Supplementary data

Supplementary methods.

Samples were composed by three fruit typically with 12-15g in total aliquots and processed with mortar and pestle. The pH parameter was determined with a pHmeter (Hanna Edge®). To determine the total soluble solids (TSS) content, a few drops of tomato juice were placed in a refractometer (Milwaukee MA871, Rocky Mount, USA) and expressed in °Brix, Total titratable acidity (TTA) was titrimetrically determined with a 0.1 mol L⁻¹ NaOH solution until reaching a pH of 8.2 (AOAC, 1980). TTA was expressed in g of citric acid per kg of fruit fresh weight.

Supplementary table 1. pH, TSS (total soluble solids), TTA (total titratable acidity) and RATIO (TSS/TTA) in control, 20 mM ascorbic acid and 10 mM H₂O₂ treatments.

		pН			TSS (°Brix)			TTA (g citric acid kg ⁻¹ FW)			Ratio (TSS/TTA)		
		Control	AA	H_2O_2	Control	AA	H_2O_2	Control	AA	H_2O_2	Control	AA	H_2O_2
WT	MG	$4.3 \pm 0.2 \text{ aA1}$	$4.2 \pm 0.2 \text{ aA1}$	$4.3 \pm 0.1 \text{ aA1}$	$4.5\pm0.2~aA1$	$4.7 \pm 0.4 \text{ aA1}$	$4.9 \pm 0.7 \text{ aA1}$	$4.7 \pm 0.6 \text{ aA1}$	$3.6\pm0.8~\text{aA1}$	$2.9 \pm 0.2 \; aA1$	$12.8 \pm 1.2 \text{ aA2}$	$10.7 \pm 0.8 \text{ aA}1$	$13.1 \pm 0.3 \text{ aA1}$
	BR	$3.9 \pm 0.1 \; aA1$	$3.9 \pm 0.1 \; aA1$	$3.9 \pm 0.1 \; aA1$	$5.6 \pm 0.3 \ aA2$	$6.3\pm1.4~aA2$	$6.1\pm1.4~aA1$	$6,9.0\pm0.2~bA2$	$6.6\pm0.7\;bA2$	$6.5\pm0.2\;bA2$	$8.0 \pm 0.4 \ aA1$	$9.6 \pm 1.2 \; aA1$	$10.3\pm1.4~aA1$
	OR	$3.9 \pm 0.1 \; aA1$	$4.0 \pm 0.0 \ aA1$	$4.0 \pm 0.1 \; aA1$	$5.8 \pm 0.6 \; aA2$	$6.7\pm1.2~aA23$	$6.3 \pm 2.1 \text{ aA1}$	$7.9 \pm 0.7 \; bA2$	$6.2\pm0.5\;bA2$	$6.6 \pm 0.6 \; aA2$	$7.2\pm1.7~aA1$	$8.3\pm1.4~aA1$	$11.5\pm1.9~aA1$
	RR	$3.9 \pm 0.1 \; aA1$	$3.9 \pm 0.1 \; aA1$	$3.8 \pm 0.0 \ aA1$	$7.0 \pm 0.2 \; aA3$	$7.2 \pm 0.3 \ aA3$	$6.9 \pm 2.3 \; aA1$	$7.3 \pm 0.1~aB2$	$7.1 \pm 0.7~aB2$	$5.9 \pm 0.6~\text{aA2}$	$9.7 \pm 0.4~aA1$	$13.8 \pm 1.3~aB2$	$16.7 \pm 1.0~aB2$
GGP-	MG	$4.3\pm0.2\;aA1$	$4.3 \pm 0.1 \; aA1$	$4.2\pm0.1\;aA1$	$4.5\pm0.4~aA1$	$4.5 \pm 0.1 \\ aA1$	$4.2 \pm 0.1 \; aA1$	$3.1\pm0.3~\text{aA1}$	$3.8 \pm 0.5 \; aA1$	$2.8 \pm 0.5 \; aA1$	$15.1\pm1.0~aA2$	$10.8 \pm 0.5 \; aA1$	$12.9 \pm 0.8 \; aA1$
