

Original Research Article

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## Combining Ability Studies on Cucumber and Snapmelon Hybrids

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### ABSTRACT

A field study with 12 x 3 Line x Tester analysis in cucumber and snapmelon revealed highly significant estimates for general combining ability and specific combining ability for all the traits thereby indicating the importance of both additive and non-additive genetic variance in the inheritance of these traits. The genotyped L3 and L2 were found to be the most promising per se for most of the trait, whereas the cross combination L3 x T3, L2 x T2 involving good x good general combiner parents was found with good desirable sca effects. The cross combination L7 x T1 was found with best SCA effect for T.S.S. To further improve yield and quality traits inclusion of F1 combinations with high SCA and parents with good GCA in multiple crosses, Line x Tester mating could be a worthwhile approach.

#### Keywords

Cucumber,  
GCA, SCA,  
F1 hybrids,  
Line x testers.

#### Article Info

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### Introduction

Cucumber is warm season vegetable grown throughout the world under tropical and subtropical conditions. It is said to be the native of northern India (Pursglove, 1969). The fruits of cucumber is said to have cooling effect, prevent constipation, checks jaundice and indigestion (Nandkarni, 1927).

Besides this, the seed of cucumber is also used in Ayurvedic preparations and raw fruits are being used for cosmetic purpose. Snap melon (*Cucumis melo* var. *momordica*) belongs to the family cucurbitaceae is used as a vegetable in variety of ways. Snap melon is rich in quality and now snapmelon juice is gaining popularity as squash. Knowledge of the nature and magnitude of variation

promotes a rational choice of the characters in which selection can be exercised.

### Materials and Methods

The present investigation entitled "Heterosis, Combining Ability and Stability in Interspecific Hybrids of Cucumis", was conducted during Kharif, 2014 at three different locations (Horticulture Farm, Department of Horticulture, Rajasthan College of Agriculture, Udaipur, Agricultural Research Station, Banswara and KVK Chittorgarh).

Twelve inbred lines (female) of cucumis melo were crossed with three testers of cucumis

sativus in line x tester mating design to develop a total 36 hybrids at Hi-Tech Horticulture unit, Department of Horticulture, Rajasthan College of Agriculture, Udaipur. These 15 parents along with 36 hybrids and three standard checks (Mamta-5002, Sedona, Kakri surya prabha) were evaluated in randomized block design with three replications at three locations viz. Udaipur, Banswara and Chittorhgarh during kharif 2014. Lines and testers accessions were collected from NBPGR, New Delhi (Table 1).

The observations were recorded for eighteen important characters namely vine length, number of branches per vine, days to anthesis of first female and male flower, number of male flower per vine, number of female flower per vine, sex ratio, number of fruits per vine, fruit weight fruit length, fruit volume, fruit diameter, pulp thickness, total yield per vine, pulp weight, seed weight (Table 2).

## Results and Discussion

Analysis of variance for combining ability reflected significant difference among crosses for all the characters in all the environments, partitioning of this variance in lines, testers and line x testers revealed significant difference, among GCA of lines for all the characters, GCA of tester for all the characters except number of branches per vine in E1, specific gravity in E3.

Significant difference among SCA of hybrids was observed for all the characters in all the environments. In all the characters tester, lines and line x tester interacted with environments significantly except days to anthesis of first female flower, fruit weight and fruit length due to testers, this indicate lack of consistency in GCA of lines and tester and SCA of crosses across the environments. It suggested that selection of the parents and

crosses for GCA and SCA in different environments should be done separately.

Analysis of GCA, SCA variance for fruit texture was reported by Yoshioka *et al.*, (2010). Olfati *et al.*, (2011), Bairagi *et al.*, (2013) and reported that both GCA and SCA variance were important for yield, yield contributing characters, quality characters and for earliness.

The estimates of GCA effects revealed that the good general combiner for yield and yield contributing characters were lines L2, L3, L5, L8 and L12 for T.S.S. L7, L11, and L12 for fruit quality traits L2, L3, L5 and L12 for plant type trait L4, L9, L10 and L12 for flowering traits L1, L3, L6, L7, and L12 in general L2, L3, L5, L7 and L12 considered for good combiners for yield and majority of the traits (Table 3).

Among the tester the tester, T3 was considered good general combiner for yield, flowering and plant type traits.

The tester T2 was considered good general combiner for quality traits. The high general combining ability effects observed is due to additive gene effect and additive x additive gene effects (Griffing, 1956 and Sprague, 1966).

