



Research Notes

Transgressive segregation for yield and yield components in some inter and intra specific crosses of desi cotton

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Desi cottons belonging to *G. arboreum* and *G. herbaceum* though have low yield potential, their consistent fibre properties are much valued and liked in the textile industry. However, due to introduction of high yielding long stapled intra hirsutum hybrids, the area under diploid cotton has been reduced drastically throughout the country. Therefore to regain the importance of desi cotton there is an urgent need to develop high yielding desi cotton varieties possessing superior fibre properties. With this view about 70 crosses involving diploid genotypes of *G. arboreum* and *G. herbaceum* species were obtained and their segregating generations studied. However, the present study is confined to F_2 and F_3 bulks from ten inter and eight intra specific crosses that were evaluated for transgressive segregates.

The experimental material consisted of 12 diverse parents, of which Mdl 1874, 1875, 2450, 2452, 2463, 2562 and 2582 were from race indicum of *G. arboreum* and PA 141 and PA 183 from race bengalense. The other three viz. Suyog, Jaydhar and Raghavendra belonged to *G. herbaceum* species. The details of 18 crosses included in the study are given in Table 1. Parents and their 18 F_2 populations were sown in 1999 and the parents and F_3 populations were sown in 2000 kharif season in a randomized block design with three replications. The parents were grown in two row plots and the F_2 's were raised in 8 rows and F_3 's in 4 row plots. The row length was 4.5 m. Five competitive plants from each of the parental lines and 30 plants from each of the F_2/F_3 population were randomly selected and observations recorded for five characters. Percentage of transgressive segregation in the F_2 and F_3 generations was obtained by defining the extreme progeny as significantly transgressive segregates (i.e. the lines that exceeded their better parent mean

and L.S.D. at 0.05 probability). Analysis of comparisons were made based on the value of individual plants. The transgressive segregates were calculated in favourable direction on

Frequency of transgressive segregates in different crosses for yield and its components in the F_2/F_3 generation of 10 interspecific and 8 intraspecific crosses is presented in Table 2.

The transgressive segregates are the result of accumulation of plus or favourable genes affecting yield and its contributing characters. In general, interspecific crosses produced more number of transgressive segregates for number of bolls/plant and kapas yield in both the generations. Four interspecific crosses involving Mdl 2450 and 1875 with Suyog and Jaydhar exhibited higher frequency of transgressants for boll weight in F_3 generation only. For mean halolens, majority of the crosses were failed to produce promising segregates except Mdl 2582 x PA 141 and Mdl 2450 x Raghavendra. On the contrary almost all the crosses gave high number for transgressive segregates ranging from 10 to 93 per cent in F_2 generations for the character ginning outturn, while the frequency of the same was reduced to almost half in F_3 generation.

While four intraspecific crosses involving the parents Mdl 1874 and Mdl 2582 with PA 141 and PA-183 and one interspecific cross Mdl 2463 with Suyog had failed to develop transgressive segregates for the character boll weight, the remaining crosses had transgressive segregates for boll weight from 2 to 5% in F_2 generation. This showed that most of the parents had similar constellation of genes for boll weight. In F_3 however, Mdl 2452 and Mdl 1875 with Jaydhar produced the highest frequency of transgressive segregates i.e. 44% and 37% respectively.

Table 1. Percentage of significant transgressive segregates for yield and yield components in the F₂ and F₃ generations of inter and intra specific crosses of desi cotton

