

# Selecting parents for developing superior hybrids in cucumber (*Cucumis sativus* L.)

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**Abstract:** Estimates of general combining ability of parents and specific combining ability of the crosses help to select desired parents for hybridization and development of superior hybrids. Crosses among eight parents were attempted in a half-diallel fashion. The material comprising eight parents, 28 F<sub>1</sub>s and one check (Pusa Sanyog) was sown at two locations in Randomized Block Design with three replications. The highest estimates of general combining ability (gca) were exhibited by G<sub>2</sub> and Gyn<sub>1</sub> for most of the characters at both the locations. In general, there was close agreement between gca effects and per se performance, but in some cases it did not hold good, which may be due to a higher degree of gene action involved. The superior cross combinations which recorded high specific combining ability (sca) estimates and per se performance for yield and number of fruits were K-90 x G<sub>2</sub> and K-90 x Gyn<sub>1</sub> and hence may be exploited for the development of F<sub>1</sub> hybrid (s) after testing their performance at multi-locations for two to three years.

## 1. Introduction

Cucumber (*Cucumis sativus* L.), a member of the Cucurbitaceae family, is grown as a summer and rainy season crop in the low and mid hills of the northwestern Himalaya from April to August and fruits are available from June to October to the plains of northern India. The crop raised in the hills, being of high quality and off-season, brings good returns to the growers.

F<sub>1</sub> hybrids in cucumber, as in many vegetable crops, have several well known advantages over open-pollinated varieties (Dogra and Kanwar, 2011) and hence provide a scope for the breeder to find more appropriate combinations to develop superior hybrids. F<sub>1</sub> hybrids are early, vigorous, high yielding, tolerant to diseases and insect-pests and more efficient in the use of water and fertilizers. Currently, farmers are purchasing hybrid seeds from private firms who charge exorbitant prices for seed. To tide over the situation, there is a need to develop F<sub>1</sub> hybrids and make their seed available to farmers at a reasonable price. For the development of superior hybrids, estimates of general combining ability of parents and specific combining ability of the crosses help to properly select parents for hybridization. Moreover, use of gynocious lines for developing cucumber hybrids makes the production of F<sub>1</sub> seed more cost effective. Furthermore, there is urgent need

to develop stable hybrids adapted to a wide range of climatic conditions.

## 2. Materials and Methods

The present investigations were carried out at two locations: Experimental Farm Nauni (L1) and Experimental Farm Chambaghat (L2) of the Department of Vegetable Crops, Dr Y S Parmar University of Horticulture and Forestry, Solan (Himachal Pradesh), India, which are 1276 m a.m.s.l. and 1300 m a.m.s.l., respectively. Both locations fall in the mid-hill sub-temperate zone of the state of Himachal Pradesh; Nauni lies at latitude and longitude of 30° 52' N and 77° 11' and Chambaghat, 30° 55' N and 77° 06'. All the parents except two gynocious lines were of monoecious type. Crosses among eight parents were attempted in a half-diallel fashion. The material comprising eight parents, 28 F<sub>1</sub>s and one check (Pusa Sanyog) was sown in Randomized Block Design with three replications. Spacing was 1.25x1.00 m. Data were recorded on randomly selected plants for yield and horticultural characters at both the locations. Griffing's (1956) method II model I was used to derive general and specific combining ability estimates. The analysis of variance for combining ability was based on following mathematical model:

$$P_{ijk} = m + g_{ii} + g_{jj} + s_{ij} + b_k + e_{ijk}$$

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where,

- $P_{ijk}$  = phenotypes of the hybrids between  $i^{\text{th}}$  and  $j^{\text{th}}$  parents in  $k^{\text{th}}$  plots
- $m$  = population mean
- $g_{ii}$  = GCA effects of  $i^{\text{th}}$  parent
- $g_{jj}$  = GCA effects of  $j^{\text{th}}$  parent
- $s_{ij}$  = SCA of the crosses between  $i^{\text{th}}$  and  $j^{\text{th}}$  parents
- $b_k$  = block effects
- $e_{ijk}$  = environmental effect associated with  $ijk^{\text{th}}$  observation

