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Heterosis and Combining Ability Studies for Quality Traits in Soybean [*Glycine max* (L.) Merrill]

Indu Rialch^{1*}, Jai Dev² and Bhupender Kumar³

¹Department of Plant Breeding and Genetics, Punjab Agricultural University, Ludhiana- 141004, India

²Chaudhary Sarwan Kumar Himachal Pradesh Krishi Vishvavidyalaya, Palampur-176062, India

³ICAR-Indian Institute of Maize Research, New Delhi- 110012, India

*Corresponding author

ABSTRACT

Soybean occupies, as importance, the first place in the group of grain legumes due to its use in human nutrition, as a source of very good protein and oil quality, as herb and because it is a very good prior plant to the ground, leaving a reserve of 80 to 140 kg N/ha. Heterosis and combining ability studies were carried in 54 crosses derived from crossing of 18 soybean lines with 3 testers. Based on various estimates viz., mean performance, heterosis and combining ability the best soybean parents and crosses for quality traits were identified. Combining ability analyses indicated that Shivalik showing the largest general combining ability (GCA) effect for protein and oil contents was the best combiner, and identified several best cross combinations with high positive specific combining ability (SCA) effects. These crosses involved geographically diverse parents and at least one parent of high General Combining Ability (GCA) effects in combination of high x high, high x average and high x low. Therefore, these crosses could throw transgressive segregants in still advance generations. The results of present study suggested that heterosis coupled with high SCA effects might be considered as criteria for selecting crosses for further improvement with respect to quality traits.

Keywords

Soybean, Protein and oil quality, GCA.

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Introduction

Soybean [*Glycine max* (L.) Merrill] is a wonderful crop gifted by the nature to mankind which is one of the richest sources of oil as well as protein. It belongs to the family Leguminosae and is a self-pollinated crop having chromosome number of $2n=40$. Soybean is one of the most important crops that have the potential to provide the world's increasing demand for food and forage. The spread of soybean from its native land of origin has been mainly due to its adaptability and predominant use as a food crop for

human nutrition, as a protein source for animals, as a medicinal plant and lately as an industrial crop (Alghamdi, 2004). Soybean is the cheapest, richest and easiest source of best quality protein. Owing to multiplicity of uses as food and industrial products, soybean is called as "Wonder crop". Soybean seed contains about 20.5% carbohydrates, 20% oil with poly unsaturated fatty acids specially Omega 6 and Omega 3 fatty acid, 4.5% minerals, 3.7% fibre, 8.1% water and 43.2% protein, its protein is complete and contains

all the essential amino acids therefore from nutritional point of view it is known as “Miracle bean”. It also contains a good amount of salts, vitamins and health promoting phyto-chemicals for human and livestock. Soybean is an environment friendly grain legume and its cultivation improves soil health because of its atmospheric nitrogen fixing ability and deep root system. Soybean is a dominant oil seed crop in the world trade accounting about 25% world’s total oil and fats production. Soybean has become the major source of edible vegetable oil and high quality proteins for food and feed supplement all over the world. Demand for soybean is still raising day by day as it has become major oil seed and also serves as raw material for various industries. To develop elite and better nutritional quality varieties such cultivars a systematic approach has to be adopted. Protein and oil are polygenic characters and their improvement is basically based upon identification and selection of superior parents or lines, which can be effectively used for developing superior quality genotypes for breeding programme. Combining ability studies are regarded useful in assessing the relative nicking ability of parents, which on crossing would produce more desirable segregants and elucidate the nature and magnitude of gene action involved. The study becomes much more reliable if a good degree of correspondence is observed between various genetic estimates over filial generations. Further, extent of heterosis will have direct effect on breeding methodology in varietal improvement programmes. Therefore, keeping aforesaid considerations in view, present study was undertaken in soybean to evaluate, identification and selection of better nutritional quality genotypes.

Materials and Methods

The experimental materials comprised 18 fixed lines, three testers of soybean and their

54 triple test cross families. Two agronomically superior and diverse genotypes, Him soya and Hara soya (P_1 and P_2) and their F_1 s were used as testers ‘ L_1 ’, ‘ L_2 ’ and ‘ L_3 ’, respectively. Details of material used are given in Table 1. The parents were selected on the basis of diverse pedigree, growth habit, geographical origin and adaptation. The fifty four F_1 hybrids, 18 lines and 3 testers were grown in a completely randomized block design with three replications during *kharif* 2013. The experimental plot of each treatment consisted of one row of 2m length. Row to row and plant to plant distances were maintained at 50 and 20 cms, respectively.