A perusal of SCA effects revealed that highest magnitude of positive significant SCA effects for total yield per vine were recorded in hybrids L2× T2 in E1 (3.10), in E3 (2.33) and on pooled basis (1.55), L10× T3 in E2 (2.06).

Two hybrids L10× T3 and L3× T3 exhibited positive significant SCA effects in all the three environments as well as on pooled basis for total yield per vine.

**Table.1** Description of parents

S.No	Symbol	Species Name	IC Number
<b>A. Lines (Female Parents)</b>			
1.	L <sub>1</sub>	<i>C.melo</i> var. <i>momoradica</i>	IC-415539
2.	L <sub>2</sub>	<i>C.melo</i> var. <i>momoradica</i>	IC-415521
3.	L <sub>3</sub>	<i>C.melo</i> var. <i>momoradica</i>	IC-433621
4.	L <sub>4</sub>	<i>C.melo</i> var. <i>utilissimus</i>	IC-315294
5.	L <sub>5</sub>	<i>C.melo</i> var. <i>utilissimus</i>	IC-258163
6.	L <sub>6</sub>	<i>C.melo</i> var. <i>utilissimus</i>	IC-313031
7.	L <sub>7</sub>	<i>C.melo</i> var. <i>momoradica</i>	VRSM-44
8.	L <sub>8</sub>	<i>C.melo</i> var. <i>agretris</i>	IC-258165
9.	L <sub>9</sub>	<i>C.melo</i> var. <i>momoradica</i>	VRSM-32
10.	L <sub>10</sub>	<i>C.melo</i> var. <i>momoradica</i>	DR/KPS/26
11.	L <sub>11</sub>	<i>C.melo</i>	BS-41
12.	L <sub>12</sub>	<i>C.melo</i> var. <i>momoradica</i>	VRSM-58
<b>B. Tester (Male Parent)</b>			
1.	T <sub>1</sub>	<i>C.sativus</i>	SKY/DR/RS
2.	T <sub>2</sub>	<i>C.sativus</i>	SPP-58
3.	T <sub>3</sub>	<i>C.sativus</i>	SPP-56

**Table.2** Grand Mean, Mean  $\pm$  SE (m) and range of eighteen characters in parents and F<sub>1</sub>

Characters	GM	Parents		F <sub>1</sub>	
		Mean $\pm$ SE(m)	Range	Mean $\pm$ SE(m)	Range
Vine length	2.39	2.68 $\pm$ 0.09	1.53 – 3.66	2.31 $\pm$ 0.09	1.30 – 3.78
Number of branches	4.31	4.01 $\pm$ 0.20	3.04 - 5.64	4.41 $\pm$ 0.20	2.69 – 5.89
Days to anthesis of first male flower	37.28	35.40 $\pm$ 0.67	31.80 – 39.09	38.11 $\pm$ 0.67	34.31 – 42.47
Days to anthesis of first female flower	43.49	41.63 $\pm$ 0.81	38.73 – 46.56	44.37 $\pm$ 0.81	39.42 – 52.18
No. of male flower/vine	132.90	137.56 $\pm$ 2.27	92.38 – 197.93	131.31 $\pm$ 2.27	94.29 – 157.89
No. of female flower/vine	16.52	16.32 $\pm$ 0.70	7.78 – 31.49	16.54 $\pm$ 0.70	8.36 – 29.27
Sex ratio	9.17	9.67 $\pm$ 0.37	4.86 – 14.44	9.06 $\pm$ 0.37	5.10 – 16.72
Number of fruit/vine	5.45	5.47 $\pm$ 0.24	3.18 – 11.98	5.52 $\pm$ 0.24	2.96 – 11.98
Fruit weight	611.30	558.99 $\pm$ 26.63	140.61 – 1441.06	653.74 $\pm$ 26.63	133.47 – 1385.67
Fruit diameter	7.53	7.87 $\pm$ 0.28	3.28 – 14.89	7.66 $\pm$ 0.28	4.40 – 11.82
Fruit length	18.15	16.34 $\pm$ 0.58	6.94 – 24.26	18.64 $\pm$ 0.58	7.91 – 34.30
Pulp thickness	14.79	14.63 $\pm$ 0.49	5.79 – 23.84	15.13 $\pm$ 0.49	8.00 – 24.28
T.S.S	5.14	4.81 $\pm$ 0.11	3.73 – 6.18	5.26 $\pm$ 0.11	3.76 – 8.17
Fruit volume	736.03	688.04 $\pm$ 20.40	204.22 – 1675.33	779.07 $\pm$ 20.40	146.22 – 1935.11
Specific gravity	0.86	0.84 $\pm$ 0.03	0.72 - 1.03	0.88 $\pm$ 0.03	0.71 – 1.03
Total yield/vine	3.26	2.93 $\pm$ 0.12	0.46 – 6.98	3.54 $\pm$ 0.12	1.04 – 7.73
Pulp weight	491.06	535.25 $\pm$ 16.62	158.61- 1656.37	495.07 $\pm$ 16.62	120.65 – 1474.43
Seed weight	120.16	126.18 $\pm$ 4.03	38.30 – 195.46	123.71 $\pm$ 4.03	34.96 – 226.91