### 3. Results and Discussion

Analysis of variance (Table 1) for combining ability revealed that the importance of  $gca$  ( $\sigma^2g$ ) was more than  $sca$  ( $\sigma^2s$ ), indicating the preponderance of additive gene action for days to first female flower appearance (DFFFA) at location 1 and days to marketable maturity (DMM) at both locations. However, in all the other traits, the  $sca$  component was higher in magnitude than  $gca$ 's, indicating the preponderance of non-additive gene effects. However, mean sum of squares for  $gca$  and  $sca$  were highly significant for all the characters except TSS, suggesting the importance of both additive and non-additive genetic variance in agreement with the findings of Om *et al.* (1978). Similar trends at both the locations proved that the conclusions on gene actions are authentic.

The parents  $G_2$ ,  $Gyn_1$  and Poinsette had negative estimates for DFFFA and node at which first female flower appears (NFFF) at both the locations (Table 2) showing earliness in fruit bearing and were good general combiners for these characters. Among  $F_1$ 's, the  $sca$  effects were significantly negative in 12 and 15 crosses, respectively, for these two traits at L1 (Table 3) whereas significantly negative in 15 crosses for each of these two traits at L2

(Table 4). The crosses LC-11 x  $Gyn_1$  (poor x high) and EC 173934 x LC-40 (poor x poor), respectively, had the highest  $sca$  effect at L1 and the crosses LC-11 x LC-40 (poor x poor) and EC 173934 x LC-40 (poor x poor), respectively, had the highest  $sca$  effects at L2 for these traits. The parents  $G_2$  and  $Gyn_1$  (L1) and  $G_2$ ,  $Gyn_1$  and Poinsette (L2) with significantly high  $gca$  estimates (with negative value) were good general combiners for DMM. Crosses LC-11 x  $Gyn_1$ , EC 173934 x LC-40, K-90 x  $G_2$  and K-90 x EC 173934 had high  $sca$  estimates at both the locations for DMM. El-Shawaf and Baker (1978), Om *et al.* (1978), and Wang and Wang (1980) also reported greater additive genetic variance for DMM. The parents  $G_2$  and  $Gyn_1$  may be used in the hybridisation programme for developing early hybrids adapted to a wide range of climate. LC-11 x  $Gyn_1$  and EC 173934 x LC-40 may be exploited as early hybrids after further multi-locational testing. These crosses may also be exploited to produce transgressive segregants in advanced generations.

With regard to fruit length, the parents  $Gyn_1$ , LC-11 and K-90 were good general combiners as is evident from their high  $gca$  estimates at both locations. Fourteen crosses exhibited significant  $sca$  effects. The  $sca$  effects were high in crosses Poinsette x LC-40 and  $G_2$  x Poinsette involving poor x poor general combiners. K-90, K-75 and EC 173934 had the highest  $gca$  with respect to fruit width and hence were good general combiners. The  $sca$  effect was maximum in  $G_2$  x  $Gyn_1$  involving poor x poor general combining parental lines (at L1) and in  $G_2$  x K-75 involving poor x high general combining parental lines (at L2). In India, slicing cucumbers are preferred, therefore lengthy fruits are desirable. Kupper and Staub (1988) and Hormuzdi and More (1989) reported contrasting results for fruit length and width due to different experimental material and environment.

Table 1 - Analysis of variance for combining ability for different characters in  $F_1$  cucumber

