum	6820	11787	95%	0.0	91.21%	65987440	NC_015445.3
CopiaSL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 3, SL3.0	lycopersicum Solanum	6457	17879	99%	0.0	91.88%	72290146	NC_015440.3
CopiaSL_05	refseq_genomes	8	Solanum pennellii chromosome 11, SPENN200	pennellii	6388	8781	99%	0.0	91.28%	66223686	NC_028647.1
CopiaSL_05	refseq_genomes	8	Solanum lycopersicum cultivar Heinz 1706 chromosome 7, SL3.0	5755	9546	98%	0.0	97.30%	68175699	NC_015444.3	
CopiaSL_05	refseq_genomes	8	Solanum pennellii chromosome 9, SPENN200	5424	6403	94%	0.0	90.11%	84057508	NC_028645.1	

CopiaSL_05	refseq_genomes	8	Solanum pennellii chromosome 4, SPENNv200	Solanum pennellii	4440	10564	98%	0.0	85.98%	77197300	NC_028640.1
CopiaSL_05	refseq_genomes	8	Solanum pennellii chromosome 2, SPENNv200	Solanum pennellii	3430	8149	92%	0.0	93.72%	59803892	NC_028638.1
CopiaSL_05	refseq_genomes	8	Solanum tuberosum cultivar DM 1-3 516 R44 unplaced genomic scaffold, SolTub_3.0 scf00577	Solanum tuberosum	3195	3984	82%	0.0	82.83%	293230	NW_006239491.1