

Research Article

General and specific combining ability for quantitative characters in castor (*Ricinus communis* L.)

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Abstract

The experimental material consisted of ten monoecious lines, their 45 F_1S developed through diallel mating excluding reciprocals, which was evaluated in Randomized Complete Block Design with three replications. The analysis of variance for combining ability revealed that mean square values due to both GCA and SCA and were significant, accordingly the estimates of both σ^2_{GCA} and σ^2_{SCA} were also significant for all the characters indicating an importance of both additive and non-additive gene effects for the inheritance of studied characters. However, values of potence ratio and predictability ratio revealed preponderance of additive genetic variance for all the characters. Parents ANDCI 10-12, ANDCI 8, ANDCI 10-4 and ANDCI 10-3 were good general combiners for seed yield per plant and other yield contributing characters. The hybrids ANDCI 10-1 x ANDCI 9, ANDCI10-11 x ANDCI 10-12 and ANDCI 8 x ANDCI 10-11, had high *per se* performance, high heterotic effects and desired sca effects for seed yield per plant and its component characters.

Key words

Castor, Ricinus communis, gca, sca, additive and non-additive, combining ability, potence ratio

Introduction

Castor (Ricinus communis L.) is industrially an important non-edible oilseed crop widely cultivated in arid and semi-arid regions of the World (Govaerts et al., 2000). At present, India is a world leader in castor production and main exporter of castor oil and its derivatives. India contributes 65 per cent of the world production and meets about 90 per cent world castor oil demands. In India during 2015-16, Castor occupied an area about 10.61 lakh ha with a production of 17.82 lakh tonnes. Interestingly, an average seed yield has increased from 1334 kg/ha in 2009-10 to 1752 kg/ha in 2015-16, which is highly encouraging (Anonymous, 2017). Castor is usually cultivated as a hybrid in India, as hybrids give significantly higher yields than varieties (Moll et al., 1962; Birchler et al., 2003; Reif et al., 2007). Higher magnitude of heterotic effects and superior hybrids can be developed by combining diverse parents. In hybrids development programme availability of monoecious inbreds as pollen parents mandatory.

In genetic improvement of different attributes of castor, selection of suitable parents is preliminary approach, which will lead to development of better hybrids, for that some indication can be obtained from *per se* performance of parents, but information based on *per se* performance of parents along with their general combining ability effect is more reliable. However, information on magnitude of heterotic effects, specific combining ability effect of hybrids, and type of gene action involved for inheritance of seed yield, and its component characters would be of immense value in selecting desired parents, and beneficial cross combinations for commercial exploitation of

hybrid vigour, and also in formulating appropriate future breeding programme of castor.

Material and Methods

The experimental material consisted of ten monoecious lines and their 45 F₁s developed through diallel mating excluding reciprocals. The experiment was conducted in a Randomized Complete Block Design with three replications at Regional Research Station, Anand Agricultural University, Anand during 2012-13. Each test genotype was grown in a single row of 10 plants with 120 cm inter row distance and 60 cm intra row distance. All the recommended agronomic practices and plant protection measures were followed to raise a good crop. The observations for all the characters under study except days to 50% flowering and days to 50% maturity of primary raceme were recorded on five randomly selected competitive plants and an average value per plant for each replication was worked out, whereas days to 50% flowering and days to 50% maturity of primary raceme were recorded on population basis. The mean values were subjected for statistical analysis as suggested by Snedecor and Cochran (1937). Combining ability analysis was performed as per Griffing (1956) Model-I andmethod-II. Estimation of relative heterosis and heterobeltosis analysis was carried out as per Turner (1953) and Fonseca and Patterson (1968).

Results and Discussion

Combining ability analysis is a powerful tool to select good combiners and for choosing desired parental material in crop breeding program. The *per se* performance of a given parent does not necessarily mean that it would be good or poor general combiner. Therefore, gathering



information on nature of gene effects and their expression in terms of combining ability is necessary. Significate value of GCA and SCA variance for a character indicates existence of additive and non-additive genetic effects, respectively. However, both variance are non significant, epistatic gene effects suppose to play an important role in determining inheritance of a character (Fehr, 1993).