Source of variation	Df	Character										
		Days to first female flower appearance	Node of first female flower	Days to marketable maturity	Fruit length	Fruit width	TSS	Flesh to seed cavity ratio	Fruit weight	No. of fruits per plant	Yield per plant	Internodal length
<b>Location 1 - Nauni</b>												
Gca	7	678.818 *	27.997 *	705.436 *	6.425 *	1.087 *	0.005	0.001 *	3787.657 *	9.898 *	0.735 *	9.512 *
Sca	28	42.264 *	3.049 *	45.029 *	3.237 *	0.243 *	0.021	0.0015 *	693.149 *	1.159 *	0.193 *	2.183 *
Error	70	0.557	0.228	0.562	0.004	0.002	0.0013	0.00004	62.357	0.112	0.0013	0.272
$\sigma^2g$		67.826	2.777	70.487	0.642	0.108	0.0004	0.0001	372.53	0.979	0.073	0.924
$\sigma^2s$		41.707	2.821	44.467	3.0233	0.240	0.020	0.002	630.79	1.047	0.191	1.911
$\sigma^2g/\sigma^2s$		1.626	0.984	1.585	0.199	0.451	0.021	0.068	0.591	0.934	0.383	0.483
<b>Location 2 - Chambaghat</b>												
Gca	7	390.457 *	35.726 *	577.811 *	7.820 *	0.993 *	0.012 *	0.0016 *	3515.486 *	14.247 *	0.786 *	7.800 *
Sca	28	67.477 *	4.551 *	37.300 *	3.895 *	0.268 *	0.028 *	0.0009 *	612.551 *	1.582 *	0.181 *	1.510 *
Error	70	0.431	0.205	0.442	0.089	0.023	0.006	0.000035	49.232	0.148	0.0096	0.358
$\sigma^2g$		39.003	3.552	57.737	0.773	0.097	0.0006	0.000159	346.630	1.409	0.078	0.744
$\sigma^2s$		67.046	4.346	36.859	3.806	0.245	0.022	0.00088	563.320	1.434	0.171	1.153
$\sigma^2g/\sigma^2s$		0.582	0.817	1.566	0.203	0.395	0.029	0.081	0.615	0.983	0.452	0.646

\* Significant at 5% level of significance.

The best general combiners for TSS at both locations in order of merit were EC 173934 and LC-40. Among 28 specific combinations, 16 (at L1) and 14 (at L2) crosses exhibited positive sca effects being maximum in K-90 x Poinsette and Poinsette x K-75 at L1 and LC-40 x Gyn<sub>1</sub>, K-90 x Poinsette and K-75 x LC-40 at L2. For flesh to seed cavity ratio (FSR), the best general combiners were Poinsette, EC 173934 and Gyn<sub>1</sub>, irrespective of locations. Cross combination K-90 x K-75 at L1 and Poinsette x EC 173934 at L2 had maximum sca among seven significant and positive specific combinations. In contradiction to the present results, importance of additive gene action for FSR has been reported (Dogra, 1995).

The parents LC-11, K-90 and K-75 depicted high *per se* performance with respect to fruit weight at both locations as is evident from their high gca effect (Table 2). These parents had maximum concentration of favourable genes for increasing fruit weight. Eleven (at L1) and 12 (at L2) specific cross combinations had significantly positive sca effects (Tables 3 and 4), being maximum in K-90 x LC-11 (high x high) and K-90 x EC 173934 (high x poor). Non-additive gene action for fruit weight was also obtained by Ghaderi and Lower (1979) in consonance with the present findings. However, Gyn<sub>1</sub> and G<sub>2</sub> were identified as good general combiners for number of fruits per plant. The top specific combinations in order of merit were

K-90 x G<sub>2</sub>, K-90 x Gyn<sub>1</sub> and K-75 x Gyn<sub>1</sub> involving medium high, medium x high and poor x high general combiners, respectively. The situation holds good for both the locations with respect to number of fruits. Importance of non additive gene action for number of fruits per plant was also reported (Om *et al.*, 1978; Ghaderi and Lower, 1979; Dogra, 1995). However, the present results with regard to fruit weight and number of fruits are in disagreement with El Hafeez *et al.* (1997). This may be due to differences in the parental material used for making diallel crosses.