**Table.3** GCA effects of parents

Parents	Vine length	No. of branches per vine	Days to anthesis of first male flower	Days to anthesis of first female Flower	No. of male flower per vine	No. of female flower per vine	Sex ratio	No. of fruit per vine
T1	0.11**	-0.03	-1.03**	-1.30**	-2.33**	0.42	-0.40**	0.82**
T2	-0.41**	0.00	0.68**	1.10**	-1.19	-1.60**	0.79**	-0.43**
T3	0.30**	0.03	0.35	0.20	3.53**	1.18**	-0.39**	-0.39**
L1	-0.19**	-0.15	-0.99*	-2.20**	-0.68	-1.31**	0.28	-0.21
L2	-0.08	-0.97**	-0.50	-0.46	4.37**	-1.20**	0.84**	0.10
L3	-0.15**	0.11	-1.59**	-1.68**	14.63**	2.79**	-0.80**	0.55**
L4	0.25**	-0.58**	0.80*	-0.03	-26.68**	-5.21**	0.92**	-1.37**
L5	-0.07	0.14	1.30**	0.23	8.55**	2.23**	-1.22**	-0.73**
L6	-0.16**	0.05	-1.84**	-0.71	-10.03**	-4.14**	1.40**	-1.25**
L7	-0.26**	0.33**	-1.85**	-0.75	-11.11**	3.63**	-2.84**	0.74**
L8	-0.87**	-0.61**	0.90*	-0.80	-14.11**	-3.79**	1.38**	-0.71**
L9	0.63**	0.42**	1.55**	1.73**	11.52**	1.09*	1.28**	-0.26
L10	0.16**	0.45**	2.44**	3.29**	14.47**	-1.33**	1.69**	-0.30*
L11	0.10	-0.45**	1.62**	1.90**	15.99**	3.08**	-0.28	3.35**
L12	0.65**	1.26**	-1.85**	-0.52	-6.91**	4.16**	-2.65**	0.10

Parents	Fruit weight	Fruit diameter	Fruit length	Pulp thickness	T.S.S	Fruit volume	SG	Total yield/vine	Pulp weight	Seed weight
T1	-22.79*	-0.11	-2.81**	0.05	-0.01	-60.29**	0.03**	0.17**	-46.64**	0.35
T2	-48.87**	-0.30**	0.23	-0.55**	-0.00	-42.44**	-0.02*	-0.28**	-29.96**	7.34**
T3	71.66**	0.41**	2.58**	0.50**	0.01	102.73**	-0.01	0.11**	76.60**	-7.69**
L1	-195.00**	-1.49**	-0.88*	-4.15**	-0.27**	-248.99**	0.02	-1.03**	-224.53**	-0.45
L2	226.97**	1.21**	2.86**	0.93**	-0.59**	250.49**	0.00	1.38**	269.40**	45.32**
L3	443.45**	2.51**	4.45**	5.19**	-0.58**	623.90**	-0.03	3.11**	727.26**	45.57**
L4	-273.94**	-2.61**	5.13**	-2.72**	-0.09	-272.85**	-0.09**	-2.06**	-152.83**	-42.98**
L5	214.37**	1.99**	-0.68	2.79**	-0.06	337.30**	-0.08**	0.68**	12.71	16.97**
L6	11.33	-1.87**	8.78**	-0.98**	-0.28**	104.49**	-0.08**	-0.67**	-46.67**	-20.46**
L7	-213.40**	0.25	-5.83**	-1.26**	0.90**	-271.29**	0.06**	-0.81**	-217.93**	-24.77**
L8	115.06**	-0.49**	0.79*	0.14	0.04	65.45**	0.06**	0.27**	-101.52**	8.66**
L9	25.13	1.10**	-2.51**	2.48**	0.07	-7.14	0.02	0.02	-40.33**	59.62**
L10	-61.97**	-0.15	-2.24**	0.88**	-0.35**	-97.22**	-0.01	-0.18*	-113.58**	-20.08**
L11	-502.37**	-2.72**	-9.99**	-6.69**	0.84**	-607.59**	0.02	-2.22**	-355.06**	-82.26**
L12	210.37**	2.27**	0.12	3.39**	0.38**	123.45**	0.11**	1.50**	243.08**	14.87**

