

Original Research Article

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Effect on Gene Action, Combining Ability and Heterosis for Yield and Yield Attributing Traits in Green Gram (*Vigna radiata* L. Wilczek)

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ABSTRACT

Keywords

Variability, Additive, Non-additive, GCA, SCA, Heterosis

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Ten genotypes as lines and three genotypes as testers of mungbean were investigated for combining ability analysis through crosses developed into line x tester passion. The study revealed that the gene action involved in controlling traits viz., plant height, number of pods per plant, number of seeds per pod and grain yield per plant was non-additive; while the characters viz., number of branches per plant and 100-grain weight had additive gene action. Genotype KM 2264 was found good general combines for most of the yield attributing traits. Among testers KM 2195 was found good general combiner for all the characters. Crosses KM 2312 × KM 2241, KM 2260 × KM 2241 and KM 2318 × KM 2241 exhibited highly significant positive SCA effects along with higher *per se* performance for yield contributing traits.

Introduction

Mungbean [*Vigna radiata* (L.) Wilczek] is one of the most important legume crops in South and Southeast Asia. It is a warm season annual, highly branches having trifoliate leaves. It is grown under rainfed condition of arid and semi-arid regions of India during *kharif* and under irrigated condition during summer season. Choice of parents is crucial for breeding high yielding varieties hence the knowledge of combining ability and heterosis

assists in selecting the suitable parents for breeding programme and the gene action determines the breeding methodology. Hence, the present investigation was undertaken to assess the gene action involved and the combining ability effects along with heterosis in the mungbean during *kharif* season.

Materials and Methods

Ten diverse mungbean genotypes viz., KM 2260, KM 2262, KM 2263, KM 2264, KM

2310, KM 2312, KM 2318, KM 2319, KM 2356 and KM 2358 were selected as lines and three genotypes KM 2195, KM 2241 and IPM99-125 as testers. These genetically diverse selected genotypes were crossed in line × tester fashion and thirty hybrids were developed during summer 2017. These hybrids along with their parents were grown in a randomized block design with three replication during *kharif* 2017. Each hybrid and parent was sown in 3m length with 45 x 10cm spacing. Recommended cultural practices were followed to grow a good crop. Observations were recorded for days to 50 per cent flowering, plant height (cm), number of branches per plant, number of pods per plant, number of seeds per pod, 100-seed weight (g) and grain yield per plant (g). The data were recorded on five randomly taken plants per entry per replication. The line x tester analysis was done as per method prescribed by Kempthorne (1957).

Results and Discussion

Analysis of variance revealed significant differences (Table 1) for all the genotypes for most of character except days to 50 per cent flowering indicating the presence of sufficient amount of variability in expression of traits. The variance of lines and testers along with their interaction was significant for plant height, number of branches per plant, number of pods per plant, number of seeds per pod, 100-seed weight and grain yield per plant and non-significant for days to 50 per cent flowering indicating the absence of hybrid vigour for this character. Similar results were observed for variances of hybrids and parents vs hybrid. These results are in agreement with earlier findings reported by Saxena and Sharma (1989).

Higher magnitude of variance for hybrids as compared to their parents was observed for plant height, number of branches per plant,

100-seed weight and grain yield per plant indicating the presence of heterosis for these characters. Ratio of SCA to GCA variances was greater than one for characters like plant height, number of pods per plant, number of seeds per pod and grain yield per plant indicating the preponderance of non-additive gene action.

While the traits number of branches per plant and 100-grain weight exhibited SCA to GCA variance ratio less than one indicating the presence of additive gene action. These finding are in agreement with the finding of Manivanna (2002), Singh and Dikshit (2003), Anbumalarmathi *et al.*, (2004), Pandiyan *et al.*, (2006) and Barad *et al.*, (2008).

Genetic mechanism controlling quantitative traits and selection of suitable parents for further hybridization programme were provided through combining ability studies. Making crosses between good combiners and selection of parents on the basis of combining ability are expected to throw out maximum desirable segregants in the upcoming generations.