For yield per plant, K-90 was the best general combiner in addition to Gyn<sub>1</sub> and G<sub>2</sub> irrespective of location (Table 2). The sca effects (Tables 3 and 4) were high for K-90 x G<sub>2</sub> (high x high), K-90 x Gyn<sub>1</sub> (high x high) and LC-11 x Gyn<sub>1</sub> (poor x high). The present results on yield per plant were similar to earlier findings of Om *et al.* (1978), Ghaderi and Lower (1979), Wang and Wang (1980) and Doligibh and Sidorova (1983) but in contradiction to the work of Gu *et al.* (2004). Parents such as G<sub>2</sub>, Gyn<sub>1</sub> and LC-40 had negative gca effects and were considered good general combiners for internodal length. Nine (at L1) and 10 (at L2) specific combinations had significant negative values with the maximum in K-90 x Poinsette and Poinsette x EC 173934, poor x poor general combiners at each location.

As is evident from the data in Tables 2, 3 and 4, environmental effect was observed as non-significant on geno-

Table 2 - Estimates of general combining ability of parents for different characters in cucumber

Source of variation	Character										
	Days to first female flower appearance	Node of first female flower	Days to market-able maturity	Fruit length	Fruit width	TSS	Flesh to seed cavity ratio	Fruit weight	No. of fruits per plant	Yield per plant	Internodal length
<b>Location 1</b>											
K-90	0.000	0.367*	-0.550*	0.361*	0.364*	-0.016*	-0.0002	20.083*	0.017	0.276*	0.021
G2	-12.133*	-2.567*	-12.217*	-1.404*	-0.041*	0.004	0.004*	-25.250*	1.317*	0.302*	-1.856*
Poinsette	-2.433*	-0.767*	-2.0183*	-0.105	-0.531*	-0.031	0.014*	-4.917*	-0.217*	-0.055*	1.048*
EC173934	8.167*	1.633*	8.517*	-0.390*	0.191*	0.037*	0.011*	7.417*	-0.617*	-0.346*	0.144
K-75	0.733*	0.733*	1.017*	-0.050*	0.320*	-0.004	0.017*	10.083*	-0.017	0.024*	1.084*
LC-11	6.600*	0.633*	6.583*	0.388*	0.136*	-0.022	0.007*	32.750*	-0.783*	-0.089*	0.604*
LC-40	9.800*	2.067*	9.950*	-0.225*	-0.008*	0.029*	0.005*	-8.417*	-1.283*	-0.379*	-0.593*
Gyn1	-10.733*	-2.100*	-11.117*	1.425*	-0.433*	0.002	0.008*	-16.917*	1.583*	0.268*	-0.453
SE (gi)	0.221	0.141	0.222	0.019	0.013	0.011	0.0019	2.336	0.099	0.011	0.154
CD0.05 (gi)	0.441	0.281	0.443	0.037	0.026	0.021	0.0038	4.658	0.197	0.022	0.307
<b>Location 2</b>											
K-90	0.075	0.258*	-0.267*	0.208*	0.269*	-0.021	-0.013*	20.492*	0.508*	0.301*	0.116
G2	-10.092*	-2.908*	-11.600*	-1.355*	0.016	-0.015	-0.016*	-23.341*	1.842*	0.285*	-1.828*
Poinsette	-1.158*	-0.375*	-1.133*	-0.285*	-0.574*	-0.008	0.018*	-7.141*	-0.325*	-0.053*	0.693*
EC173934	7.642*	1.192*	7.867*	-0.592*	0.196*	0.065*	0.014*	-5.342*	-0.858*	-0.384*	0.489*
K-75	2.908*	0.792*	1.100*	-0.025	0.309*	-0.013	-0.010*	8.825*	-0.258*	0.058*	0.869*
LC-11	2.875*	1.325*	6.300*	0.495*	0.083*	-0.028*	-0.0001	30.825*	-1.092*	-0.125*	0.513
LC-40	5.675*	2.358*	7.900*	-0.148*	0.083*	0.045*	-0.0007	-5.342*	-1.358*	-0.363*	-0.364*
Gyn1	-7.925*	-2.642*	-10.167*	1.702*	-0.381*	-0.026*	0.009*	-18.375*	1.542*	0.279*	-0.488*
SE (gi)	0.194	0.134	0.197	0.088	0.044	0.023	0.0018	2.076	0.114	0.029	0.177
CD0.05 (gi)	0.387	0.267	0.393	0.175	0.088	0.046	0.0036	4.139	0.227	0.058	0.353

\* Significant at 5% level of significance.

