# Influence of Ploidy Levels on Phenotypic and Cytogenetic Traits in Maize and Zea perennis Hybrids

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**Summary** The aim of this work was to analyse: a) the phenotypic and cytogenetic differences of *Zea mays* and *Zea perennis* hybrids 2n=30 and 40; b) the effect of colchicine on their chromosomes pairing in meiosis. The plant materials used for these studies were *Z. mays* (Zm20) 2n=20, *Z. mays* (Zm40) 2n=40 and *Z. perennis* (Zp) 2n=40 and the hybrids  $Zm20\times Zp$  (MP30) 2n=30 and Zm40×Zp (MP40) 2n=40. Some characteristics of *Z. perennis* such as rhizomes, distic ears, perennially and profuse tillering were only observed in MP40 plants. In the other hand, MP30 plants had maize like phenotype; they were annual, with 4 round ears and low tillering. Therefore, phenotypic expression of several traits depends upon the dose of genes from each parent in the hybrid. The genomic formulae proposed for these species and hybrids were Zm20: AmAm BmBm; Zm40: AmAmAmAm BmBmBmBm; Zp: ApApApAp Bp<sub>1</sub>Bp<sub>1</sub> Bp<sub>2</sub>Bp<sub>2</sub>; MP30: AmApAp Bp<sub>1</sub>Bp<sub>2</sub> Bm; MP40: AmAmApAp BmBm Bp<sub>1</sub>Bp<sub>2</sub>. Prezigotene colchicine treatment increased the frequency of quadrivalents in Zm20, Zp and Mp40 and trivalents in MP30. These results suggest homoeology between Bm and Bp<sub>1</sub>Bp<sub>2</sub> genomes from maize and Zp, respectively.

Key words Zea, Maize hybrids, Genomic formulae, Ploidy, Embryo rescue.

Zea is an important genus of the tribe Maydeae. According to Doebley and Iltis (1980) and Iltis and Doebley (1980) it is composed of 2 sections: Section Luxuriantes, which includes the annual Z. luxurians and the perennials Z. diploperennis and Z. perennis and Section Zea comprises only an annual species (Z. mays L.) which can be divided into three subspecies: Z. mays ssp. mays, Z. mays ssp. mexicana and Z. mays ssp. parviglumis. All the species mentioned above have 2n=20 chromosomes except Z. perennis which has 2n=40.

In previous works, Molina and Naranjo (1987), Naranjo *et al.* (1990, 1994), Poggio *et al.* (1990) and Molina and García (1994) obtained cytological evidence supporting a polyploid condition for 2n=20 for the taxa of the genus Zea, and proposed the genomic formulae for them. Moore *et al.* (1995) confirmed that maize is a complete tetraploid, although there are two separate genomes. Most of the larger chromosomes (1, 2, 3, 4, 6) arrange a genome and the remaining shorter ones (5, 7, 8, 9, 10) arrange another.

Thomas and Kaltsikes (1977) and Thomas and Ingram (1987) demonstrated that a dilute solution of colchicine applied to the outset of meiosis promotes the formation of multivalents from bivalents. Jackson and Murray (1983) suggested that colchicine can disrupt the genetically controlled position of the genomes promoting intergenomic pairing and revealing criptic homology in amphidiploids. Furthermore, pairing between heterologous chromosomes was observed in rye after dilute colchicine treatment. This could be due to a homology segment shared by those chromosomes (Puertas *et al.* 1984).

Maize and Z. perennis showed criptic homology when cells were treated in pre-zygotene with diluted colchicine during a 12 h period. Quadrivalents appeared in maize and its number increased

in Z. perennis. These results support the allotetraploid nature of the genus Zea and suggests a basic genome x=5 (Poggio *et al.* 1990). It is also suggested that the genus has homoeologous genomes that probably do not pair when a Ph like gene is present.

The purpose of this work was to analyze the phenotypic and cytogenetic behaviour of hybrids between *Zea mays* and *Z. perennis* with different ploidy levels as well as the effect of premeiotic treatment with diluted colchicine solutions on the meiotic behaviour.