Days for 50 per cent flowering where negative GCA effect is desirable (Table 2 and Figure 1), the lines-KM 2264 and KM 2319; among tester KM 2195 had significantly higher negative GCA effect in the desired direction. For plant height, the lines KM 2318, KM 2312 and KM 2319; among testers none of them had positive significant GCA effect in the desired direction. Rupal Dhoot *et al.*, (2017) reported significant positive correlation between plant height and seed yield per plant. For number of branches per plant, the line KM 2264 and none of the tester had positive GCA effect. For number of pods per plant the line KM 2264 and tester KM 2195 had significant positive effect. Number of seeds per pod, the lines KM 2264, KM 2262 and KM 2260 and none of testers had positive significant GCA

effect. 100-seed weight KM 2263, KM 2264, KM 2262, KM 2356, KM 2310 and KM 2260 and among tester KM 2195 had positive significant GCA effect. Lines KM 2264, KM 2356, KM 2260 and KM 2262 and among testers KM 2195 and IPM-99-125 had higher and significant positive GCA effect.

Among the lines KM 2264, KM 2262 and KM 2260 and among the testers KM 2195 were found to have significant GCA effect for most of the character in desired characters. Hence, these genotypes are good combiner and hence can be exploited in the hybridization programme.

Table.1 Analysis of variance, estimates of combining ability variance and degree of dominance for line x tester mating design of six metric traits in mungbean

Source of Variation	d.f.	Plant height (cm)	Number of branches per plant	Number of pods per plant	Number of seeds per pod	100-gra in weight (g)	Grain yield per plant (g)
Replications	2	1.93	0.69**	8.91	4.28	0.56	2.41
Genotypes	42	19.14**	0.23**	69.32**	4.32**	0.30**	12.87**
Parents	12	0.11**	0.17**	41.26**	4.32**	0.14**	4.83**
Female	9	8.12**	0.21**	50.88**	4.42**	0.17**	4.85**
Males	2	21.94**	0.42**	29.44**	3.33**	0.04*	2.31**
Female vs males	1	1.09	0.47**	0.87	4.43**	0.008*	13.59**
Hybrids	29	14.87**	0.38**	32.61**	3.11**	0.17**	9.24**
Parent vs hybrid	1	231.20**	0.41**	587.76**	6.85**	6.04**	231.4**
Error	84	1.61	0.12	1.50	1.11	0.5	0.78
σ_g^2		3.27	0.07	8.96	0.73	0.06	4.63
σ_s^2		3.99	0.5	10.32	1.09	0.05	5.11
$(\sigma_s^2/\sigma_g^2)^{0.5}$		1.08	0.81	1.06	1.18	0.89	1.17

*, ** Significant at P=0.05 and P=0.01, respectively.

Table.2 Estimates of gca effects of parents for six metric traits of a line x tester mating design in mungbean

Source of Variation	Days to 50 per cent flowering	Plant Height (cm)	Number of branches per plant	Number of pods per plant	Number of seeds per pod	100-seed weight (g)	Grain yield per plant (g)
Line							
KM 2260	1.59**	-2.46**	0.05	-1.79**	0.42*	0.06**	0.83*
KM 2262	-0.38	0.15	-0.05	1.52**	0.97**	0.15**	0.23**
KM 2263	0.32	0.03	0.03	-0.32	-0.82**	0.21**	-0.89**
KM 2264	-1.42**	-1.73**	0.31**	1.41**	0.72**	0.18**	0.93**
KM 2310	-0.73*	0.59*	0.08	1.58	-0.24	0.08**	-0.32
KM 2312	0.36	1.79**	0.06	0.92	0.30	-0.19**	-0.49
KM 2318	0.93**	2.77**	-0.19	0.31	-0.37	-0.14**	0.96
KM 2319	-1.02**	0.93**	-0.21	1.92	-0.48	-0.20**	-0.21
KM 2356	0.28	-0.35	0.06	-0.39	0.30	0.15**	0.93**
KM 2358	0.19	0.14	-0.18*	-4.81**	-0.07**	-0.15**	-1.47**
Tester							
KM 2195	-0.39**	-0.48**	0.08	1.41**	-0.04	0.02**	0.59**
KM 2241	0.21	0.29	-0.12*	-0.89**	-0.04	-0.01*	-0.26**
IPM 99-125	0.16	0.21	0.04	-0.52**	0.9	-0.01*	0.22*
S.E (gi) ₋ ⁺	0.11	0.15	0.06	0.15	0.12	0.01	0.11
S.E (gj) ₋ ⁺	0.30	0.33	0.10	0.32	0.26	0.02	0.24
S.E (gi-gi) ₋ ⁺	0.28	0.32	0.09	0.31	0.27	0.01	0.23
S.E (gj-gj) ₋ ⁺	0.52	0.59	0.16	0.33	0.49	0.03	0.42

