

STUDIES ON IN VIVO INTERSPECIFIC HYBRIDISATION IN BRASSICA

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SUMMARY

Studies were undertaken to explore the possibility of exploiting interspecific variability by species hybridisation in the genus Brassica. Six divergent species, viz. B. juncea, B. napus, B. campestris, B. chinensis, B. nigra and B. japonica were used in an attempt to produce interspecific hybrids by following conventional (in vivo) method of hybridisation. The resultant F_1 hybrids showed considerable variations for morphological traits contributing to yield. The meiotic analysis of F_1 hybrids revealed the presence of bivalents and univalents which can be attributed to the difference in the genetic constitution of the parental species. The pollen fertility of various cross combinations ranged from 57.00% (B. campestris x B. chinensis) to 16.00% (B. chinensis x B. juncea). The variation for pod setting was from 4.76% (B. chinensis x B. napus) to 71.05% (B. juncea x B. napus). The hybrid of cross, B. juncea x B. napus was recorded for maximum seed set (74.69%), whereas the hybrid of cross, B. chinensis x B. napus was recorded for minimum seed set (50.00%). The per cent germination of hybrid seed was maximum (54.54) again in the cross, B. juncea x B. napus and minimum (9.09) in the cross, B. napus x B. juncea. The present investigation thus showed the potential of using in vivo method of interspecific hybridisation for induction of genetic variability in Brassica species and introgression of desirable genes from related/ non cultivated species to commercially cultivated species.

INTRODUCTION

The oilseed crops have an important role in country's economy. In the world oiliferous Brassica ranks fifth in importance among oilseeds. There are 159 species in the genus Brassica, which includes a number of wild species also. The study of chromosome number and meiotic behaviour indicated polyploid nature of Brassica species. The elementary species are B. nigra, B. oleracea and B. campestris with $2n = 16, 18$ and 20 respectively, whereas the

natural allopolyploids B. carinata, B. juncea and B. napus have $2n = 34$, 36 and 38 respectively. A-genome was assigned to B. campestris, B to B. nigra and C to B. oleracea. The genomes of higher chromosomal species were designated AB for B. juncea, BC for B. carinata and AC for B. napus. A new species of B. tournefortii with $2n = 20$ and D genome has also been identified. All these species are chromosomally balanced and show regular bivalent formation.

MATERIAL AND METHODS

The experimental material consisted of different cultivars of following species :

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| 1. <u>B. nigra</u> | 2. <u>B. campestris</u> cv. TL-15 |
| 3. <u>B. chinensis</u> | 4. <u>B. japonica</u> |
| 5. <u>B. juncea</u> cv. RLM 198 | 6. <u>B. napus</u> cv. Tower |

In each case, hand pollinations were made in the field. The emasculations were done generally in the evening and pollinations in the morning at about 10 o'clock. Observations on different morphological traits like plant height, length of main shoot, number of primary and secondary branches, days to first flowering, number of pods on main shoot, pod length and number of seeds per pod were carried out. The crossability of different crosses was observed by recording per cent pod set, per cent seed set and per cent germination of hybrid (F_1) seeds. For cytological studies, young buds were collected in the morning and fixed in freshly prepared 3:1 alcohol & acetic acid fixative. Anthers were macerated in 2 per cent acetocarmine and studied under microscope to determine different meiotic stages (diakinesis, metaphase-I, anaphase-I). Pollen fertility was recorded by using acetocarmine staining technique.

RESULTS

The differences for various morphological characters, viz. plant height, length of main shoot, number of primary and secondary branches, days to first flowering, number of pods on main shoot, pod length and number of seeds per pod were recorded in case of Brassica species and interspecific crosses. Highly significant differences for most of the morphological traits were observed between different species and their crosses, except for number of primary and secondary branches.

