# Combining ability and its relationship with gene action in slicing cucumber

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### ABSTRACT

A study was conducted during 2005-06 and 2006-07 on the combining ability analysis in a 6 × 6 half-diallel cross of cucumber (Cucumis sativus L.) excluding reciprocals. The 15 F, hybrids along with six parental lines were grown in randomized block design with three replications. The mean square due to gca and sca were highly significant for almost all the characters indicating the importance of both additive and non-additive genetic components for the characters under study. The predictability ratio and average degree of dominance was observed to be less than 0.5 and more than 1 for important yield contributing characters, viz., earliness, number of fruits per plant and total yield per plant suggesting the importance of non-additive components of variance for improvements of these characters. The parents P<sub>1</sub>, P<sub>2</sub>, P<sub>5</sub> and P<sub>6</sub> were observed to be good combiners for a number of characters, including yield per plant. The crosses,  $P_1 \times P_5$ ,  $P_2 \times P_5$  and  $P_2 \times P_5$  were most promising combinations for earliness and other desirable characters including yield per plant. The results indicated the importance of heterosis breeding for effective utilization of non-additive genetic variance in cucumber.

Key words: Cucumber, combining ability, gca, sca, predictability ratio.

## INTRODUCTION

India being native place of cucumber (Cucumis sativus L.) possesses high genetic variability for vegetative and fruit characters. Low fruiting ability and yield suppression due to its inherent fruiting habits are the major factors limiting fruit yield in slicing and processing cucumber (Lower et al., 5). Development of high yielding varieties mainly depends upon genetically superior parents, coupled with suitable breeding methodology. Combining ability analysis is one of the powerful tools available which give the estimates of combining ability effects and aids in selecting desirable parents and crosses for further exploitation. Such studies in cucumber are very few. Therefore, the present investigation was undertaken to identify the best combiners among the existing germplasm as well as to study the gene action of different quantitative characters in 6 × 6 half-diallel set for formulation of a sound breeding programme in cucumber.

#### MATERIALS AND METHODS

The experiment was carried out at Indian Agricultural Research Institute during summer seasons of 2005-06 and 2006-07. Six genetically diverse inbreds of cucumber, viz., CRC-8 (P1), CHC-2 (P2), G-338 ( $P_{a}$ ), CH-20 ( $P_{a}$ ), Pusa Uday ( $P_{a}$ ) and DC-1 ( $P_{a}$ ) were crossed in  $6 \times 6$  half-diallel (excluding reciprocal)

mating scheme (Hayman, 4). The resulting 15 F. hybrids along with 6 parental lines were evaluated in a randomized block design with three replications. The seeds were sown in rows of 1.5 m with 0.50 m spacing between the plants. All the recommended package of practices was followed to grow a successful crop. Out of 12 plants, 10 were marked for taking observations. Observations on individual plant basis were recorded on eight quantitative characters, viz., days to first female flower opening, node number of first female flower, days to first fruit harvest, fruit weight (g), number of fruits per plant, fruit length (cm), fruit diameter (cm) and total yield per plant (g). The combining ability estimates were calculated according to Method-2, Model 1 (Griffing, 2). For the gene action, relative importance of general combining ability (gca) and specific combining ability (sca) was estimated by predictability ratio 2  $\sigma^2 g/2 \sigma^2 g + \sigma^2 s$  (Baker, 1) for fixed effect model, where  $\sigma^2 \tilde{g}$  is the additive component of variance and  $\sigma^2$ s is the non-additive component of variance. Average degree of dominance was estimated by using formula  $\sigma^2$ s / 2  $\sigma^2$ g.

## **RESULTS AND DISCUSSION**

The mean square due to gca and sca were highly significant for all the characters indicating the importance of both additive and non-additive genetic components for the characters under study (Table 1). Sca components of variance was higher than gca components of variance for five traits, viz., days to first female flower opening, node number of first female

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Source	d.f.	Days to 1 <sup>st</sup>	Node No. of	Days to first	Fruit weight	No. of fruits	Fruit	Fruit	Total yield per
		female flower	1 <sup>st</sup> female	fruit harvest	(g)	per plant	length	dia.	plant
		opening	flower				(cm)	(cm)	(g)
gca	5	36.97**	12.08**	51.70**	276.12**	6.92**	7.39**	1.31**	670790.35**
sca	15	12.09**	3.12**	14.66**	30.68**	1.56**	0.91**	0.21**	254584.35**
Error	40	0.55	0.02	0.17	1.82	0.06	0.24	0.03	1739.17
σ <sup>2</sup> g		3.11	1.12	4.63	34.46	0.67	0.83	0.11	52025.75
σ <sup>2</sup> s		11.54	3.10	14.49	32.51	1.62	0.67	0.18	252845.18
PR		0.35	0.42	0.39	0.68	0.46	0.71	0.61	0.14
Add		1.81	1.36	1.68	0.82	1.21	0.41	0.90	2.43