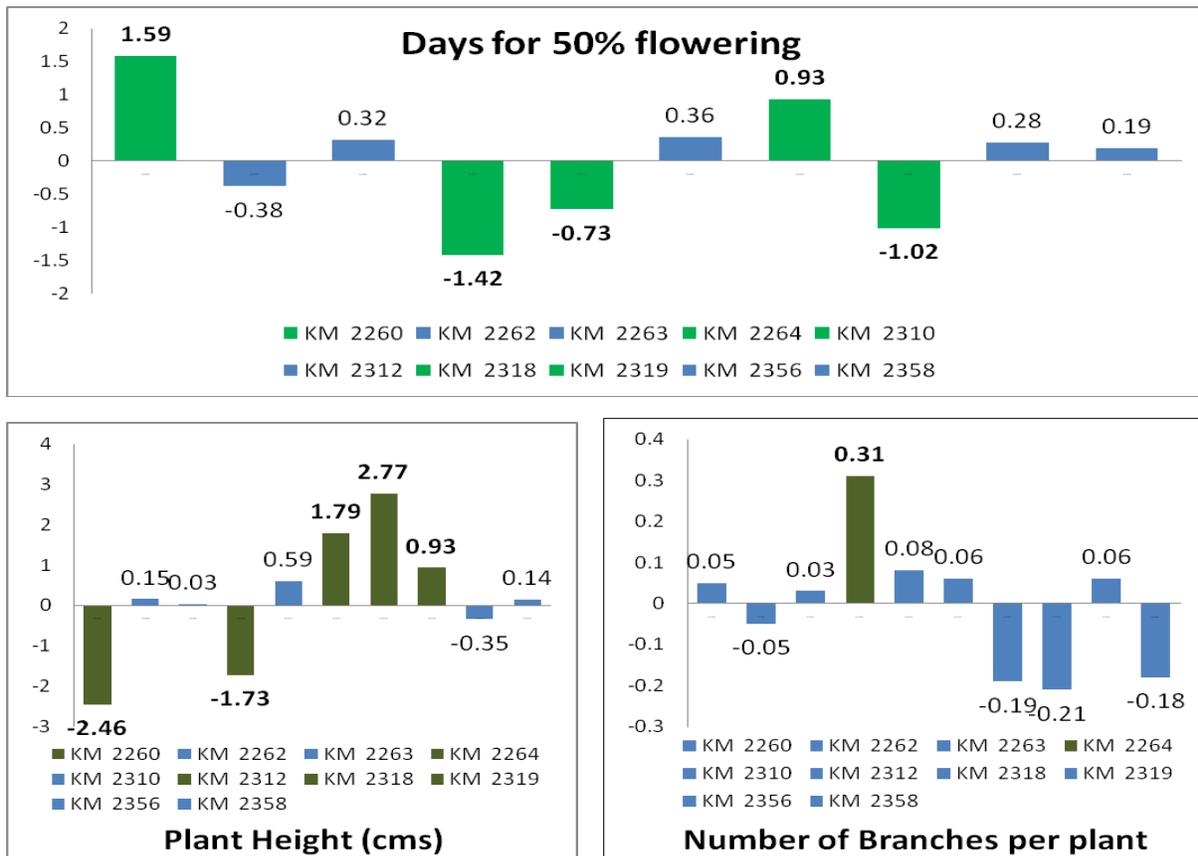
*, ** Significant at P=0.05 and P=0.01, respectively.

