

Morphology and Chromosome Pairing of a Hybrid Between *Triticum durum* Desf. and *Haynaldia villosa* (L.) Schur.

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Summary. Intergeneric hybrids between T. durum Desf. (2n = 4x = 28, AABB) and Haynaldia villosa (L.) Schur. (2n = 2x = 14, VV) was obtained at a frequency of about 5.6% of pollinated florets. Phenotypically the F₁ plants resemble more the maternal parent than the H. villosa and are almost completely sterile. However, some seeds were obtained on selfed and backcrossed heads with the durum wheat parent. The hybrid had a somatic complement of 2n = 3x = 21, ABV, with a mean chromosomal relationship of 13.62 univalents, 3.30 bivalents, and 0.26 trivalents. The high pairing was likely due to gene(s) of H. villosa interacting with the 5B homoeologous restricting system of wheat.

Key words: Intergeneric hybrids – *Triticum* – *Haynaldia* – Homoeologous pairing – Pairing promoters

Introduction

Interest in the interspecific hybridization in *Triticinae* has arisen from the need to exploit the variability of the related species for genetic improvement of *durum* and common wheat (McFadden and Sears 1947; Riley and Kimber 1966; Feldman 1979) and to investigate evolutionary relationships between species (Kihara 1954; Sears 1975; Alonso and Kimber 1981).

Haynaldia villosa (L.) Schur. is an allogamous annual plant native of the Mediterranean region; it is a diploid (2n = 14) with a genome composition of VV. As a germplasm donor this species has been found to be interesting because of marked variability of some of its morphological characteristics, drought tolerance as well as resistance to disease such as powdery mildew, stem and leaf rusts (Hyde 1953; Pasquini et al. 1978; Scarascia et al. 1982). Hybridization of *H. villosa* with diploid, tetraploid and hexaploid species of *Triticum* has been investigated (Strampelli cited in Raineri 1914; Sando 1935; Sears 1953; Kostoff and Arutinova 1937; McFadden and Sears 1947; Nakajima 1959; Halloran 1966; Gupta 1972; Meletti et al. 1977; Piralov 1981; Ibraginov 1979; Chen et al. 1982) and found to have different but usually low degrees of crossability (from 1.2 to 17.2% of pollinated florets). As in other instances of interspecific hybridization the crossability depends both on the specific genotypes involved and on the effects due to changes in seasonal conditions and crossing procedure. The *H. villosa* chromosomes have, none the less, been successfully added, collectively and individually, to the full chromosome complement of hexaploid wheat (Hyde 1953; Sears 1953).

This study was carried out to investigate the morphology, the level of chromosome pairing and the fertility of T. durum $\times H$. villosa hybrids.

Materials and Methods

The Triticum durum Desf. (2n = 4x = 28, AABB) parent used in this study was the Italian semi-dwarf cultivar 'Creso'. The *H. villosa* (L.) Schur. (2n = 2x = 14, VV) strain was collected at the experimental farm of the Faculty of Agriculture, University of Bari (Italy), located 15 km from Bari.

The parental lines were grown in the field and during the spring of 1981; 250 *durum* wheat florets were emasculated and pollinated with *H. villosa*. Five embryos were excised from developing seeds 12–15 days after pollination and cultured on sterile B 5 medium (Gamborg et al. 1968). Of these, three embryos developed into plantlets on the medium and were transplanted into pots at the three leaf stage; two of them grew into vigorous F_1 hybrids. Ten other developing seeds were left to ripen on the plants and they yielded nine hybrid seeds. Although shrivelled, five of these germinated normally on moist filter paper in petri-dishes in the laboratory.

The root-tip of the F_1 hybrids were pre-treated for 4 h with 0.05% colchicine and then fixed in Carnoy's 3:1 fixative for mitotic studies. Individual anthers with PMC's at metaphase I were fixed in Carnoy's 6:3:1 fixative for meiotic studies. A standard Feulgen staining and squash procedure was used with both root-tips and PMC's to prepare material for cytological examination. Spike characteristics of the parental lines and F_1 hybrid were measured from 10 spikes.