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

flower, days to first fruit harvest, number of fruits per plant and total yield per plant while gca components of variance was higher than sca components of variance for three traits, viz., fruit weight, fruit length and fruit diameter which indicated the predominance of both and non-additive gene action for the characters studied. The relevant results regarding relative importance of gca and sca (predictability ratio) observed to be < 0.5 for days to first female flower opening, node number of first female flower, days to first fruit harvest, number of fruits per plant and total yield per plant, thus reflecting the predominant role of non-additive components of variance for improvement of these characters. Average degree of dominance which was found to be >1 for these characters further confirmed the results. Predictability ratio and average degree of dominance was observed to be >0.5 and <1 for fruit weight, fruit length and fruit diameter, indicating the predominance of additive gene action for improvement of these characters. Among the six parental lines, the parent P1 showed highest gca for days to first female flower opening (-1.98), node number of first female flower (-0.69), days to first fruit harvest (-1.97); P<sub>6</sub> for fruit weight (13.82), number of fruits per plant (0.46), fruit length (1.20);  $P_{f}$  for fruit diameter (0.31) and total yield per plant (328.41) (Table 2). In most of the cases it was observed that per se performance of parents bear direct reflection of their respective gca effects, i.e. parents showing highest gca effects for a character, were also observed to be good performer with respect to that particular character. The results are in conformity with the findings of Munshi et al. (7), Yudhvir and Sharma (15), and Sarkar and Sirohi (11) in cucumber.

The number of crosses having significant sca effects were 10 for days to first female flower opening, 8 for node number of first female flower, 8 for days to first fruit harvest, 8 for fruit weight, 6 for number of fruits per plant, 9 for fruit length, 7 for fruit diameter and 7 for total yield per plant. The crosses showing highest significant sca effects for various characters in order of merit were  $P_1 \times P_5$  for days to first female flower opening and total yield per plant; P<sub>5</sub> × P<sub>6</sub> for node number of first female flower;  $P_2 \times P_5$  for days to first fruit harvest, number of fruits per plant and fruit length;  $P_1 \times P_6$  for fruit weight and  $P_2 \times P_6$  for fruit diameter. In most of the cases, these 5 hybrids which showed best per se performance also possessed desirable significant sca effects (Table 3). This indicated the per se performance of hybrids had a direct relation with respective sca effects. A comparison of the sca effects of the crosses and gca effects of the parents involved indicated that in most of the cases gca effects were reflected in the sca effects of the cross combination. It is apparent that in almost all the hybrids which showed the best sca effects, the parental lines involved were at least one of the four outstanding parental lines, viz., CRC-8 ( $P_1$ ), CHC-2 ( $P_2$ ), Pusa Uday ( $P_5$ ) and DC-1 ( $P_6$ ), also had high gca effects for one or more characters contributing towards yield. This indicated that there was strong tendency of transmission of higher gain from the parents to the offspring. The present findings corroborated the earlier work of Munshi et al. (6, 7), Navazio and Simon (10), Gulamuddin and Ahmed (3), Sarkar and Sirohi (12), Shushir et al. (13), and Verma et al. (14) in cucumber; Munshi and Verma (9) in muskmelon and Munshi and Sirohi (8) in bittergourd.

The results indicated the importance of heterosis breeding for effective utilization of non-additive genetic variance which had a predominant role for the improvement of five most yield attributing traits, *viz.*, the days to first female flower opening, node number of first female flower, days to first fruit harvest, number of fruits per plant and total yield per plant. The other three characters, *viz.*, fruit weight, fruit length and fruit diameter where additive gene action is predominant can be improved through selection. The combination  $P_1 \times P_5$  (CRC-8 × Pusa Uday), which exhibited highest sca effects for days to first female flower opening and total yield per plant may be exploited for commercial cultivation.