Table 3 - Estimates of specific combining ability of  $F_1$  for different characters in cucumber at Nauni (L1)

Crosses	Characters										
	Days to first female flower appearance	Node of first female flower	Days to marketable maturity	Fruit length	Fruit width	TSS	Flesh to seed cavity ratio	Fruit weight	No. of fruits per plant	Yield per plant	Inter-nodal length
K-90x G2	-7.422*	-1.059*	-7.252*	-2.481*	-0.411*	-0.082*	0.122*	-13.259*	2.685*	1.023*	-0.627*
K-90x Poinsette	-4.789*	-0.526*	4.618*	-0.440*	0.086*	0.275*	-0.046*	-38.593*	0.848*	-0.324*	-2.530*
K-90x EC173934	-7.056*	-1.259*	-7.612*	-0.445*	-0.573*	-0.073*	-0.040*	40.574*	-0.752*	-0.059*	-1.894*
K-90x K-75	10.378*	1.974*	9.484*	1.382*	0.451*	0.011	0.012*	-35.259*	-0.352*	-0.543*	0.466
K-90x LC-11	9.178*	-0.259	9.948*	-0.406*	0.516*	-0.061*	-0.031*	55.407*	-0.252	-0.309*	2.246*
K-90x LC-40	-5.356*	-1.693*	-5.085*	1.324*	-0.794*	0.006	0.006*	-1.759	-0.085	-0.303*	2.043
K-90x Gyn1	-4.489*	-0.526*	-5.352*	-1.343*	-0.182*	-0.051*	-0.013*	-23.259*	2.382*	0.509*	-0.030
G2 x Poinsette	2.011*	1.074*	3.715*	2.334*	-0.393*	0.002*	-0.045*	0.074	-0.119	-0.316*	0.680*
G2 x EC173934	3.411*	0.007	3.348*	1.819*	0.482*	-0.032	-0.029	-4.093	-1.052*	-0.398*	0.883*
G2x K-75	0.178	-1.093*	0.515	2.113*	0.306*	0.112*	-0.009*	21.074*	1.348*	0.494*	1.143*
G2 x LC-11	-3.356*	0.674*	-4.052*	-1.142*	-0.367*	0.043	-0.014*	24.074*	0.115	0.268*	-0.044
G2 x LC-40	9.444*	0.574*	9.248*	0.638*	-0.240*	0.160*	0.003	-26.426*	-1.385*	-0.659*	0.453
G2 x Gyn1	-0.356*	0.741*	0.982*	1.421*	0.756*	0.093*	-0.023*	-14.593*	0.082	-0.140*	1.013*
Poinsette x EC173934	1.044*	-1.126*	1.315*	2.070*	0.192*	0.085*	0.080*	5.574	0.482*	0.065*	-2.387*
Poinsettex K-75	-0.522*	0.107	-1.185*	2.002*	0.343*	0.223*	-0.039*	16.407*	-0.118	0.201*	2.006*
Poinsettex LC-11	-5.056*	-1.126*	-5.418*	-0.325*	-0.527*	-0.023*	-0.035*	10.407*	0.648*	0.478*	1.419*
Poinsettex LC-40	-4.922*	-1.226*	-5.452*	2.622*	0.654*	-0.256*	0.032*	-28.426*	0.481*	0.088*	-1.517*
Poinsettex Gyn1	14.944*	0.941*	15.615*	-1.995*	0.126*	0.120*	0.032*	18.407*	-0.385*	0.260*	1.109*
EC173934x K-75	-1.456*	1.041*	-2.885*	0.033	-0.276*	-0.205*	0.0006	-7.759*	-0.718*	-0.314*	1.009*
EC173934x LC-11	-0.322*	-2.526*	-0.785*	1.161	0.139*	-0.074*	0.044*	9.574*	0.048	0.189*	0.489*
EC173934x LC-40	-9.189*	-3.293*	-8.818*	1.174*	-0.525*	-0.107*	-0.002	-5.926	1.548*	0.446*	0.353
EC173934x Gyn1	9.678*	3.541*	8.915*	-0.092*	-0.059*	0.036*	-0.018*	-14.093*	-1.985*	-0.602*	0.179
K-75x LC-11	3.778*	0.374	3.715*	1.238*	0.436*	0.020	-0.014	-36.259*	0.448*	-0.258*	-2.084*
K-75x LC-40	4.244*	-2.059*	4.682*	0.901*	-0.677*	0.134*	-0.013*	-0.093	-0.052	-0.028*	-1.954*
K-75x Gyn1	-6.556*	-0.893*	-6.585*	-1.116	-0.379*	-0.213*	-0.020*	13.407*	1.415*	0.538*	0.006
LC-11x LC-40	4.044*	4.041*	4.115*	0.496*	-0.593*	-0.005	-0.003	22.241*	-0.285	0.049*	1.259*
LC-11x Gyn1	-9.422*	-1.459*	-9.818*	-0.454*	-0.158*	0.128*	-0.030*	25.741*	0.181	0.518*	-0.614*
LC-40x Gyn1	3.378*	-0.226	2.482*	-1.707*	0.173*	0.211	-0.019*	-16.426*	-1.318*	-0.352*	-2.084*
SE (ij)±	0.676	0.433	0.680	0.058	0.044	0.032	0.0057	7.160	0.303	0.033	0.472
CD0.05	1.994	0.883	1.356	0.116	0.088	0.064	0.011	14.280	0.604	0.066	0.941