Protein and oil content in seed samples of each genotype for each replication was estimated with the help of Infratec™1241 grain analyzer which uses near-infrared transmittance technology to test multiple parameters, *viz.*, moisture, protein, oil, starch etc. in a broad range of grain and oilseed commodities (Sudar *et al.*, 2007). Different types of heterosis *viz.*, heterosis over mid parent, heterobeltiosis over better parent and economic heterosis over standard check were estimated using mean values. Analysis of line x tester data for combining ability was done according to Kempthorne (1957) and Arunachalam (1974).

Results and Discussion

Analysis of variance was carried out for lines, tester, their crosses and that revealed highly significant differences for all the lines and crosses indicating sufficient variation in the material for both protein and oil content. However, mean squares due to testers were significant for only protein content. The study for mean and range revealed that among parents there was great variability in the lines than testers for both the characters. It was reflected that the hybrids superior in F_1

involved both or at least one parent of high *per se* performance.

For protein content, the highest positive heterosis over both mid parent and better parent was recorded for H-330 x Him Soya. Out of 36 hybrids, 25 exhibited positive heterosis over mid parent and 22 exhibited positive heterosis over better parent. Out of these crosses exhibiting positive heterosis, there were 20 and 13 crosses exhibiting significant heterosis over mid and better parent, respectively.

Gadag and Upadhyaya (1995) and Sabbouh *et al.*, (1998) also observed significant positive heterosis for protein content. The range of heterosis over mid parent was from -6.06 to 8.65 per cent and over better parent it was from -11.15 to 8.58 per cent (Table 2).

In case of oil content, the range of heterosis over mid parent was from -17.89 to 12.44 per cent and over better parent it was from -24.07 to 11.12 per cent for this trait. Cross H-330 x Him Soya exhibited the highest significant heterosis over mid parent and better parent. There were 10 hybrids which showed positive heterosis over mid parent and five hybrids indicated positive heterosis over better parent. Four crosses, *viz.*, H-330 x Him Soya, P6-1 x Him Soya, P169-3 x Him Soya and H-330 x Hara Soya over mid parent and only one cross H-330 x Him Soya over better parent had significant heterosis. Similar results have been reported by Chen (1983), Wehrmann *et al.*, (1987), Fahmi *et al.*, (1999) and several other workers.

In autogamous crops like soybean, combining ability provides required information about the nature and magnitude of genetic variation along with nicking ability of genotypes. Over and above, the study become much more precise and reliable if the relative combining ability is analyzed in F1, F2, and F3

generations so as to obtained a good degree of correspondence between these estimates which will help in identifying superior parents and crosses that could be effectively utilized in breeding programme. Such study may also reflect the possibility of obtaining rare favourable combinations of characters in segregating generations.

Analysis of variance for combining ability for both the quality traits indicated that variance due to crosses, lines and line x tester interaction were significant for protein and oil content. Significance of lines and testers indicated higher contribution towards general combining ability, while for lines x tester interaction towards specific combining ability for both the characters. Predominance of estimated component of Specific Combining Ability (SCA) variance over General Combining Ability (GCA) indicated importance of non-additive genetic effects for protein and oil content. Contribution of lines was higher for protein content, whereas higher contribution due to interaction between line and tester was for oil content.

On the basis of heterosis, it can be concluded that heterosis was displayed for almost all the characters. Seed yield is the sum total of contribution made by individual component and hence a complex character itself. The breeder has to simplify complex situation to breed for high yield by handling number of related traits simultaneously. A good number of crosses showed the presence of desirable heterotic response for the different characters over the standard cultivars.

Among the parents lines Shivalik, SL-795, P2-2, P6-1 and H-330 recorded positive significant GCA effects for protein content. While for oil content Shivalik, PS-1466, SL-679, P13-4 and PS-1469 could be considered superior as these exhibited high GCA effects (Table 3).