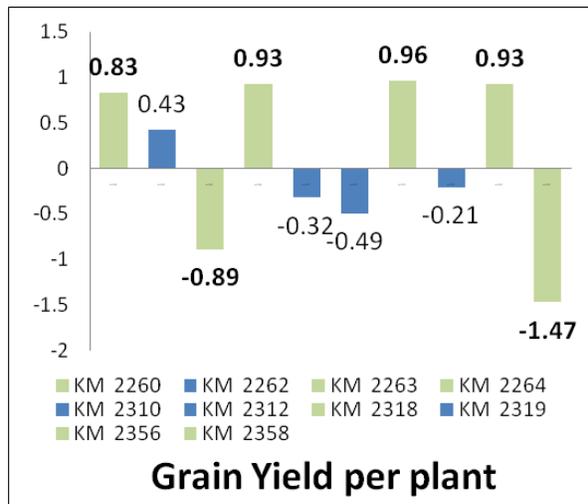
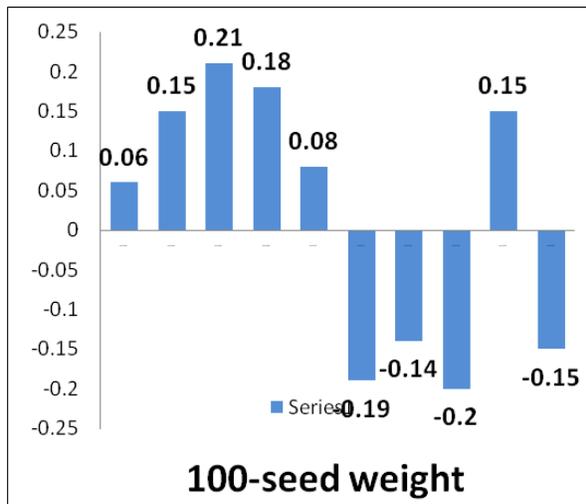
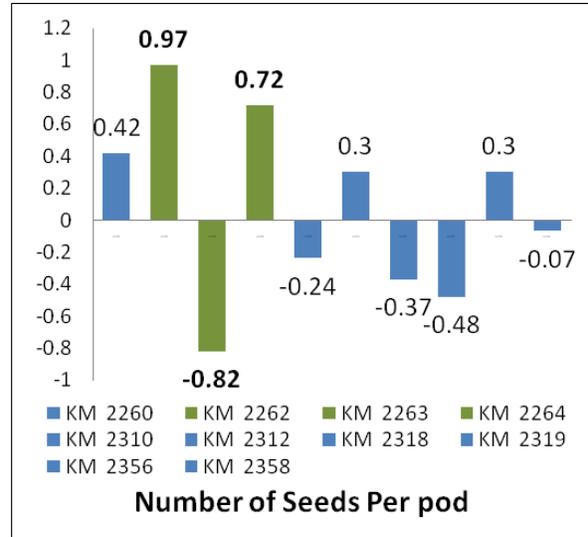
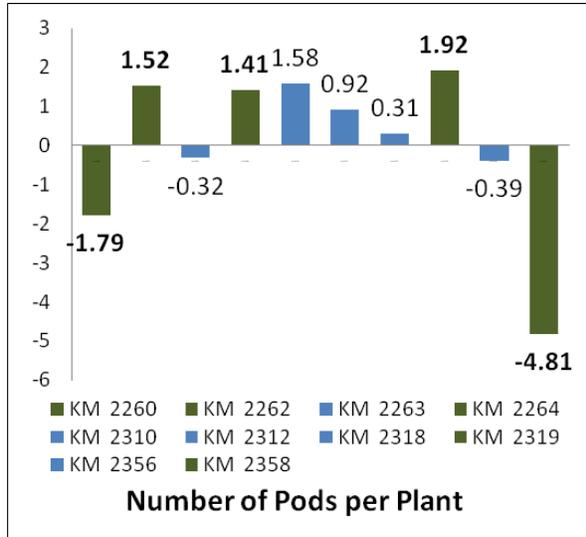
Table.3 Good specific combiners for grain yield, their performance for some other traits and gca effect of the parents involved under line x tester mating design in mungbean