\*Significant at 5% level of significance.

Table 4 - Estimates of specific combining ability of F<sub>1</sub> for different characters in cucumber at Chambaghat (L2)

Crosses	Characters										
	Days to first female flower appearance	Node of first female flower	Days to marketable maturity	Fruit length	Fruit width	TSS	Flesh to seed cavity ratio	Fruit weight	No. of fruits per plant	Yield per plant	Inter-nodal length
K-90x G2	-7.826*	-0.915*	-7.207*	-2.255*	-0.745*	-0.095*	0.009*	-8.641*	3.696*	1.058*	-1.097*
K-90x Poinsette	-3.759*	0.885*	-3.674*	-0.025	0.012	0.265*	-0.390*	-28.174*	-0.137	-0.361*	-0.917*
K-90x EC173934	-6.893*	-2.015*	-6.674*	-0.152*	-0.625*	-0.142*	-0.028*	43.359*	-1.270*	-0.140*	-1.580*
K-90x K-75	9.174*	1.719*	11.426*	1.081	0.295*	0.070	0.042*	-27.474*	-0.537*	-0.565*	1.140*
K-90x LC-11	11.207*	-0.148	7.893*	-0.172	0.088	-0.049	-0.014*	33.859*	-0.370*	-0.252*	2.263*
K-90x LC-40	-1.593*	-1.515*	-3.374*	1.705*	-0.412*	-0.022	-0.004	-1.641	-0.437*	-0.288*	-0.193
K-90x Gyn1	-7.659*	-1.182*	-5.974*	-1.578*	-0.082*	-0.050	-0.007*	-19.674*	1.663*	0.431*	-0.737*
G2 x Poinsette	3.741*	1.385*	4.659*	2.838*	-0.502*	-0.009	-0.036*	-17.674*	-0.470*	-0.255*	0.027
G2 x EC173934	0.607*	0.819*	2.326*	2.845*	0.428*	0.018	-0.022*	-14.474*	-1.937*	-0.434*	1.363*
G2x K-75	-5.659*	-1.448*	-2.907*	2.278*	0.782*	0.196*	-0.001	23.026*	0.796*	0.334*	-0.583*
G2 x LC-11	-1.293*	0.352	-3.774*	-1.275*	-0.392*	0.045	-0.008*	24.359*	-0.704*	0.184*	-0.327
G2 x LC-40	18.574*	1.319*	7.293*	0.702*	0.008	-0.095*	0.023*	-12.807*	-1.437*	-0.515*	0.717*
G2 x Gyn1	-0.826*	0.652*	2.026*	1.018*	0.738*	0.143*	-0.014*	-7.507*	-0.670*	-0.087	0.807*
Poinsette x EC173934	2.674*	-1.048*	2.859*	0.908*	0.318*	0.145*	0.077*	14.326*	0.896*	0.147*	-1.957*
Poinsette K-75	-3.593*	-0.315	-1.374*	-1.792*	0.072	0.190*	-0.042*	30.159*	0.296	0.255*	1.330*
Poinsette LC-11	-2.893*	0.485*	-6.574*	-0.478*	-0.302*	0.005	-0.012*	11.493*	0.796*	0.492*	1.587*
Poinsette LC-40	-0.693*	-1.548*	-2.841*	2.665*	1.065*	-0.170*	0.029*	-29.007*	0.729*	0.066	-1.903*
Poinsette Gyn1	11.574*	1.118*	13.893*	-2.252*	-0.372*	0.069	0.015*	9.959*	-0.170	0.344*	0.753*