5261	BR	$4.1\pm0.1\;aA1$	$4.0 \pm 0.1 \; aA1$	$3.9 \pm 0.2 \; aA1$	$5.2\pm0.4~aA12$	$4.8 \pm 0.1 \; aA1$	$4.8\pm1.8~\text{aA12}$	$4.8 \pm 0.2 \; aA2$	$5.0 \pm 0.1 \; aA2$	$4.8 \pm 0.1 \ aA2$	$12.2 \pm 1.1 \ bA1$	$10.7 \pm 1.2 \; aA1$	$11.7 \pm 0.6~\text{aA1}$
	OR	$3.9 \pm 0.1 \; aA1$	$3.9 \pm 0.1 \; aA1$	$3.9 \pm 0.1 \; aA1$	$6.5\pm0.5~\text{aA2}$	$6.1 \pm 0.4~aA2$	$6.2\pm1.1~aA2$	$5.5 \pm 0.3 \; aA2$	$6.2\pm0.2~bA3$	$5.0 \pm 0.6 \; aA2$	$12.8\pm1.7~bA2$	$9.4 \pm 0.3 \; aA1$	$11.6 \pm 0.7 \text{ aA1}$
	RR	$3.9 \pm 0.4 \; aA1$	$4.0 \pm 0.1 \; aA1$	$4.0 \pm 0.1 \; aA1$	$7.9 \pm 0.6~\text{aA3}$	$7.9 \pm 1.4~aA2$	$7.5\pm0.9~aA1$	$7.9 \pm 0.5~aB3$	$5.7 \pm 0.1 \; aA3$	$5.0 \pm 0.3 \; aA2$	$12.4\pm1.2\;bA1$	$14.7 \pm 1.6~aB2$	$16.9 \pm 0.7~aB1$
GGP-	MG	$4.4\pm0.2\;aA1$	$4.4\pm0.2\;aA1$	$4.3\pm0.2\;aA1$	$3.9 \pm 0.5 \; aA1$	$3.7 \pm 0.6 \ a \ A1$	$3.8 \pm 0.8 \; aA1$	$3.2 \pm 0.2 \; aA1$	$2.7 \pm 0.4 \; aA1$	$2.4 \pm 0.3 \; aA1$	$12.7 \pm 1.4 \text{ aA}12$	$13.8 \pm 0.8 \; aA1$	14.1 ± 0.8
49C12													aA12
	BR	$4.0 \pm 0.1 \; aA1$	$4.0 \pm 0.1 \; aA1$	$3.9 \pm 0.1 \; aA1$	$4.0 \pm 0.6 \; aA1$	$4.1\pm0.9\;aA1$	$4.7 \pm 0.7 \; aA1$	$4.3\pm0.3~aA2$	$4.5\pm0.4~aA2$	$4.8 \pm 0.6 \; aA2$	$15.0 \pm 0.3~bB2$	$9.3\pm1.5~aA1$	$10.1\pm0.5~\text{aA1}$
	OR	$3.9 \pm 0.1 \; aA1$	$3.7 \pm 0.4 \; aA1$	$3.7 \pm 0.1 \; aA1$	$5.1\pm0.5~aA12$	$5.5\pm0.8~aA12$	$5.5\pm0.6~aA12$	$5.1 \pm 0.1 \; aA3$	$4.8 \pm 0.2 \; aA2$	$5.0 \pm 0.45~aA2$	$9.4 \pm 1.3 \text{ bA}1$	$11.3 \pm 0.2 \; aA1$	$12.4\pm0.2\;aA1$
	RR	$3.9 \pm 0.1 \; aA1$	$3.7 \pm 0.5 \; aA1$	$3.7\pm0.5~aA1$	$7.2 \pm 0.4~aA2$	$6.5\pm0.3~aA2$	$6.6 \pm 0.8 \; aA2$	$7.8 \pm 0.4 \; aB4$	$4.6\pm0.5~aA2$	$4.8 \pm 0.4 \; aA2$	$11.2\pm1.6~bA1$	$13.0\pm0.3~aB1$	$15.5\pm0.3~aB2$

Lower-case letters denote statistical differences between different genotypes in the same ripening stage and treatment, capital letters denote statistical differences between same genotype and ripening stage on different treatments and numbers denote statistical differences between ripening stages on the same genotype (ANOVA, $P \le 0.05$). Values are the mean of three-independent experiments.