Cross Grain yield	Per se performance	sca effect	gca effect		Traits for which cross also exhibited desirable sca effects
			P ₁	P ₂	
KM 2312 × KM 2241	16.21	3.92**	-0.29	0.53	number of pods per plant (5.03**) & 100-grain weight (0.25**)
KM 2260 × KM 2241	15.89	2.83**	-0.85	0.62	number of grains per pod (1.93**) & 100-grain weight (0.16**)
KM 2243 × KM 2241	11.47	2.04**	0.64	0.59	number of branches (0.37**) & 100-grain weight (0.07**)
KM 2264 × KM 99-125	12.31	1.92**	-0.50	-0.27	number of grains per pod (1.93**) & 100-grain weight (0.16**)
KM 2318 × KM 2241	8.62	1.62**	-1.47	0.15	number of branches per plant (0.48**) & number of pods per plant (1.48**)
KM 2264 × KM 2195	11.35	1.39**	-0.51	-0.28	100-grain weight (0.28**) & number of pods per plant (2.89*)
KM 2248 × KM 2241	12.27	0.99**	0.93	0.49	number of pods per plant (1.67**)
KM 2310 × KM 2195	11.83	0.98**	0.21	-0.28	number of branches (0.37**) & 100-grain weight (0.07**)
KM 2262 × KM 2241	11.21	1.02**	0.86	0.51	Number of pods per plant (3.19**), number of grains per pod (1.47**), 100-seed weight (0.14**), & number of branches per plant (0.37**)
KM 2318 × KM 2241	13.20	0.83*	0.83	0.49	number of grains per pod (1.13**), 100-grain weight (0.17**) & number of branches per plant (0.51**)

*, ** Significant at P=0.05 and P=0.01, respectively.

Fig.1 Bar graph indicates the gca effects of ten-lines for various agro-morphological characters





In the present investigation desirable SCA effects for different characters were evaluated to identify potential hybrids (Table 3). The hybrid KM 2312 × KM 2241 has showed highest SCA effect for number of pods per plant and 100-seed weight. The cross KM 2260 × KM 2241 had showed higher SCA effect for number of grains per pod and 100-seed weight. KM 2243 × KM 2241 had showed highest SCA effects for the number of branches and 100-seed weight. The hybrid KM 2264 × KM 99-125 exhibited desirable SCA effect for number of grains per pod and 100-seed weight. The hybrid KM 2318 × KM 2241 had significant SCA effect for number of branches per plant and number of pods per

plant. Superior SCA effect for traits 100-seed weight and number of pods per plant was observed for KM 2264 × KM 2195. The cross KM 2248 × KM 2241 had significant SCA effect for only number of pods per plant. The crossing between KM 2310 and KM 2195 had significant SCA effect for number of branches and 100-seed weight. Significant SCA effect in desired direction was observed in hybrid KM 2262 × KM 2241 for number of pods per plant, number of grains per pod, 100-seed weight and number of branches per plant. The lowest significant SCA effect was observed for hybrid KM 2318 × KM 2241 for number of grains per pod, 100-seed weight and number of branches per plant.

Based on the *per se* performance (Table 3) the following hybrids arranged in descending order were found to have superior performance: KM 2312 × KM 2241; KM 2260 × KM 2241; KM 2318 × KM 2241; KM 2264 × KM 99-125 and KM 2310 × KM 2195.

Based on the present investigation carried out in mungbean under Line × Tester design, the traits plant height, number of branches per plant, 100-seed weight and grain yield per plant showed heterosis. Non-additive gene action was shown for the traits plant height, number of pods per plant, number of seeds per pod and grain yield per plant. Based on the GCA effect the lines KM 2264, KM 2262 and KM 2260 were found to have good combining ability for most of the agronomically important traits, hence can be exploited in further hybridization programme in mungbean improvement. Based on the *per se* performance and specific combining ability, the hybrid KM 2312 × KM 2241; KM 2260 × KM 2241 and KM 2318 × KM 2241 were found to be superior and hence these hybrids can be utilized in further studies to know the stability of these hybrids in various environmental conditions.

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