**Table.4** SCA effects of hybrids

Parents	Vine length	No. of branches per vine	Days to anthesis of first male flower	Days to anthesis of first female flower	No. of male flower per vine	No. of female flower per vine	Sex ratio	No. of fruit per vine
L1 x T1	-0.58**	0.19	2.04*	0.62	5.65*	3.57**	-1.87**	-0.11
L2 x T1	0.05	-0.50*	0.62	0.66	6.79*	0.81	-0.83	-0.02
L3 x T1	-0.33**	-1.01**	1.20	-0.04	14.28**	-4.64**	2.98**	-1.07**
L4 x T1	0.03	0.45	-2.02*	-1.62	-0.17	-2.55**	1.99**	0.37
L5 x T1	-0.12	-0.17	-0.76	2.03*	1.59	-1.69*	0.83	-1.25**
L6 x T1	0.24*	-0.32	-0.93	-0.90	-24.66**	-4.21**	1.74**	-1.60**
L7 x T1	0.22*	0.13	0.03	1.42	-0.25	-3.46**	1.32**	-0.10
L8 x T1	-0.16	0.81**	-0.94	-0.92	-2.98	-3.58**	2.16**	0.50
L9 x T1	-0.47**	0.35	2.21**	1.96*	1.37	-0.45	-1.45**	0.30
L10 x T1	0.35**	0.22	0.10	0.93	-2.38	5.61**	-3.48**	0.48
L11 x T1	0.48**	-0.31	-1.32	-1.01	4.43	9.22**	-3.16**	2.29**
L12 x T1	0.28**	0.18	-0.23	-3.12**	-3.67	1.34	-0.23	0.21
L1 x T2	0.08	0.91**	-2.33**	-2.36*	-10.47**	0.57	-0.79	0.81**
L2 x T2	0.21	1.29**	-1.71*	-2.82**	0.85	4.70**	-3.16**	2.12**
L3 x T2	-0.03	0.72**	-0.53	-0.35	-13.64**	3.51**	-2.49**	1.21**
L4 x T2	0.53**	-0.52*	-0.35	0.23	-5.89*	4.18**	-3.33**	-0.76**
L5 x T2	-0.02	-1.14**	-1.73*	-4.28**	1.49	-0.40	-0.17	0.56
L6 x T2	0.30**	0.29	0.34	0.95	16.78**	2.00*	-0.35	1.12**
L7 x T2	-0.17	-0.02	1.24	-1.92	-6.88*	3.63**	-1.91**	0.66*
L8 x T2	0.28*	-1.07**	2.27**	0.07	13.73**	-0.33	0.90*	-1.09**
L9 x T2	0.66**	-0.60*	1.55	2.81**	-6.28*	-7.67**	5.59**	-0.65*
L10 x T2	-0.25*	-0.53*	-0.52	0.69	10.21**	-2.79**	3.08**	-0.87**
L11 x T2	-0.78**	0.61*	2.06*	4.81**	-1.44	-4.75**	2.18**	-3.15**
L12 x T2	-0.81**	0.06	-0.29	2.17*	1.55	-2.66**	0.44	0.04
L1 x T3	0.50**	-1.10**	0.29	1.74	4.81	-4.15**	2.65**	-0.70*
L2 x T3	-0.25*	-0.78**	1.09	2.16*	-7.63**	-5.51**	3.99**	-2.10**
L3 x T3	0.36**	0.29	-0.66	0.39	-0.63	1.13	-0.49	-0.14
L4 x T3	-0.57**	0.08	2.36**	1.39	6.06*	-1.63	1.33**	0.39
L5 x T3	0.15	1.30**	2.49**	2.24*	-3.08	2.08*	-0.66	0.69*
L6 x T3	-0.54**	0.04	0.59	-0.06	7.88**	2.21**	-1.39**	0.48
L7 x T3	-0.05	-0.11	-1.27	0.49	7.13**	-0.18	0.59	-0.56
L8 x T3	-0.12	0.26	-1.33	0.86	-10.75**	3.91**	-3.06**	0.58*
L9 x T3	-0.19	0.25	-3.76**	-4.76**	4.91	8.12**	-4.13**	0.35
L10 x T3	-0.10	0.31	0.42	-1.61	-7.82**	-2.83**	0.41	0.40
L11 x T3	0.30**	-0.30	-0.74	-3.80**	-2.99	-4.47**	0.97*	0.86**
L12 x T3	0.52**	-0.24	0.53	0.96	2.12	1.31	-0.21	-0.25