Table 2. Estimates of general combining ability (gca) effects of parents for eight quantitative characters in cucumber	timates (	of gener	al combir	ning abilit	y (gca) €	effects of	parents	for eight	quantit	ative cha	aracters	in cucum	iber.			
Genotype	Days	to1 <sup>st</sup>	Node N	Days to 1 <sup>st</sup> Node No. of 1 <sup>st</sup>	Days to	Days to1 <sup>st</sup> fruit		Fruit weight	No. o	No. of fruits	Fruit length	ength	Fruit diameter	ameter	Total y	Total yield per
	female	flower	female flower female flower	e flower	harvest	vest	(g)	(f	per	per plant	(cm)	u)	(cm)	ц)	plant (g)	t (g)
	obe	opening														
	Mean	Effects	Mean Effects Mean Effects	Effects	Mean	Mean Effects	Mean	Mean Effects	Mean	Mean Effects Mean Effects	Mean I	Effects	Mean Effects	Effects	Mean	Mean Effects
CRC-8	48.33	-1.98**	4.60	48.33 -1.98** 4.60 -0.69**	60.20	60.20 -1.97**	120.83	120.83 -5.77**		8.20 0.14** 14.61 0.25	14.61	0.25	4.74	0.05	943.78	-98.25**
CHC-2	50.31	50.31 -0.52**	4.03	-0.42**	62.33	-0.48**	116.30	116.30 -7.54**	7.80	-0.33**	12.70	-0.33**	4.71	-0.06	860.53	-54.42**
G-338	50.20	0.71**	5.07	0.23**	63.57	0.72**	129.57	129.57 -9.42**	7.60	-0.80**	13.23	-0.84**	4.64	-0.12*	672.97	-171.66**
CH-20	52.44	52.44 1.16**	4.81	0.13**	63.51	1.25**	142.80	142.80 -2.89*	7.07	-0.43**	14.34	-0.74**	4.77	-0.10	1128.80	-74.60**
Pusa-Uday 51.60 0.27*	51.60	0.27*	4.94	0.06**	64.56	0.38**	152.13	152.13 11.81**	6.62	0.46*	14.60	0.46*	5.24	0.31**	1513.80	328.41**
DC-1	52.71	52.71 0.41**	5.11	0.18**	62.43	0.10	159.80	159.80 13.82**	5.64	0.31**	16.43	1.20**	4.45	0.17	1265.30	267.52**
SE(gi)		0.13		0.01		0.14		1.37		0.08		0.16		0.06		8.78
*,** Significant at 5 and 1% levels.	int at 5	and 1%	levels.													

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Character	Crosses with per-se performance	Crosses with significant sca effects
Days to first female flower opening	1×5 (47.10)	1×5 (-3.07)
	1×6 (47.76)	1×6 (-2.97)
	2×5 (50.01)	2×5 (-2.61)
	2×6 (50.09)	3×5 (-2.53)
	3×5 (51.40)	2×6 (-2.50)
Node number of first female flower	2×5 (3.50)	1×5 (-0.58)
	2×6 (3.60)	1×6 (-0.54)
	1×5 (3.80)	2×5 (-0.39)
	1×6 (3.86)	2×6 (-0.33)
	3×5 (4.42)	3×5 (-0.24)
Days to first fruit harvest	2×5 (56.47)	2×5 (-2.82)
	2×6 (56.53)	2×6 (-2.64)
	1×5 (57.93)	1×5 (-2.60)
	3×5 (58.03)	3×5 (-2.49)
	1× 6 (59.27)	1× 6(-2.25)
Fruit weight (g)	1× 6 (192.17)	1× 6 (25.80)
	1×5 (187.07)	2×5 (22.95)
	2×6 (187.01)	1×5 (22.70)
	2×5 (185.53)	2×6 (22.41)
	3×5 (177.87)	3×5 (16.56)
Number of fruits per plant	1×5 (10.27)	2×5 (1.68)
	2×5 (9.77)	1×5 (1.64)
	2×6 (9.67)	2×6 (1.42)
	1× 6 (9.30)	5× 6 (1.18)
	5× 6 (7.77)	1× 6 (1.12)
Fruit length (cm)	2×6 (21.13)	2×5 (2.91)
	2×5 (19.43)	1×5 (2.67)
	1×5 (19.27)	2×6 (2.37)
	1× 6 (18.50)	4×5 (1.18)
	4×5 (16.80)	1× 6 (1.15)
Fruit diameter (cm)	2×6 (6.18)	2×6 (0.68)
	1× 6 (5.77)	1× 6 (0.63)
	2×5 (5.40)	2×5 (0.51)
	1×5 (5.33)	1×5 (0.49)
	3×4 (4.77)	3×4 (0.40)
Total yield per plant (g)	1×5 (2192.28)	1×5 (420.38)
	2×5 (2067.85)	2×5 (363.06)
	1× 6 (1767.08)	1× 6 (380.94)
	2×6 (1692.13)	2×6 (345.62)
	2×3 (1156.68)	2×4 (331.66)

**Table 3.** Ranking of best crosses for *per-se* performance and desirable crosses for significant sca effects with respect to eight quantitative traits in cucumber.

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