### Materials and methods

The plant materials used for these studies were Zea mays spp. mays cv. "Colorado Klein" (Zm20) 2n=20, the maize inbreeds 2n=40 (Zm40) N103A, N104B, N107C, N107B, and 90–2189–2190 supplied by the Maize Genetics Coop. Stock Center (Urbana, Illinois, USA); Zea perennis (Zp) 2n=40, from Ciudad Guzmán Jalisco, México, demised by Dr. Prywer and grown since then at the Instituto Fitotécnico de Santa Catalina. The hybrids studied were MP30 (Z. mays cv. "Colorado Klein"×Z. perennis) 2n=30 and MP40 (Z. mays N107B×Z. perennis and N107C×Z. perennis) 2n=40.

MP30 hybrid plants were obtained through embryo rescue. In autumn, maize silks were cut off to 2 and 3 cm and pollinated with Zp. Two days after pollination, the maize ears were sprayed with a 0.45 mM 2,4-dichlorophenoxiacetic acid (2,4-D) solution (Furini and Jewel 1995) to help grain formation. The ears were harvested 21 days after pollination and caryopsis were disinfected with a 2.5% sodium hypochlorite solution. Embryos with a maximum length of 0.5 mm were plated on the basic medium of García and Molina (1992) supplemented with 4.5  $\mu$ mol L<sup>-1</sup> 2,4-D and then incubated at 28–30°C with a 16 h photoperiod. Plants were regenerated on 2,4-D free medium.

The colchicine treatment was done according to Poggio *et al.* (1990). The stems were cut under a dilute solution of colchicine  $(0.5 \times 10^{-4} \text{ M})$  and kept therein for 12 h. Before anthers fixation, the stems were placed in tap water during 24 h. Control materials were fixed in 3 : 1 (absolute ethanol : acetic acid) solution.

Anthers were squashed in 2% acetic haematoxyline (Nuñez 1968). The pairing configuration was determined at diakinesis (metaphase I). Only cells with well spread plates were scored.

### Results

Only two caryopsis developed on one cv. "Colorado Klein" ear were obtained when Zm20 was outcrossed by Zp. Twenty one days after pollination these caryopsis showed deficient endosperm development and their embryos had a length of 0.5 mm. Both embryos gave rise to green organogenic calli. Five plants with profuse radicular systems were regenerated over a 5 months period.

The number of developed caryopsis changed significantly according to the maize genotype used as female parent (Table 1). The best results were undoubtedly attained with the inbred N107C. Furthermore when the reciprocal crosses were done inbred N107C was the only one that induced

Maize genotype	Pollinated ears	Developed caryopsis	Hybrid seed germination (%)	Hybrid pollen fertility (%)	
N 103 A	10	0	0	0	
N 104 B	10	39	0	0	
N 107 B	10	400	20	50-60	
N 107 C	10	2450	100	70-90	
90-2189/2190	10	0	0	0	

Table 1. Results of crossing different maize inbreeds (2n=40) with Zp

MP30			MP40			
1.	Perennial	1.	Annual of Biennial			
2.	Seeds viability=1%	2.	Seeds viability=70-90%			
3.	Pollen fertility=3–5%	3.	Pollen fertility=70–95%			
4.	Distic ears. The rachis dissarticulates at maturity scattering the seeds.	4.	Small 2 to 4 rowed ears. Nacked seeds disposed on a rigid rachis.			
5.	Average number of tillers/plant=20	5.	Average number of tillers/plant=4			
6.	Short and defined rhizomes	6.	No rhizomes			
7.	Flowering response to photoperiod: short day qualitative	7.	Flowering response to photoperiod: short day quantitative.			
8.	Multiplication through rhizomes.	8.	Cannot be multiply asexually			
9.	Does not hybridise naturally	9.	Naturally hybridise to some maize genotypes			
0.	Shows clorantia under stress conditions	10.	Does not show clorantia			

Table 2. Pollen fertility and phenotypic differences of Zm×Zp hybrids with different ploidy levels

the development of viable hybrid seeds.

Ploidy level or the number of chromosomes contributed by each parent, notably affected the phenotype of the hybrid plants (Table 2, Fig. 1).

The most frequent meiotic configurations observed in diakynesis-metaphase I (Table 3) were the following:

Zm20 (Table 3, Fig. 2A). A 100% of the analysed cells showed 10II. Secondary association, separation of the chromosomes in two groups of 5 each and an average of 14 chiasma/cell were observed in a 40% of the cells. Molina and Naranjo (1987), Naranjo *et al.* (1990) according to cytogenetic studies fulfilled in Zm20, Zp, *Z. diploperennis* and their hybrids proposed a basic chromosome number x=5 for genus *Zea* and the genomic formulae AmAm BmBm for Zm20.