Table.1 Soybean lines used to generate hybrids in this study

S. No.	Lines	S. No.	Lines
1	SL-682	12	P2-2
2	P6-1	13	H-330
3	SL-679	14	PS-1469
4	P9-2-2	15	VLS-59
5	DS1213	16	JS-335
6	PK-472	17	P169-3
7	Hardee	18	P13-4
8	Bragg		Testers
9	SL-795	1	Him Soya (L ₁)
10	Shivalik	2	Hara Soya (L ₂)
11	PS-1466	3	Hara Soya x Him Soya (L ₃)

Table.2 Estimation of heterosis (MP), heterobeltiosis (BP), economic heterosis (EP) and in breeding depression for protein and oil content in soybean

Cross	Seed protein content (%)			Seed oil content (%)		
	MP	BP	EP	MP	BP	EP
P9-2-2 x Him Soya	7.13*	6.67*	7.61*	-2.47	-4.65*	-4.65*
P9-2-2 x Hara Soya	0.93	-0.65	0.23	-5.24*	-13.18*	-0.36
PS-1466 x Him Soya	-6.06*	-10.57*	-1.06	1.66	-0.42	-0.42
PS-1466 x Hara Soya	-5.65*	-11.15*	-1.70	-4.64*	-12.47*	0.46
P6-1 x Him Soya	-2.38*	-4.48*	-0.19	7.83*	2.81	2.81
P6-1 x Hara Soya	-0.35	-3.58*	0.76	1.03	-9.58*	3.78*
PK-472 x Him Soya	0.00	-0.90	0.92	2.29	1.75	1.75
PK-472 x Hara Soya	5.29*	3.17*	5.07*	-11.30*	-17.42*	-5.22*
VLS-59 x Him Soya	3.20*	2.89*	2.89*	-1.07	-2.79	-2.79
VLS-59 x Hara Soya	3.40*	2.53*	1.92*	-8.74*	-15.99*	-3.59*
P13-4 x Him Soya	0.65	0.45	0.84	0.64	-1.61	-1.61
P13-4 x Hara Soya	3.17*	1.81	2.20*	-5.80*	-13.70*	-0.95
PS-1469 x Him Soya	3.58*	3.53*	3.63*	1.27	1.12	1.12
PS-1469 x Hara Soya	3.67*	2.44*	2.54*	-4.39*	-10.67*	2.52
DS-1213 x Him Soya	2.01*	0.98	3.06*	-3.83*	-6.90*	-0.55
DS-1213 x Hara Soya	-0.29	-2.40*	-0.39	-7.75*	-10.95*	2.20
SL-679 x Him Soya	5.23*	4.14*	6.33*	-1.11	-5.09*	-5.09*
SL-679 x Hara Soya	7.77*	5.47*	7.69*	-15.68*	-24.07*	-12.85*
SL-682 x Him Soya	2.32*	0.29	4.42*	-3.32*	-4.45*	-2.16
SL-682 x Hara Soya	3.84*	0.65	4.80*	-8.97*	-13.88*	-1.16
H-330 x Him Soya	8.65*	8.58*	8.58*	12.44*	11.12*	13.80*
H-330 x Hara Soya	7.65*	6.49*	6.36*	4.06*	-1.54	13.00*
P169-3 x Him Soya	1.96*	1.25	1.25	-2.78*	-2.94	-2.94
P169-3 x Hara Soya	1.51	1.06	-0.35	6.50*	-0.51	14.18*
P2-2 x Him Soya	3.68*	3.35*	4.02*	-7.64*	-9.39*	-5.83*
P2-2 x Hara Soya	3.70*	2.20*	2.86*	-15.18*	-19.19*	-7.25*
Shivalik x Him Soya	-1.58	-2.59*	-0.55	-2.63*	-5.18*	0.06
Shivalik x Hara Soya	2.40*	1.53*	1.92*	-5.14*	-8.96*	4.48*
Hardee x Him Soya	-0.51	-1.88*	0.91	-0.01	-1.95	2.01
Hardee x Hara Soya	1.43	-1.09	1.72	-8.48*	-12.75*	0.13
JS-335 x Him Soya	0.32	-0.56	1.22	-0.97	-1.84	-1.84
JS-335 x Hara Soya	2.61*	0.57	2.37*	-10.15*	-16.62*	-4.31*
SL-795 x Him Soya	2.24*	1.22	1.22	-7.48*	-12.81*	-1.44
SL-795 x Hara Soya	4.45*	4.30*	2.22*	-17.89*	-18.51*	-6.47*
Bragg x Him Soya	-3.39*	-6.20*	-0.41	0.36	0.06	0.66
Bragg x Hara Soya	-2.18*	-6.07*	-0.27	-1.80	-7.86*	5.75*