The analysis of variance for combining ability (Table 1) revealed that mean square values due to GCA and SCA were significant, accordingly the estimates of both σ^2_{GCA} and σ^2_{SCA} were significant for all the characters, suggesting an importance of both additive and non-additive genetic variances for the inheritance of all the characters under investigation. Whereas, above one values of potence ratio and above one half value of predictability ratio indicated preponderance of additive genetic variance for all the characters except days to 50% maturity of primary raceme and shelling out turn. The results are in conformity with the findings of Patel (2010), Patel et al. (2012), Ramesh et al. (2013), Chaudhari and Patel (2014) and Punewar et al. (2017) as they reported importance of additive genetic variance for seed yield per plant and other important seed yield component characters. Although there was preponderance of additive gene action for all the characters, but presence of a considerable amount of non-additive gene effect could not be totally neglected. The estimates of average degree of dominance for all the characters except days to 50% maturity of primary raceme, shelling out turn were less than one revealing existence of partial dominance behavior of interacting alleles.

The perusal of the results (Table:2) in respect to gca effect of parents revealed that for seed yield per plant, parents ANDCI 10-12, ANDCI 8, ANDCI 10-4, ANDCI 10-3 and SKI 215 were good general combiners and parent JI 360 was average general combiner. Among good general combiners for seed yield, ANDCI 8 was the good general combiner for effective length of primary raceme, while it was good general combiner for number of capsules on primary raceme, test weight and oil content; parent ANDCI 10-12 was also good general combiner for days to 50% flowering of primary raceme, length of primary raceme, effective length of primary raceme, number of capsules on primary raceme and days to 50% maturity of primary raceme, while it was average general combiner for shelling out turn and test weight.

Among the good general combiners for seed yield, ANDCI 10-4 was also good general combiner for effective length of primary raceme and number of capsule on primary raceme, ANDCI 10-3 was also good general combiner for days to 50% flowering of primary raceme, number of effective branches per plant, length of primary spike, effective length of primary spike and number of capsules on primary raceme and SKI 215 was also good general combiner for number of effective branches per plant, test weight and oil content.

The top ranking three parental genotypes on the basis of their per se performance and general combing ability effects as well as crosses in respect to per se performance, specific combining ability, relative heterosis and heterobeltiosis for seed yield per plant and its component characters are presented in table 3. The per se performance of parents along with their gca effect could be a better criteria for selection of superior parent/s in future breeding programme. In present investigation, the results revealed that the most of the parents had relatively high degree of correspondence between per se performance and their gca effects for almost all the characters, which could be ascribed to existence of genes, which showed additivity. Therefore, in selection of parents for varietal development programme due weightage should also be given to per se performance along with their *gca* effect.

The estimates of specific combining ability effect provide information on role of intra and interallelic interactions in the expression of heterotic effects and inheritance of a character. The results revealed that crosses, which had high *per se* performance also depicted higher heterotic effects and high estimate of *sca* effect for all the growth and developmental attributes.

Among the evaluated crosses, none of the cross combinations had desirable significant *sca* effects simultaneously for all the characters under study; however, good specific combiners for seed yield also depicted desirable *sca* effects for most of the yield component characters.

In respect to gca effect of parents involved in a particular cross, crosses could be grouped in to resultant of six different categories of good, average and poor general combiner parents viz., G x G, G x A, A x A, A x P and P x P. In general, the crosses, which exhibited high sca effect did not always involved both good general combining parents with high gca effect, there by suggesting importance of intra as well as inter-allelic interactions. The high sca effect of crosses in general corresponded to their high heterotic response, but these might also be accompanied by poor and/or average gca effect of the parents. For seed vield per plant, total five crosses exhibited significant positive sca effect; and out of ten parents, seven parents involved in these crosses, of which four parents viz. ANDCI-8, ANDCI 10-04, ANDCI 10-3 and ANDCI 10-12 were good general combiners, and three parents viz., ANDCI 10-1,



ANDCI 10-11 and ANDCI 9 were poor general combiners, therefore, cross combinations were of resultant of P x P, G x P and G x G *gca* effect of parents, and high *sca* or heterotic effects could be because of intra and inter allelic interactions.