1. Per cent pod set : Percentage of pod set ranged from 4.76 (B. chinensis x B. napus) to 71.05 (B. juncea x B. napus), whereas it was 14.54 and 23.80 in the crosses of B. napus x B. juncea and B. campestris x B. chinensis respectively (Table 1).
2. Per cent seed set : It was highest in cross of B. juncea x B. napus (74.69%) and lowest in case of B. chinensis x B. napus cross (50.00%). Except for the first cross, percentage was almost same in other three crosses (Table 1).
3. Per cent germination of hybrid seeds : It ranged from 9.09% (B. napus x B. juncea and B. chinensis and B. napus) to 54.54%

(B. juncea x B. napus). It was 45.45% in cross of B. campestris x B. chinensis).

4. Per cent pollen fertility : Pollen fertility ranged from 57% in the cross of B. campestris x B. chinensis to 16% in the cross of B. napus x B. juncea. The pollen fertility in the crosses, B. juncea x B. napus and B. chinensis x B. napus was 39% and 28% respectively.

Meiotic behaviour of the interspecific hybrids of the cross of B. juncea x B. napus revealed the presence of 37 chromosomes in the form of bivalents and univalents. At anaphase-I, they were unequally distributed to both the poles with some laggards. The cross of B. campestris x B. chinensis showed the presence of 10 bivalents, whereas in case of cross B. chinensis x B. napus 10 bivalents and 9 univalents were present.

DISCUSSION

Existing variability in a species can be supplemented with new variability by wide hybridisation. The tribe Brassicaceae offers a great potential to incorporate genes from one species to another. Variability for morphological characters can be utilized in increasing yield. Variability for different morphological characters was also recorded by many other workers. However, several crossability barriers are encountered in wide hybridisation, so it is essential to know about crossability and genomic relations between different species of Brassica. The findings indicated substantial variability for various economic characters. The observed differences in crossability may be due to the differences in the genetic constitution of the parental species. Variability for this character has also been reported by Khanna and Chowdhury (1974 and 1981) while working with B. juncea, B. campestris, B. chinensis and B. carinata and their crosses. The differences in per cent pod set and seed set may be attributed to the differences in pollen fertility, rate of pollen tube growth and penetration and abortion rate of embryos. Low germination percentage was mainly due to seed sterility of F_1 's.

Meiotic studies also are in conformity with the studies of Morinaga (1934), Sikka (1940), Ramanujam and Srinivasachar (1943) and Nwankit (1970 and 1971). They recorded the presence of bivalents and univalents. The presence of bivalents and univalents depends upon the genetic constitution of the parental species. The formation of bivalents in crosses shows the possibilities of transferring the desirable characters from one species to other, by the means of crossing over that takes place during the meiosis. So, desirable lines could be developed by extensive crossings. It will lead to the improvement in the agronomically desirable varieties of rapeseed and mustard.

Various aspects of crossability studied in F₁ hybrids of crosses involving different Brassica species (Tables 1, 2 and 3) are :

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Table 1 : Per cent pod and seed set in different crosses of Brassica species.

Cross	No. of buds pollinated	No. of pods formed	Total No. of seeds	Pod set (%)	Seed set (%)
<u>B. juncea</u> x <u>B. napus</u>	950	675	3025	71.05	74.69
<u>B. napus</u> x <u>B. juncea</u>	495	72	245	14.54	58.05
<u>B. juncea</u> x <u>B. campestris</u>	1050	250	785	23.80	52.33
<u>B. chinensis</u> x <u>B. chinensis</u>	840	40	100	4.76	50.00
<u>B. chinensis</u> x <u>B. napus</u>					

Table 2 : Germination percentage of hybrid seeds of different crosses

Cross	No. of seeds sown	No. of seeds germinated	Per cent germination
<u>B. juncea</u> x <u>B. napus</u>	44	24	54.54
<u>B. napus</u> x <u>B. juncea</u>	11	1	9.09
<u>B. campestris</u> x <u>B. chinensis</u>	22	10	45.45
<u>B. chinensis</u> x <u>B. napus</u>	11	1	9.09

Table 3 : Per cent pollen fertility of F₁ hybrids

Cross	% pollen fertility
<u>B. juncea</u> x <u>B. napus</u>	39
<u>B. napus</u> x <u>B. juncea</u>	16
<u>B. campestris</u> x <u>B. chinensis</u>	57
<u>B. chinensis</u> x <u>B. napus</u>	28