Results and Discussions

The hybrid *T. durum* \times *H. villosa* was produced with a relatively low frequency. In fact two seedlings were obtained from the five cultured embryos and five others were obtained from nine mature seeds. This represents a seed set of 5.6% and a hybrid seedling frequency of 2.8% of pollinated florets in the field conditions, since 250 pollinations were carried out. It is worth noting that the in vitro culture of the hybrid embryos excised from developing seeds 12–15 days after pollination is not needed to obtain hybrid plants, since most of the mature seeds had enough endosperm for a normal germination in petri-dishes.

All the plants analysed had the expected somatic chromosome number (2n=21) and no chromosome

Table 1. Mean spike characteristics from 10 spikes of *T. durum* Desf. cv. "Creso", *T. durum* Desf. cv. "Creso" \times *H. villosa* (L.) Schur. hybrid, and *H. villosa* (L.) Schur.

Character	T. durum	T. durum × H. villosa	H. villosa	
Spike lenght, cm	6.1	9.2	8.4	
Spike width, cm	1.6	1.4	1.5	
Spikelet length, cm	1.7	1.9	2.0	
Spikelet/spike, n.	16.6	23.4	25.1	
Florets/spikelet, n.	4.8	5.0	4.2	
Kernels/spikelet, n.	3.60	0.07	2.00	
Pollen fertility, %	95.4	0.8	88.7	
Glume body length, cm	1.1	1.0	0.7	
Glume awn length, cm	0.1	1.4	3.7	
Lemma body length, cm	1.1	1.3	1.3	
Lemma awn length, cm	13.1	9.7	4.6	
Anther length, mm	3.6	4.2	10.9	
Glume hairiness	absent	absent	present	
Rachis	tough	fragile	fragile	

instability was observed. Phenotypically the hybrid plants resemble the maternal parent more than the *H. villosa*. The hybrid spike characteristics are presented in Table 1, as are those for the *durum* wheat and *H. villosa* parents. The spikes of the hybrid (Fig. 1) were longer and with more spikelets than those of the *durum* wheat. Their glumes were provided with short awns, while the lemma awns were very long (about 10 cm). The rachis of the spike was as fragile and hairy as that of the *H. villosa*.

The pollen fertility, estimated as the percentage of pollen grains stainable with aceto-carmine, was less than 1% and the spikes were almost completely sterile; only 1 or 2 seeds per selfed and back-crossed heads with the *durum* wheat pollen were obtained. The chromosome number of the F_2 and BC_1 plants (2n = 42) and 2n = 35, respectively), along with the meiotic analysis of the PMC's of the F_1 plants (2n=21), indicated that the fertility of the F₁'s was very likely due to meiotic non reduction in a number of cells that resulted in male and female gametes with unreduced chromosome number. This process, that determines a certain fertility in interspecific and intergeneric hybrids expected to have little or no chromosome pairing, has been recently reviewed and demonstrated for T. eldreichii \times T. durum F₁ hybrid by Mann and Sasakuma (1977).

Table 2 shows the results of the chromosome association at the first meiotic methaphase over 100 pollen mother cells of the F_1 hybrids *T. durum* × *H. villosa*, along with the chromosome pairing of the haploid of *T. durum* (Lacadena and Ramos 1968), and of the hybrids *T. dicoccoides* × *H. villosa* (McFadden and Sears 1947) and *T. aestivum* × *H.villosa* (euhaploid and nulli-5B haploid; Halloran 1966).

Table 2. Chromosome pairing at metaphase I in F_1 hybrids among T. durum, T. dicoccoides, T. aestivum and H. villosa, and in the haploid of durum wheat

Hybrid	Chromosome	No. of cells	Chromosome associations					References	
	number 2n		I	II		111	Other		
				Rod	Ring	Total			
T. durum × H. villosa	21	100	13.62 (8 – 19)	3.15 (1–6)	0.15 (0-2)	3.30 (1-6)	0.26 (0-2)	-	Authors
T. durum (euhaploid)	14	94	13.19		-	0.37	-	_	Lacadena and Ramos 1968
T. dicoccoides \times H. villosa	21	100	19.35	-	-	0.79		0.07	McFadden and Sears 1947
T. aestivum $ imes$ H. villosa	28	100	27.02	0.47	0.02	0.49		_	Halloran 1966
T. aestivum $ imes$ H. villosa	27 (5B absent)	100	11.96	3.81	1.00	4.81	0.86	0.71	Halloran 1966