No.	Crosses	Generations	No. of bolls/ plant	Kapas yield/ plant (g)	Boll weight (g)	Halo length (mm)	Ginning outturn (%)
	Mdl 1874 x PA 141	F ₂	83	33	0	0	53
		F ₃	64	85	9	0	0
	Mdl 1874 x PA 183	F ₂	30	7	0	0	80
		F ₃	72	66	2	0	40
	Mdl 1874 x Suyog	F ₂	77	37	7	0	93
		F ₃	75	42	0	0	82
	Mdl 2452 x PA 183	F ₂	47	40	3	7	73
		F ₃	73	46	3	15	27
	Mdl 2452 x Suyog	F ₂	73	43	3	17	73
		F ₃	66	45	26	31	50
	Mdl 2452 x Jaydhar	F ₂	60	40	16	20	64
		F ₃	56	58	44	27	33
	Mdl 2582 x PA 141	F ₂	70	10	0	7	10
		F ₃	72	46	4	76	4
	Mdl 2582 x PA 183	F ₂	20	10	0	0	73
		F ₃	52	48	6	3	63
	Mdl 2582 x Suyog	F ₂	71	48	7	7	65
		F ₃	66	68	0	36	18
10.	Mdl 2450 x PA 141	F ₂	68	49	24	15	73
		F ₃	62	70	4	21	11
11.	Mdl 2450 x PA 183	F ₂	23	43	10	0	90
		F ₃	65	58	10	26	48
12.	Mdl 2450 x Raghavendra	F ₂	71	41	4	37	61
		F ₃	98	97	9	83	20
13.	Mdl 2463 x PA 141	F ₂	31	35	10	0	62
		F ₃	65	72	3	15	30
14.	Mdl 2463 x Suyog	F ₂	59	41	0	0	88
		F ₃	75	68	5	7	66
15.	Mdl 1875 x Suyog	F ₂	83	67	10	0	23
		F ₃	89	96	36	10	5
16.	Mdl 1875 x Raghavendra	F ₂	75	50	5	15	15
		F ₃	90	63	5	0	0
17.	Mdl 1875 x Jaydhar	F ₂	72	50	2	7	22
		F ₃	58	79	37	22	0
18.	Mdl 2562 x Suyog	F ₂	52	29	52	0	93
		F ₃	71	21	7	0	0

Breeding for yield generally aims at recovery of transgressive segregates. All the interspecific crosses had thrown a good number of transgressive segregates for kapas yield per plant ranging from 29 to 67% in the F_2 generation. Similar trend has been repeated in F_3 generation also for this character along with number of bolls per plant.

Highest number of transgressive segregates in the F_3 generation for yield per plant was observed in the intraspecific crosses of Mdl 1874 and Mdl 2450 with the testers PA - 141 and PA-183. Amongst intraspecific crosses Mdl 2452 x PA-183 and Mdl 2450 x PA-141 were promising because in addition to yield per plant these crosses expressed transgressive segregates for important yield associated with fibre components in both the generations.

Transgressive segregates in the F_2 may arise due to dominance and dominance interactions in addition to additive x additive interaction which is fixable, due to recombination of genes

with positive effects and responsible for the production of transgressive segregates in the F_3 generation. The findings, therefore, also revealed that the parents differed for many genes and introgression of genes from herbaceous germplasm lines created large amount of genetic variability for yield and fibre components in most of the crosses suggesting the scope to use this material and the crosses in future breeding programme. Similar results were also reported in lentil by Kant and Singh (1998).

The crosses which gave high frequency of transgressive segregates for yield per plant accompanied with desirable fibre properties in both the generations may be preferred over the other crosses in on-going breeding programme

Reference

- Kant, L. and Singh, D.P. (1998). Transgressive segregation of yield and yield components in lentil. *Indian J. Genet.* 58: 343-347.

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Research Notes

Heterosis for rooting characters in maize (*Zea mays* L.)

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Maize occupies a pride of place among coarse cereals in India. About 80 per cent of the maize belt being rain dependent, experience vagaries of weather in terms of either water logging or drought. Several researchers have suggested that breeding for improved root systems in annual crops such as cereals and grain legumes could significantly improved their yield under drought (Hurd, 1976 and Blum, 1982). Turner *et al.* (1978) stated that the variation in root growth between species determines differences in drought tolerance. Utilisation of genetic variation in improving crop varieties requires knowledge of the heritability, heterosis and genetic control

of root system traits. Present study involves the estimation of heterosis for number of roots, root length and root dry weight in 15 crosses of maize hybrids.

Six outstanding maize inbreds viz. UMI 112, UMI 285, UMI 130, UMI 467, UMI 810 and UMI 90 were chosen as parents, based on grain yield with good agronomic characteristics but differing in genetic background. These lines were crossed in all possible combinations, excluding reciprocals, resulting in deriving of 15 hybrids. The root potential of the six parents and fifteen F_1 progenies were evaluated in the greenhouse