Among the crosses, which depicted significant and positive *sca* effect for seed yield per plant, crosses ANDCI 10-1 x ANDCI 9 and ANDCI 10-12 x ANDCI 10-11 were good specific combiner for days to 50 % flowering of primary raceme, days to 50 % maturity of primary raceme, number of nodes up to primary raceme, , and plant height up to primary raceme .The cross ANDCI 10-12 x ANDCI 10-11 was also good specific combiner for number of effective branches per plant, effective length of primary raceme, number of capsules on primary spikes, and oil content, while it was average specific combiner for rest of the characters.

The cross ANDCI-8 x ANDCI 10-11 was good specific combiner for number of effective branches per plant, effective length of primary raceme, test weight and oil content. The crosses which exhibited high sca effect for seed yield per plant also registered desirable sca effect for at least two yield component characters, but those might not necessarily have higher sca effect for the said characters, which suggested cumulative effect of various yield contributing attributes towards high sca effect for seed yield, and thereby high heterotic effects as well. This also appropriately suggest that yield is a complex character dependent on number component characters, and suitable of recombination of genes governing these characters would have produced promising crosses, therefore, none of the crosses had desirable sca effect for all the characters under study.

The monoecious parents ANDCI 10-12, ANDCI 8, ANDCI 10-4, ANDCI 10-3 and SKI 215 were good general combiners for seed yield per plant and other contributing characters suggesting these parental lines need to be used extensively as pollen parents in heterosis breeding and for making normal crosses and those crosses may be advanced for development of superior recombinants to strengthens monoecious gene pool and thereby for development of promising inbreds as improved varieties and/or male parents. Similarly, the cross combinations, ANDCI 10-1 x ANDCI 9, ANDCI10-11 x ANDCI 10-12 and ANDCI 8 x ANDCI 10-11, which showed high per se performance, high heterotic effects and desired sca effects for seed yield per plant and its component characters including developmental characters need to be advanced for development of inbreds having short plant stature, earliness, profused branches and resistant to major biotic stresses.

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Source of variation	d.f.	Days to 50 % flowering of primary raceme	Plant height up to primary raceme	No. of nodes up to primary raceme	No. of effective branches per plant	Effective length of primary raceme	No. of capsules on primary raceme	Days to 50 % maturity of primary raceme	Shelling out turn	Test weight	Seed yield per plant	Oil content
Mean squares												
GCA	9	59.18**	1109.88**	12.21**	40.35**	346.22**	867.66**	160.77**	5.47**	31.80**	2933.43**	11.90**
SCA	44	7.25**	46.57**	0.35**	4.56**	32.51**	63.93**	28.95**	1.61**	2.31**	309.64**	1.15**
Error	106	0.81	10.32	0.10	0.77	5.66	12.02	2.544	0.91	0.200	134.77	0.09
Estimates of co	mponen	ts of genetic varia	nce and relate	d parameters								
$\sigma^2_{GCA} (\sum g i^2)$		4.86**	91.63**	1.01**	3.29**	28.38**	71.30**	15.42**	0.37**	2.63**	233.22**	0.98**
σ^2_{SCA} ($\sum sij^2$)		6.44**	36.25**	0.25**	3.79**	26.84**	51.91**	26.45**	0.69**	2.11**	174.86**	1.05**
Potence ratio		3.69	12.35	19.75	4.24	5.17	6.72	2.85	2.62	6.09	6.52	4.56
Predectibility ra	itio	0.60	0.83	0.88	0.63	0.67	0.73	0.54	0.52	0.71	0.72	0.65
$\sigma^2 A$		9.73	183.26	2.01	6.59	56.76	142.60	30.83	0.75	5.26	466.44	1.96
$\sigma^2 D$		6.44	36.25	0.25	3.79	26.84	51.91	26.45	0.69	2.11	174.86	1.05
$(\sigma^2 D/\sigma^2 A)^{0.5}$		0.81	0.44	0.35	0.75	0.69	0.60	0.93	0.95	0.63	0.61	0.73