Figs. 1 – 3. 1 (left to right) *T. durum* cv "Creso"; *T. durum* \times *H. villosa*; *H. villosa*. **2** Somatic metaphase of *T. durum* \times *H. villosa* hybrid. **3** Meiotic metaphase in *T. durum* \times *H. villosa* hybrid with $1^{III} + 6^{II} + 6^{I}$

The main chromosome pairing observed in the *durum* wheat \times *Haynaldia* hybrid (3.15 rod and 0.15 ring bivalents, with a range from 1 to 6, and 0.26 trivalents, with a range from 0 to 2) was much higher than the haploid of *durum* wheat (Lacadena and Ramos 1968) which has an average of 0.37 bivalents

per cell (with a range from 0 to 2 bivalents) due to autosyndesis within the wheat complement.

Previous studies on the association of *Haynaldia* and *Triticum* chromosomes at meiosis have indicated that the pairing is low and variable. McFadden and Sears (1947) observed 0.79 bivalents per cell in the

T. dicoccoides \times *H. villosa* hybrid. In the crosses with *T. aestivum*, Halloran (1966) observed 0.49 bivalents, while Chen et al. (1982) detected 1.66 e 2.29 bivalents respectively in two hybrids having different wheat parents.

It is interesting to compare the meiotic behaviour of the durum \times H. villosa hybrid with that of the T. aestivum \times H. villosa nulli-5B haploid (Halloran 1966), considering that the latter has 7 chromosomes of the D genome. The pairing in the durum hybrid was only slightly lower than in that of the above cited hybrid which lacks the Ph gene (suppressor of homoeologous pairing) located on the long arm of the 5B chromosome (Sears and Okamoto 1958; Riley and Chapman 1958). The high pairing observed in the hybrid examined can be due to the presence of a gene (or genes) in the H. villosa genotype that interacts with the 5B homoeologous restricting system of wheat.

Genotypes that promote pairing or suppress the Ph effects have been found in Ae. speltoides (Riley et al. 1961), Ae. mutica (Riley and Law 1965) and Ae. longissima (Mello-Sampayo 1971). These promoters of pairing have no effect in the diploids themselves, where complete pairing of homologues is the rule and there are no homoeologues; however, in hybrids with wheat they increase the pairing potential of homoeologues. The level of pairing induced by the highpairing Ae. speltoides and Ae. mutica strains is almost similar to the one taking place in the same hybrids in absence of chromosome 5B. Moreover, strains of these species were found which affected pairing less strongly or not at all (Dover and Riley 1972; Kimber and Athwal 1972; Dvorak 1972). Partial suppression of Ph evidently occurs in hybrids with tetraploid Ae. caudata (Riley 1968) and Ae. umbellulata (Riley et al. 1973). Mochizucki (1962), also, found that the addition of a particular pair of Agropyron elongatum chromosomes to durum wheat resulted in a substantial amount of homoeologous pairing.

If the *H. villosa* genotype suppresses the *Ph* effects, the pairing observed in the hybrid examined is presumed to be mainly between the A genome chromosomes with the corresponding homoeologues of genome B, such as in the nulli-5B haploid of common wheat (Riley and Law 1965) and in the hybrids of wheats lacking the Ph gene and rye (Dhaliwal et al. 1977; Giorgi and Cuozzo 1980). However, on the basis of trivalent frequency at meiosis in the durum wheat \times H. villosa hybrid, it can be assumed that some chromosomes of H. villosa are pairing with those of Triticum durum, as already observed by Halloran (1966) in common wheat. The analysis of homoeologous pairing by Giemsa C-banding of the hybrid meiotic chromosomes can be establish the pairing degree between wheat and Haynaldia chromosomes. The fact that the Haynaldia villosa chromosomes can pair with those of wheat indicates that every useful character of Haynaldia can be transferred to the wheat complement by homoeologous recombination.

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