Parents	Fruit weight	Fruit diameter	Fruit length	Pulp thickness	T.S.S	Fruit volume	SG	Total yield/vine	Pulp weight	Seed weight
L1 x T1	-28.27	-0.61	1.03	0.08	0.03	-43.12	0.01	-0.16	21.27	-29.32**
L2 x T1	-92.10**	-0.68*	1.78*	-5.51**	0.27*	-116.15**	-0.01	0.03	-166.35**	57.53**
L3 x T1	-	-1.07**	1.10	3.92**	0.03	-380.23**	0.08*	-1.41**	298.73**	11.32*
L4 x T1	151.37**									
L5 x T1	-57.33	0.17	-0.05	-0.64	-0.25	-107.93**	0.05	-0.19	-43.96*	-10.37*
L6 x T1	130.69**	1.21**	1.87**	-1.78**	-0.37**	151.70**	0.01	-0.04	-176.85**	-24.59**
L7 x T1	146.30**	0.12	-6.69**	3.27**	0.25	104.07**	0.05	-0.28	200.82**	-6.92
L8 x T1	3.52	0.03	0.22	0.67	2.02**	9.40	0.03	-0.10	-4.95	-10.84*
L9 x T1	13.86	-0.30	1.13	-0.40	-0.03	44.88	-0.03	0.69**	-7.53	8.21
L10 x T1	-10.21	0.24	-1.33	1.51*	-0.33*	-10.08	0.01	0.26	-67.02**	-25.14**
L11 x T1	36.94	0.37	0.43	-1.76**	-0.36**	181.77**	-0.09*	0.59**	-82.44**	2.64
L12 x T1	4.89	-0.43	2.83**	-0.49	-0.29*	35.03	-0.01	0.12	27.27	-6.84
L1 x T2	3.06	0.96**	-2.32**	1.13	-0.97**	130.66**	-0.10**	0.51**	1.02	34.32**
L2 x T2	11.12	0.73*	-3.01**	0.56	0.82**	-39.19	0.05	0.29*	38.60	32.32**
L3 x T2	-7.65	-0.27	1.33	0.15	-0.90**	-16.90	0.03	1.55**	-62.28**	-32.64**
L4 x T2	-65.46*	-0.17	-1.01	-1.90**	0.12	-49.19*	-0.02	0.45**	-164.85**	-3.62
L5 x T2	131.12**	0.33	3.38**	0.75	-0.35**	256.88**	-0.09*	0.16	180.15**	38.41**
L6 x T2	-88.01**	0.09	-3.03**	1.86**	-0.23	-144.60**	0.02	-0.22	137.84**	17.08**
L7 x T2	18.49	0.24	6.64**	-1.33*	-0.13	152.44**	-0.06	0.63**	-61.55**	19.79**
L8 x T2	84.79**	0.68*	-2.70**	0.89	-0.92**	51.99*	0.01	0.61**	31.99	-1.15
L9 x T2	70.70*	-0.03	-1.08	1.63**	0.26*	11.03	0.04	-0.72**	16.53	-43.96**
L10 x T2	97.03**	0.34	2.99**	0.38	0.33*	95.62**	0.02	-0.18	138.51**	34.35**
L11 x T2	-	-1.16**	-4.27**	-3.17**	0.20	-322.30**	-0.04	-2.04**	-167.37**	-18.64**
L12 x T2	286.90**									
L1 x T3	51.30	0.28	0.50	0.34	0.26*	52.51*	-0.01	-0.23	21.67	-6.71
L2 x T3	-16.54	-1.04**	0.26	-0.16	0.53**	-48.30	0.05	-0.31*	-109.23**	-35.23**
L3 x T3	17.15	-0.11	1.97**	-0.64	-0.86**	82.31**	-0.07	-0.13	-59.87**	-3.00
L4 x T3	99.74**	0.96**	-3.11**	5.37**	0.63**	133.05**	-0.02	-1.58**	228.63**	-24.89**
L5 x T3	216.83**	1.24**	-0.09	-2.01**	-0.15	429.42**	-0.07	0.97**	-133.88**	-7.69
L6 x T3	-73.80*	-0.50	-3.33**	-0.11	0.60**	-148.95**	0.04	0.03	-136.18**	-28.05**
L7 x T3	-42.68	-1.29**	1.16	-0.08	0.60**	-7.10	-0.02	0.26	39.02	7.51
L8 x T3	-	-0.36	0.05	-1.95**	-0.12	-256.51**	0.02	-0.35*	-139.27**	-12.87**
L9 x T3	164.79**									
L10 x T3	-88.31**	-0.71*	2.48**	-1.56**	-1.10**	-61.40*	-0.04	-0.51**	-27.04	11.99*
L11 x T3	-84.56**	0.33	-0.05	-1.23*	-0.23	-55.91*	-0.02	0.03	-9.00	35.75**
L12 x T3	-86.82**	-0.57	-1.66*	-1.89**	-0.01	-85.54**	-0.03	-0.08	-71.49**	-9.21
L1 x T4	249.95**	0.78*	3.84**	4.92**	0.16	140.53**	0.13**	1.46**	249.81**	16.00**
L2 x T4	-56.19	0.14	-3.33**	0.14	0.03	-87.54**	0.02	0.10	-48.93*	13.55**
L3 x T4	13.48	0.09	2.06**	-0.97	0.44**	-82.36**	0.05	-0.20	108.21**	0.90