Table 1. Analysis of variance for combining ability for various characters of castor

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



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Table 2. Estimates of general combining ability (gca) effect of parents for various characters of castor

Parents	Days to 50 % flowering of primary raceme	Plant height up to primary raceme	No. of nodes up to primary raceme	No. of effective branches per plant	Effective length of primary raceme	No. of capsules on primary raceme	Days to 50 % maturity of primary raceme	Shelling out turn	Test weight	Seed yield per plant	Oil content
SKI 215	3.11**	10.17**	0.83**	3.00**	-2.30**	-7.71**	6.77**	0.40	1.92**	6.25*	1.07**
JI 360	-0.92**	-5.27**	-0.30**	-1.65**	-0.57	-4.52**	2.27**	1.25**	2.84**	-2.39	0.39**
ANDCI 8	0.50*	5.97**	0.81**	-1.19**	9.14**	8.84**	-0.39	0.41	0.61**	16.72**	0.28**
ANDCI 10-04	0.94**	8.90**	0.76**	-0.03	4.17**	9.52**	-0.84	-0.75**	-1.62**	10.72**	-0.57**
ANDCI 10-3	-0.72**	1.30	0.21*	1.57**	2.12**	6.62**	-0.42	-0.04	-2.04**	7.86*	0.13
ANDCI 10-12	-0.53*	8.88**	0.55**	-1.24**	3.69**	12.27**	-5.06**	-0.06	0.15	17.45**	-1.09**
ANDCI 10-1	2.03**	-10.92**	0.21*	-1.19**	-0.66	-4.21**	4.82**	0.59*	0.26*	-6.63*	0.82**
ANDCI 10-11	-5.06**	-19.63**	-2.50**	-2.48**	-11.39**	-12.08**	-5.28**	-0.91**	-1.34**	-35.21**	-2.12**
ANDCI 9	1.33**	0.97	0.14	2.06**	-1.86**	-2.64**	1.30**	-0.66*	-1.57**	-6.61*	0.90**
ANDCI 1	-0.69**	-0.35	-0.71**	1.16**	-2.34**	-6.10**	-3.17**	-0.22	0.78**	-8.16*	0.18*
Range of GCA	-5.06	-19.63	-2.50	-2.48	-11.39	-12.08	-5.28	-0.91	-2.04	-35.21	-2.12
effects	3.11	10.17	0.83	3.00	9.14	12.27	6.77	1.25	2.84	17.45	1.07
S.E (gi)±	0.24	0.88	0.08	0.24	0.65	0.94	0.43	0.26	0.12	3.17	0.08
S.E. $(gi - gj) \pm$	0.36	1.31	0.13	0.35	0.97	1.41	0.65	0.39	0.18	4.73	0.12
C.D. 0.05 (gi)	0.47	1.72	0.15	0.47	1.27	1.84	0.84	0.50	0.23	6.21	0.15
C.D. 0.05 (gi - gj)	0.71	2.57	0.25	0.69	1.90	2.76	1.27	0.76	0.35	9.27	0.23

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



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Table 3. Top three crosses with respect to their *per se* performance, RH, HB and sca effect and *per se* performance of parents for various characters in castor

