

Combining ability studies for yield and yield components over environments in American cotton (*G. hirsutum* L.)

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Abstract : A 10 x 10 diallel analysis was made to assess the combining ability for yield and yield contributing characters involving divergent varieties of American cotton (*Gossypium hirsutum* L.) at Regl. Agri. Research Station, Lam in four environments. The pooled analysis of variance revealed that GCA x E (except for monopodia per plant and sympodia per plant) and SCA x E (except for number of monopodia per plant) was important for all the characters. Though the magnitude of *gca* variance was considerably higher than the *sca* variance, both additive and non additive genetic variances were important for number of bolls per plant, boll weight and seed cotton yield per plant. Non additive genetic variance was found important for days to 50% flowering, number of monopodia per plant, number of sympodia per plant and number of seeds per boll. The parents NA 1325, NDH 1678 and CWROK 165 were the best general combiners for number of sympodia per plant, number of bolls per plant, boll weight and seed cotton yield per plant. NA 1325 and NDH 1678 also exhibited good general combining ability for days to 50% flowering in desired direction. The crosses NDH 1678 x CWROK 165, L 604 X NDH 1678 and MCU 5 x NA 1325 were identified as the best crosses on the basis of *per se* performance and combining ability.

Key words : Combining ability, Cotton, Additive, Non-additive.

Introduction

Cotton is one of the most important commercial crops and forms the back bone of Indian textile industry. Choice of suitable parents for hybridization and adoption of proper breeding procedure are very important in any crop improvement programme. Combining ability is a powerful tool to discriminate between good and poor general combiners and for choosing appropriate parental lines to produce superior hybrids. The present study therefore was undertaken to obtain information on general and specific combining ability effects and their interaction with environmental variation with respect to yield and yield component characters and the results discussed.

Materials and Methods

The experimental material for the present investigation consisted of 45 F_1 crosses obtained by involving 10 diverse parents *viz.* MCU 5, BN 1, L 389, NA 1325, AC 738, L 604,

NDH 1678, ARB 8824, CWROK 165 and ICMF 82 obtained from different cotton research stations, in a diallel fashion (excluding reciprocals). The 45 F_1 's along with 10 parents were grown in a Randomized Block Design with 3 replication in 4 environments *viz.* E_1 - Normal sowing - Irrigated; E_2 - Normal sowing - Rainfed; E_3 - Delayed sowing - Irrigated and E_4 - Delayed sowing - Rainfed during 1999-2000 season at Regional Agricultural Research Station, Lam. Recommended package of practices for the region were followed. Each genotype was grown in a 2 row plot of 6 meters length adopting a spacing of 120 cm between rows and 60 cm within a row. Five competitive plants were randomly selected for recording data on days to 50 per cent flowering, number of monopodia per plant, number of sympodia per plant, number of bolls per plant, boll weight (g) and number of seeds per boll and seed cotton yield plant⁻¹ (g). The data pooled over environments were subjected to statistical analysis following Griffing (1956)

Table 1. Analysis of variance for combining ability for yield and yield components

Source	d.f.	Days to 50% flowering	No. of monopodia plant ⁻¹	No. of sympodia plant ⁻¹	No. of bolls plant ⁻¹	Boll weight (g)	No. of seeds boll ⁻¹	Seed cotton yield plant ⁻¹ (g)
GCA	9	10.725**	0.583**	6.188**	176.359**	1.178**	10.735**	6825.117**
SCA	45	5.785**	0.165**	3.168**	29.927**	0.273**	4.179**	834.428**
ENVIRONMENTS	3	378.818**	0.317**	347.122**	1158.669**	12.885**	0.875	27819.234**
GCA x E	27	0.678**	0.013	0.267	5.032**	0.046**	1.280**	390.499**
SCA x E	135	0.733**	0.007	0.265*	3.421**	0.040**	1.536**	102.358**
Error	432	0.446	0.012	0.204	1.985	0.018	0.594	30.969
σ^2 gca	0.10	0.01	0.06	3.03	0.02	0.14	119.22	183.02
σ^2 sca		1.26	0.04	0.73	6.63	0.06	0.66	0.57
$2 \sigma^2$ gca / $2 \sigma^2$ sca		0.14	0.30	0.15	0.48	0.40	0.30	

*, ** Significant at 5 and 1% levels, respectively

Table 2. Estimates of general combining ability effects (pooled) for yield and yield components in cotton