Hybrid L2× T2 besides total yield per vine also exhibited significant positive SCA effects for number of fruits per vine, number of female flower, and number of branches per vine. Hybrid L10× T3 in addition to yield per vine also exhibited significant positive SCA effects for fruit weight, fruit diameter, fruit length, pulp thickness, fruit volume, specific gravity, and pulp weight. Hybrid L3× T3 besides yield per vine also exhibited significant positive SCA effects for fruit volume, fruit weight, fruit diameter and for vine length. Hybrid L12× T1 in addition to yield per vine also exhibited significant positive SCA effects for vine length, fruit diameter and fruit volume (Table 4).

Out of total thirty six hybrids five best hybrids which exhibited highest positive significant SCA effect on pooled basis for total yield per vine were viz., L2× T2, L10× T3, L3× T3, L8× T1 and L6× T2. These hybrids also exhibited higher magnitude of economic heterosis with high mean performance. Similar findings for identification of superior parental lines, tester and hybrids based on GCA and SCA effects for fruit yield and morphological characters in cucumber were reported by Golabadi *et al.*, (2015), Tasdighi and Baker (1981), Musmade *et al.*, (1986) and Kumar *et al.*, (2013) for yield and its components and for fruit texture by Yoshioka *et al.*, (2010).

Out of total thirty six hybrids five best hybrids exhibited highest magnitude of positive significant SCA effects for T.S.S. on pooled basis were viz., L7× T1, L1× T2, L2× T3, L4× T3, and L5× T3 (Table 4). Similarly five best hybrids exhibiting highest positive significant SCA effects for fruit weight are viz., L10× T3, L3× T3, L6× T1, L4× T2, and L5× T1.

On the basis of SCA/GCA effect, per se performance, economic heterosis, heterosis,

heterobeltiosis best three economic hybrids were identified *i.e.* L3 x T3, L7 xT1, L4 x T2 for total yield per vine, T.S.S and fruit length respectively.

The mean performance of crosses could be envisaged as a criterion of SCA effect and selection of promising crosses based on per se performance. The present finding corroborated the earlier work of (Musmade *et al.*, 1986, Srivastava and Srivastava, 1976; and Tasdighi and Baker, 1981). From the results of this experiment, it may be suggested that it is possible to predict the best hybrid combination for yield from the GCA values of the parental lines involved; at least in this population. Therefore, it may be concluded that hybrid breeding programme aimed at yield improvement in cucumber should be based on high GCA for yield in parental arrays (Bairagi *et al.*, 2013).

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