Characters	<i>Per se</i> performance	GCA effects	Per se performance	Relative heterosis	Heterobeltiosis	SCA effects
Days to 50 %	ANDCI 10-11	ANDCI 10-11	ANDCI 10-12 x ANDCI 10-11	ANDCI 10-12 x ANDCI 10-11	JI 360 x ANDCI 1	ANDCI 10-12 x ANDCI 10-11
flowering of	JI 360	JI 360	ANDCI 10-04 x ANDCI 10-11	JI 360 x ANDCI 1	ANDCI 10-3 x ANDCI 1	JI 360 x ANDCI 1
primary raceme	ANDCI 10-12	ANDCI 10-3	ANDCI 10-3 x ANDCI 10-11	ANDCI 10-3 x ANDCI 1	JI 360 x ANDCI 10-3	ANDCI 10-04 x ANDCI 10-11
Plant height up to primary raceme	ANDCI 10-11	ANDCI 10-11	ANDCI 10-3 x ANDCI 10-11	ANDCI 10-12 x ANDCI 10-11	ANDCI 8 x ANDCI 10-12	ANDCI 10-12 x ANDCI 10-11
	ANDCI 10-1	ANDCI -10-1	ANDCI 10-12 x ANDCI 10-11	ANDCI 8 x ANDCI 10-12	SKI 215 x ANDCI 10-04	ANDCI 8 x ANDCI 10-12
	JI 360	JI 360	ANDCI 10-11 x ANDCI 1	JI 360 x ANDCI 8	ANDCI 8 x ANDCI 10-04	SKI 215 x ANDCI 10-04
	ANDCI 10-11	ANDCI 10-11	ANDCI 10-11 x ANDCI 1	ANDCI 10-1 x ANDCI 1	ANDCI 10-1 x ANDCI 9	SKI 215 x ANDCI 10-11
Number of nodes up to primary raceme	JI 360	ANDCI 1	SKI 215 x ANDCI 10-11	ANDCI 10-11 x ANDCI 1	ANDCI 10-1 x ANDCI 1	ANDCI 10-1 x ANDCI 9
to primary facefile	ANDCI 1	JI 360	ANDCI 10-12 x ANDCI 10-11	ANDCI 10-1 x ANDCI 9	SKI 215 x ANDCI 1	SKI 215 x ANDCI 1
	SKI 215	SKI 215	SKI 215 x ANDCI 9	ANDCI 8 x ANDCI 10-11	ANDCI 8 x ANDCI 10-11	ANDCI 10-04 x ANDCI 10-1
Number of effective branches per plant	ANDCI 9	ANDCI 9	ANDCI 9 x ANDCI 1	ANDCI 10-12 x ANDCI 10-11	ANDCI 10-04 x ANDCI 10-1	ANDCI 10-12 x ANDCI 10-11
branches per plant	ANDCI 10-3	ANDCI 10-3	ANDCI 10-04 x ANDCI 10-3	ANDCI 10-04 x ANDCI 10-1	JI 360 x ANDCI 8	ANDCI 8 x ANDCI 10-11
Effective length of primary raceme	ANDCI 8	ANDCI 8	ANDCI 8 x ANDCI 10-3	ANDCI 10-04 x ANDCI 10-11	JI 360 x ANDCI 10-3	ANDCI 10-04 x ANDCI 10-11
	ANDCI 10-04	ANDCI 10-04	ANDCI 10-3 x ANDCI 10-12	ANDCI 10-12 x ANDCI 10-11	SKI 215 x ANDCI 10-3	ANDCI 10-04 x ANDCI 10-1
	ANDCI 10-12	ANDCI 10-12	ANDCI 8 x ANDCI 10-12	JI 360 x ANDCI 10-11	ANDCI 10-3 x ANDCI 10-12	ANDCI 10-12 x ANDCI 10-11