Parents	Days to 50% flowering	No. of monopodia plant ⁻¹	No. of sympodia plant ⁻¹	No. of bolls plant ⁻¹	Boll weight (g)	No. of seeds boll ⁻¹	Seed cotton yield plant ⁻¹ (g)
MCU 5	0.33**	0.17**	0.25**	0.08	-0.04**	-0.11	-1.89**
BN I	-0.18**	-0.19**	-0.25**	0.27	-0.29**	0.95**	-6.14**
L 389	0.30**	-0.07**	-0.26**	-1.88**	0.04*	0.07	-7.50**
NA 1325	-0.98**	0.04**	0.39**	2.73**	0.06**	-0.36**	16.08**
AC 738	0.67**	0.04**	-0.07	-2.44**	-0.03	-0.58**	-13.96**
L 604	0.16	-0.04**	-0.12*	1.11**	-0.15**	0.65**	0.59
NDLH 1678	-0.39**	0.03**	0.26**	1.45**	0.22**	-0.02	16.04**
ARB 8824	-0.12	-0.04**	0.14*	0.04	-0.03	-0.39**	-1.64**
CWROK 165	-0.18**	-0.10**	0.39**	1.74**	0.24**	-0.22**	14.49**
ICMF 82	0.38**	0.15**	-0.75**	-3.11**	-0.02	0.01	-16.07**
SE (gi)	0.09	0.01	0.06	0.19	0.02	0.11	0.76
SE (gi-gj)	0.14	0.02	0.09	0.29	0.03	0.16	1.14

*, ** Significant at 5 and 1% levels, respectively

Table 3. Crosses having high mean performance and significant positive specific combining ability effects

Character	Mean Performance	SCA effects
1. Days to 50 flowering	NA 1325 X L 604 AC 738 X CWROK 165 L 389 X NA 1325	AC 738 X CWROK 165 NA 1325 X L 604 MCU 5 X NDH 1678
2. No. of monopodia plant ⁻¹	MCU 5 X ICMF 82 MCU 5 X L 604 NA 1325 X ICMF 82	BN 1 X ICMF 82 MCU 5 X L 604 L 604 X ARB 8824
3. No. of sympodia plant ⁻¹	NDH 1678 X CWROK 165 NA 1325 X L 604 NA 1325 X AC 738	NA 1325 X L 604 NDH 1678 X CWROK 165 NA 1325 X AC 738
4. No. of bolls plant ⁻¹	MCU 5 X NA 1325 BN 1 X AC 738 NDH 1678 X CWROK 165	BN 1 X AC 738 L 389 X NDH 1678 MCU 5 X NA 1325
5. Boll weight (g)	AC 738 X NDH 1678 NDH 1678 X CWROK 165 AC 738 X CWROK 165	AC 738 X NDH 1678 BN 1 X ARB 8824 MCU 5 X ARB 8824
6. No. of seeds boll ⁻¹	BN 1 X NDH 1678 MCU 5 X CWROK 165 L 604 X ICMF 82	MCU 5 X CWROK 165 BN 1 X NDH 1678 ARB 8824 X CWROK 165
7. Seed cotton yield plant ⁻¹ (g)	NDH 1678 X CWROK 165 L 604 X NDH 1678 MCU 5 X NA 1325	BN 1 X AC 738 MCU 5 X NA 1325 L 604 X NDH 1678

and as extended by Singh (1973, 1979) using Method - 2, Model - 1.

Results and Discussion

The analysis of variance for combining ability over environments (Table 1) revealed highly significant differences among the genotypes for all the traits. The pooled analysis also revealed that environments were found to be significant for all the characters and had highly significant interaction variance involving *gca* x environments (except for No. of monopodia and no. of sympodia) and *sca* x environments (No. of monopodia), which revealed that both *gca* effects of parents and *sca* effects of the hybrids were influenced by the sampled environments. The magnitude of the former was greater than the latter for majority of the traits suggesting greater influence of environment on *gca* than *sca* variance. Similar results for number of bolls plant⁻¹, boll weight and seed

cotton yield per plant were earlier reported by Pavasia *et al.* (1998).

The combining ability analysis is frequently used for testing the performance of parent and hybrid combinations and also for understanding the nature and magnitude of gene action involved in the expression of different traits beside isolating good general combiners for use in breeding programmes. Gupta and Singh (1974) Singh *et al.* (1988) and Pavasia *et al.* (1998) reported significant relationship between the *per se* performance and *gca* effects of the parents in upland cotton. Similar relationship was also found in the present study between *per se* performance and *gca* effects of the parents for days to 50 per cent flowering, number of monopodia plant⁻¹, number of seeds per boll and seed cotton yield per plant. Whereas such relationship was not observed for number of sympodia per plant; number of boll

plant⁻¹ and boll weight. Therefore, it can be stated for these characters that the ability of the parents to nick well can only be judged from combining ability studies.