Zm40 (Table 3) revealed 10IV (30%) and 9IV+2II (23.94%) with an average of 8.34IV+3.24II and 33.75 chiasma/cell. According to these results and the fact that the basic number for the genus *Zea* is x=5, it is proposed for Zm40 the genomic formulae AmAmAmBmBmBmB.

Fig. 1. Ears of *Z. mays*, *Z. perennis* and their hybrids: A) Zm; B) Zp; C) MP30; D) MP40. Scale=3 cm.

Zp (Table 3, Fig. 2B) shown 5IV+10II

(54.47%) and 4IV+12II (20.15%). Only in a 5% of the cells more than 5IV were observed with an average of 4.44IV+11.02II. The average number of chiasma/cell was 15.69 in bivalents and 18.87 in tetravalents with a whole of 34.56. The most frequent chromosomic pairing found in Zp suggests for this species the genomic formulae ApApApAp Bp<sub>1</sub>Bp<sub>2</sub>Bp<sub>2</sub>.

Hybrid MP30 (Table 3, Fig. 2C) had 5III+5II+5I (53.71%) and 4III+6II+6I (25.71%) with an average/cell of 4.71III+5.39II+5.07I. More than 5IV were observed just in a 6% of the cells, being the average number of chiasma/cell=23.06. The trivalents were arranged by the pairing of

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Species and hybrids	2n	diakinesis-metaphase configurations			%	No. of cells studied	
		I	II	III	IV	70	No. of cens studied
Zm20	20		10			100	200
Zm40	40		0		10	30.28	142
			2		9	23.94	
			4		8	16.90	
			6		7	10.52	
			8		6	7.75	
			10		5	4.23	
			12		4	0.70	
			14		3	1.40	
		1	6	1	6	0.70	
		1	8	1	5	0.70	
		1	-	1	9	1.40	
		1	2	1	8	1.40	
		X=0.04	3.24	0.04	8.34		
Zp	40		18		1	1.49	1.34
чr			16		2	3.73	
			14		3	6.72	
			12		4	20.15	
			10		5	54.47	
			8		6	5.22	
		1	16	1	1	0.75	
		2	15	-	2	0.75	
		2	13		3	0.75	
		2	11		4	2.98	
		1	10	1	4	0.75	
		2	9		5	1,49	
		4	8		5	0.75	
		X=0.16	11.02	0.01	4,44		
MP30	30	7	7	3		2.14	140
		6	6	4		25.71	
		4	7	4		5.00	
		7	4	5		1.43	
		5	5	5		53.57	
		3	6	5		5.71	
		1	7	5		0.71	
		4	4	6		5.00	
		4	3	7		0.71	
		X=5.07	5.39	4.71			
MP40	40		20		0	0.45	224
			18		1	3.12	
			16		2	8.48	
			14		2 3	16.96	
			12		4	21.43	
			10		5	27.23	
		1	16	1	1	0.45	
		1	8	1	5	2.23	
		1	12	1	3	2.23	
		2	11		4	0.45	
		2	9		5	2.23	
		2	7		6	0.45	
		2	13		3	1.80	
		4	12		3	0.45	
		2	7		6	0.45	
		$\bar{X} = 0.20$	11.62	0.03	4.12		

### Table 3. Meiotic configurations of Zea species and F<sub>1</sub> studied

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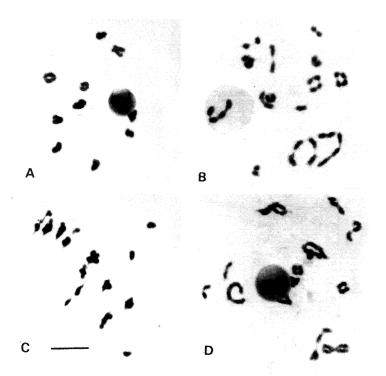


Fig. 2. Meiotic configurations observed in diakynesis–metaphase I in A) Zm20 (10II); B) Zp (51V+ 10II); C) MP30 (51II+5II+5I); D) MP40 (51V+10II). Scale=10  $\mu$ m.