	ANDCI 8	ANDCI 10-12	ANDCI 10-3 x ANDCI 10-12	ANDCI 10-12 x ANDCI 10-11	SKI 215 x ANDCI 1	ANDCI 10-12 x ANDCI 10-11
Number of capsules on primary raceme	ANDCI 10-12	ANDCI 10-04	ANDCI 8 x ANDCI 10-3	ANDCI 10-04 x ANDCI 1	ANDCI 10-3 x ANDCI 10-12	JI 360 x ANDCI 10-12
	ANDCI 10-04	ANDCI 8	ANDCI 10-12 x ANDCI 10-11	SKI 215 x ANDCI 1	ANDCI 9 x ANDCI 1	ANDCI 10-1 x ANDCI 10-11
Days to 50 %	ANDCI 10-12	ANDCI 10-11	ANDCI 10-12 x ANDCI 10-11	ANDCI 8 x ANDCI 10-12	ANDCI 8 x ANDCI 10-1	ANDCI 8 x ANDCI 10-12
maturity of primary	ANDCI 10-11	ANDCI 10-12	ANDCI 8 x ANDCI 10-12	ANDCI 8 x ANDCI 10-1	JI 360 x ANDCI 8	ANDCI 10-04 x ANDCI 10-11
raceme	ANDCI 1	ANDCI 1	ANDCI 10-04 x ANDCI 10-11	JI 360 x ANDCI 8	ANDCI 8 x ANDCI 9	ANDCI 10-12 x ANDCI 10-11
	JI 360	JI 360	JI 360 x ANDCI 1	ANDCI 10-11 x ANDCI 1	ANDCI 10-11 x ANDCI 1	ANDCI 10-11 x ANDCI 9
Shelling out turn	ANDCI 10-1	ANDCI 10-1	ANDCI 10-3 x ANDCI 10-1	SKI 215 x ANDCI 1	SKI 215 x ANDCI 1	ANDCI 10-04 x ANDCI 10-3
-	ANDCI 8	ANDCI 8	JI 360 x ANDCI 10-1	JI 360 x ANDCI 1	ANDCI 10-11 x ANDCI 9	ANDCI 10-3 x ANDCI 10-1
	JI 360	JI 360	SKI 215 x ANDCI 1	ANDCI 8 x ANDCI 10-3	SKI 215 x ANDCI 1	ANDCI 8 x ANDCI 10-12
Test weight	ANDCI 1	SKI 215	JI 360 x ANDCI 10-1	ANDCI 8 x ANDCI 10-11	ANDCI 8 x ANDCI 10-3	SKI 215 x ANDCI 1
	SKI 215	ANDCI 1	SKI 215 x JI 360	SKI 215 x ANDCI 1	ANDCI 10-12 x ANDCI 10-1	ANDCI 10-11 x ANDCI 9
Seed yield per plant	ANDCI 10-12	ANDCI 10-12	ANDCI 8 x ANDCI 10-04	ANDCI 10-1 x ANDCI 9	ANDCI 10-1 x ANDCI 9	ANDCI 10-1 x ANDCI 9
	ANDCI 8	ANDCI 8	ANDCI 8 x ANDCI 10-3	ANDCI 10-12 x ANDCI 10-11	ANDCI 8 x ANDCI 10-04	ANDCI 10-12 x ANDCI 10-11
	SKI 215	ANDCI 10-04	ANDCI 8 x ANDCI 10-12	ANDCI 8 x ANDCI 10-11	ANDCI 8 x ANDCI 10-3	ANDCI 8 x ANDCI 10-04
	SKI 215	SKI 215	SKI 215 x ANDCI 9	ANDCI 10-11 x ANDCI 1	ANDCI 10-11 x ANDCI 1	ANDCI 10-11 x ANDCI 1
Oil content	ANDCI 8	ANDCI 9	JI 360 x ANDCI 10-1	ANDCI 10-12 x ANDCI 10-11	ANDCI 10-04 x ANDCI 1	ANDCI 10-12 x ANDCI 10-11
	ANDCI 9	ANDCI 10-1	ANDCI 10-1 x ANDCI 10-11	JI 360 x ANDCI 10-12	JI 360 x ANDCI 1	JI 360 x ANDCI 10-12