EC173934x K-75	-1.726*	3.452*	-0.041	-0.085	-0.165*	-0.250*	0.002	-3.307	-0.170	-0.247*	-0.200
EC173934x LC-11	4.307*	-2.082	0.759*	1.095*	0.195*	-0.069	0.042*	19.693*	0.330*	0.236*	-0.343
EC173934x LC-40	-6.159*	-4.115*	-8.507*	1.105*	-0.205*	-0.175*	-0.004	-19.141*	1.263*	0.340*	1.600*
EC173934x Gyn1	2.774*	0.219	4.893*	-0.412*	-0.109*	-0.070*	-0.014*	-18.841*	-1.970*	-0.618*	1.423*
K-75x LC-11	5.707*	-1.015*	4.193*	1.062*	0.582*	-0.157*	-0.014*	-41.141*	0.396*	-0.309*	-0.857*
K-75x LC-40	4.907*	-3.715*	4.259*	0.938*	-0.885*	0.203*	-0.017*	-6.041	-0.004	-0.131*	-1.613*
K-75x Gyn1	8.507*	-1.048*	-5.674*	-1.578*	-0.255*	-0.242*	-0.020*	10.326*	1.429*	0.307*	0.077
LC-11x LC-40	-22.726*	3.085*	6.393*	0.618*	-0.825*	-0.015	-0.014*	23.026*	0.163	0.025*	0.077
LC-11x Gyn1	-5.793*	-2.248*	-7.541*	-0.865*	0.238*	0.123*	-0.030*	28.326*	-0.404*	0.574*	-1.467*
LC-40x Gyn1	2.074*	5.052*	1.859*	-2.322*	0.105	0.317*	-0.023*	-12.174*	-1.137*	-0.302*	-1.322*
SE (ij)±	0.595	0.411	0.603	0.271	0.134	0.070	0.0054	6.362	0.349	0.089	0.542
CD0.05	1.186	0.819	1.202	0.540	0.267	0.139	0.011	12.685	0.696	0.177	1.081

\*Significant at 5% level of significance.

types and hybrid combinations for most of the characters. The results are similar at both locations with developed hybrid combinations and hence hybrids K-90 x G<sub>2</sub> and K-90 x Gyn<sub>1</sub> can be exploited in similar types of climates.

K-90, G<sub>2</sub> and Gyn<sub>1</sub> may be used in hybridisation for developing high yielding hybrids with higher number of fruits per vine, long fruits and high TSS on the basis of results from location 1, whereas G<sub>2</sub> and Gyn<sub>1</sub> are promising for developing high yielding hybrids with higher number of fruits per vine and short inter-nodal length on the basis of results from location 2. It can be concluded that G<sub>2</sub> and Gyn<sub>1</sub> may be used in hybridisation for developing high yielding hybrids with more fruits per vine and wider adaptability. The crosses K-90 x G<sub>2</sub> and K-90 x Gyn<sub>1</sub> can be released as hybrids after further testing.

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