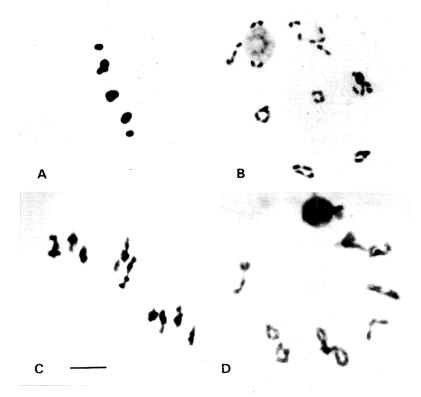


Fig. 3. Meiotic configurations induced by colchicine treatment in: A) Zm20 (5IV); B) Zp (8IV+4II); C) MP30 (10III); D) MP40 (9IV+2II). Scale=10 μm.

one Zm20 chromosome with two Zp chromosomes (AmApAp). The bivalents were the result of the pairing among the homoeologous chromosomes of Zp  $(Bp_1Bp_2)$  and the univalents were the chromosomes of Zm20 (Bm). The genomic formulae proposed for the hybrid MP30 is AmApAp  $Bp_1Bp_2$  Bm.

Hybrid MP40 (Table 3, Fig. 2D) had 5IV + 10II (27.23%) and 4IV + 12II (21.43%) with an average of 4.12IV + 11.62II per cell. More than 5IV were observed only in a 1% of the cells. The average of chiasma per cell was 33.59. The genomic formulae proposed is AmAmApAp BmBm Bp<sub>1</sub>Bp<sub>2</sub>.

The effects of colchicine treatment on the meiotic configurations of analysed species and hybrids are shown in Figs. 3–5. Control Zm20 plants formed only bivalents, but treated plants showed at least one quadrivalent in the 61% of the cells, with a maximum of 5IV in the 11% of the cells (Figs. 3A, 4a). In Zm20, colchicine promoted pairing between homoeologous chromosomes from both genomes of *Zea* (AmAm BmBm). It was also observed an enlargement of the chiasma average number, from 14 (control) to 20 (treated) per cell.

Zm40 treated plants showed a little increase of quadrivalents if compared to control plants, but the difference was not significant. Neither hexavalents nor octovalents were observed (Fig. 4b).

In Zp (Figs. 3B, 4c), colchicine treatment induced a remarkable increase of the average number of IV per cell, from 4.44IV (control) to 5.83IV (treated) up to a maximum of 10IV per cell. In this case, the increment of quadrivalents would be due to the pairing induction amongst homoeologous chromosomes that normally do not pair. The average number of chiasma/cell is 41.80, a 21% higher than control plants.

In the hybrid MP40 (Figs. 3C, 4d) a remarkable increase in the number of IV was observed in treated plants ( $\overline{X}$ =5.24) if compared to control plants ( $\overline{X}$ =4.12). In this case, the increment in the number of IV would be due to pairing between homoeologous genomes of both species (BmBm Bp<sub>1</sub>Bp<sub>2</sub>).

In the hybrid MP30 (Figs. 3D, 5) a highly significant increase of III was observed in the treated material ( $\overline{X}$ =7.85) when compared to control plants ( $\overline{X}$ =4.71). The colchicine treatment induced pairing of homoeologous genomes from both species (BmBm Bp<sub>1</sub>Bp<sub>2</sub>) which notably increased the percentage of trivalents and the number of chiasma/cell from 23.06 (control) to 27.09 (treated).

### Discussion

Zm40 and Zp hybridize without restrictions under natural conditions. Their hybrids have meiotic regularity and fertile offspring. On the other hand, crossings between Zm20 and Zp only yield 0.1 to 1% of viable seeds. Although a few hybrid caryopsis developed until 21 days after fecundation, the endosperm collapses becoming in the abortion of the embryos. This problem can be overcome through rescue and *in vitro* culture of the immature embryos.

Z. mays and Z. perennis hybrid phenotypes differ according to their chromosome number (Table 2). This fact suggests that phenotypic expression of several traits depends upon the dose of genes from each parent in the hybrid. This would explain the great variability observed in the next generations (Molina 1978).