The relative importance of *gca* and *sca* variance was assessed by the ratio $2\sigma^2_{gca}/\sigma^2_{gca}+2\sigma^2_{sca}$ as suggested by Baker (1978). The estimates of this ratio indicated that the relative magnitude of non-additive gene effects over additive gene effects for days to 50% flowering, number of monopodia plant⁻¹, number of sympodia plant⁻¹ and number of seeds bolls⁻¹. Both additive and non-additive gene actions were observed for number of boll per plant, boll weight and seed cotton yield per plant. So far as earlier findings regarding nature of gene action is concerned, Shanti and Selvaraj (1995), Surana *et al.* (1997), Murthy and Ranganathacharyulu (1998), Valarmathi and Jehangir (1998) have reported the predominance of non-additive gene action for number of sympodia/plant. Murthy and Ranganathacharyulu (1998) and Pavasia *et al.* (1998) reported the importance of both additive and non-additive gene action for number of bolls plant⁻¹, boll weight and seed cotton yield per plant.

The estimates of *gca* effects of the parents are presented in Table 2. The parents NA 1325, NDLH 1678 and CWROK 165 showed significant *gca* effects for seed cotton yield. These parents also showed significant *gca* effects for days to 50% flowering, number of monopodia plant⁻¹, number of sympodia plant⁻¹, number of bolls plant⁻¹ and boll weight. The parent NDLH 1678 also recorded significant *gca* effects for number of monopodia plant⁻¹, number of bolls plant⁻¹ and boll weight besides seed cotton yield. The parents NA 1325, NDLH 1678 and CWROK 165 were adjudged as good general combiners for and yield component traits. The *gca* of yield was influenced by *gca* of yield contributing characters.

The estimates of *sca* effects of top performing crosses for different traits are presented in Table 3. The cross NDLH 1678 X CWROK

165 recorded high *sca* effects for seed cotton yield, number of sympodia plant⁻¹, number of bolls plant⁻¹ and boll weight. The cross MCU 5 X NA 1325 showed significant *sca* effects for number of sympodia plant⁻¹, number of bolls plant⁻¹ and seed cotton yield. The crosses L 389 X NA 1325 and MCU 5 X NA 1325 also showed significant *sca* effects for days to 50% flowering in the desired direction. The cross NDLH 167 X CWROK 165, which had high *sca* effects for seed cotton yield and various yield components *viz.* number of sympodia plant⁻¹, number of bolls plant⁻¹ and boll weight also possessed significant and high *gca* effects indicating *gca* is an indicator for specific cross effects. This close relationship may be due to predominance of additive and non-additive gene action for number of bolls plant⁻¹ and boll weight and seed cotton yield.

In the case of crosses, which exhibited significant *sca* effects in the pooled data, the grading of the parents based upon their *gca* effects indicated that for number of monopodia/plant, number of seeds boll⁻¹ and seed cotton yield, some of the crosses involved both the parents with poor combining ability. The superiority of the poor x poor combiners may be attributed to the genetic diversity of the parents. The cross NDLH 1678 X CWROK 165, which involved H X H general combiners is expected to release desirable segregants in the subsequent generations, since these involve additive and additive x additive interactions. The crosses L 604 X NDLH 1678 and MCU 5 X NA 1325, which involve high x low or low x high general combiners are likely to produce desirable transgressive segregants. El-Adl and Miller (1971) obtained transgressive segregants that were better than the original F₁. Kalsy and Garg (1980) and Gururajan and Basu (1992) also stressed the importance of H x L crosses in obtaining superior combinations.

The present study indicated that the parents NA 1325, NDLH 1678 and CWROK 165, which had high *gca* effects for different economic characters can be utilized in conventional breeding

programme and the crosses NDH 1678 X CWROK 165, L 604 X NDH 1678 and MCU 5 X NA 1325 with high sea can be exploited for hybrid vigour. Both additive and non-additive components of variance was found important for number of bolls plant⁻¹, boll weight and seed cotton yield. Under such situation, recurrent selection methods would be appropriate for rapid improvement of yield. The relatively high interaction of non additive genetic variance with the environments suggested that specific hybrids may be identified for specific situations.

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