Genetic, cytogenetic and biochemical studies provided enough evidence to suggest that the genus Zea has a basic chromosome number x=5 and that Zm20 is an allotetraploid with two homoeologous genomes Am and Bm (Rhoades 1951, Ghatnekar 1965, Vijendra Das 1970, Gottlieb 1982, Bennett 1983, 1984, Molina and Naranjo 1987, Naranjo *et al.* 1990). The maize Am genome five chromosomes pair arranging II with the homoeologous Bm chromosomes only when their respective homologous are not present in the cell, as for example during diakynesis-metaphase I of haploid plants (Mc Clintock 1933, Chaganti 1965, Ting 1966, 1969).

According to Molina and Naranjo (1987), Zp has 3 genomes Ap Bp<sub>1</sub> and Bp<sub>2</sub>. The Ap genome

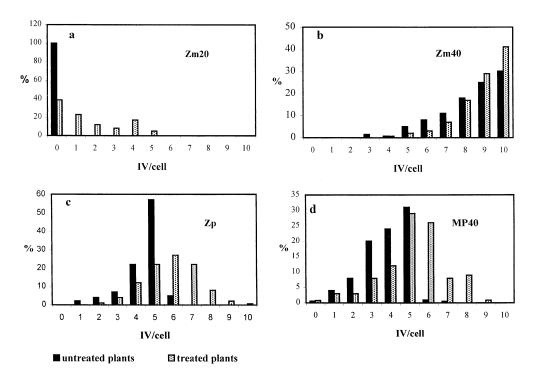


Fig. 4. Number of IV/cell in colchicine  $(0.5 \times 10^{-4} \text{ M})$  treated and control plants of a) Zm20, b) Zm40, c) Zp and d) MP40.

pairs building 5IV whilst Bp<sub>1</sub> and Bp<sub>2</sub> genomes are homoeologous and pair between them only when their respective homologous are not present in the nucleus as it is observed in the hybrids MP30 and MP40. Otherwise it was seen that A genomes from maize and Zp pair arranging trivalents (AmApAp) or quadrivalents (AmAmApAp) depending if the hybrid is 2n=30 or 40, respectively. The chromosomes shared by both species are 1, 2, 3, 4 and 6 from maize and 1–2, 3–4, 7–8, 9–10 and 13–14 from Zp (Molina 1986). Jackson and Murray (1983) demonstrated that the application of a dilute solution of colchicine  $(0.5 \times 10^{-4} \text{ M})$  during

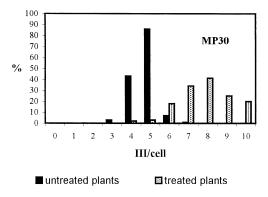


Fig. 5. Number of III/cell in MP30 plants treated with a colchicine solution  $(0.5 \times 10^{-4} \text{ M})$  and control plants.

pre-meiosis promotes intergenomic pairing and reveals cryptic homology. With this technique, Poggio *et al.* (1990) and Naranjo *et al.* (1994) confirmed the existence of cryptic homology and the polyploid nature of Zm20, Zp and the hybrids between Zp and Z. *diploperennis* (2n=30-40). The 11% of the Zm20 cells showed 5IV, which points out that all the Am and Bm homoeologous chromosomes pair in these cells (Figs. 4a, 3A). All the opposite, colchicine treatment did not promote any significant increase in the number of quadrivalents per cell in Zm40 (Fig. 4b).

Colchicine treated Zp plants showed a significant increase in the number of quadrivalents per cell because of  $Bp_1$  and  $Bp_2$  genomes pairing (Fig. 4c). In Zm and Zp hybrids, colchicine treatment notably increased the number of III (MP30) and IV (MP40) per cell because of pairing between Bm genome from maize and  $Bp_1Bp_2$  genomes from Zp. This confirms that Bm and  $Bp_1Bp_2$  genomes are homoeologous.

#### Conclusions

Beginning with the study of the hybrids between Zm with two ploidy levels (2n=20 or 40) and Zp, the following statements can be expressed:

Both species naturally outcross whether they have the same chromosome number therefore they show chromosome compatibility.

The plant phenotype is determined by the dose of genes donoured by each parent to the hybrid.

The colchicine treatment reveals cryptic homology between Bm genome from maize and  $Bp_1$  Bp<sub>2</sub> genomes from Zp.

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