* Significant at P ≤ 0.05

Table.3 General Combining Ability (GCA) effects for quality traits in soybean

Genotype	Seed protein content (%)	Seed oil content (%)
P6-1	0.62*	-0.43*
SL-682	-1.36*	0.02
SL-679	-0.74*	0.59*
P9-2-2	0.27	-0.29*
DS-1213	0.05	-0.55*
PK-472	-0.27	-0.21
Hardee	0.31	0.33*
Bragg	-0.35	0.16
SL-795	1.77*	-1.56*
P2-2	0.88*	-0.28*
Shivalik	1.95*	2.36*
PS-1466	-0.67*	1.00*
H-330	0.44*	-1.13*
PS-1469	-1.19*	0.41*
VLS-59	-0.35	0.20
JS-335	-0.17	-0.53*
P169-3	-0.20	-0.68*
P13-4	-0.97*	0.57*
SE(gi) ±	0.18	0.13
SE(gi-gj) ±	0.25	0.18
Testers		
Him Soya	0.08	-0.06
Hara Soya	-0.08	0.06
SE (gi) ±	0.06	0.04
SE (gi-gk) ±	0.08	0.06

* Significant at $P \leq 0.05$

Table.4 Specific Combining Ability (SPA) effects for quality traits in soybean

S.N.	Crosses	Protein content (%)	Oil content (%)	S.N.	Crosses	Protein content (%)	Oil content (%)
1	P9-2-2 x Him Soya	1.29*	-0.32	20	SL-682 x Hara Soya	0.15	0.03
2	P9-2-2 x Hara Soya	-1.29*	0.32	21	H-330 x Him Soya	0.33	0.13
3	PS1466 x Him Soya	0.04	-0.02	22	H-330 x Hara Soya	-0.33	-0.13
4	PS1466 x Hara Soya	-0.04	0.02	23	P169-3 x Him Soya	0.21	-1.45*
5	P6-1 x Him Soya	-0.26	-0.03	24	P169-3 x Hara Soya	-0.21	1.45*
6	P6-1 x Hara Soya	0.26	0.03	25	P2-2 x Him Soya	0.13	0.18
7	PK-472 x Him Soya	0.86*	0.67*	26	P2-2 x Hara Soya	-0.13	-0.18
8	PK-472 x Hara Soya	-0.86*	-0.67*	27	Shivalik x Him Soya	-0.96	-0.53
9	VLS-59 x Him Soya	0.09	0.13	28	Shivalik x Hara Soya	0.96	0.53*
10	VLS-59 x Hara Soya	-0.09	-0.13	29	Hardee x Him Soya	-0.23	0.22
11	P13-4 x Him Soya	-0.34	-0.002	30	Hardee x Hara Soya	0.23	-0.22
12	P13-4 x Hara Soya	0.34	0.002	31	JS-335 x Him Soya	-0.29	0.27
13	PS1469 x Him Soya	0.12	-0.07	32	JS-335 x Hara Soya	0.29	-0.27
14	PS1469 x Hara Soya	-0.12	0.07	33	SL-795 x Him Soya	-0.27	0.49*
15	DS1213 x Him Soya	0.56*	-0.18	34	SL-795 x Hara Soya	0.27	-0.49*
16	DS1213 x Hara Soya	-0.56*	0.18	35	Bragg x Him Soya	-0.11	-0.39*
17	SL-679 x Him Soya	-0.34	0.74*	36	Bragg x Hara Soya	0.11	0.39*
18	SL-679 x Hara Soya	0.34	-0.74*		SE (Sij) ±	0.25	0.18
19	SL-682 x Him Soya	-0.15	-0.03		SE (Sij-Skl) ±	0.36	0.26

* Significant at $P \leq 0.05$

Table.5 Best parents and crosses selected on the basis of their *per se* performance, GCA effects, SCA effects and heterotic response for protein and oil content in soybean

Characters	Protein content	Oil content
Best parents <i>per se</i>	Shivalik, SL-795, P2-2, P6-1 and H-330	Shivalik, PS-1466, SL-679, P13-4 and PS-1469
Best crosses <i>per se</i>	H-330 x Him Soya, SL-679 x Hara Soya, P9-2-2 x Him Soya, H-330 x Hara Soya, SL-679 x Him Soya	P169-3 x Him Soya, H-330 x Him Soya, H-330 x Hara Soya, Bragg x Hara Soya, P6-1 x Hara Soya
Best heterotic Crosses	H-330 x Him Soya, SL-679 x Hara Soya, P9-2-2 x Him Soya, Shivalik x Hara Soya	P169-3 x Him Soya, H-330 x Him Soya, H-330 x Hara Soya, Bragg x Hara Soya, Shivalik x Hara Soya
Best GCA parents	H-330 x Hara Soya, SL-679 x Him Soya Shivalik, SL-795, P2-2, P6-1 and H-330	Shivalik, PS-1466, SL-679, P13-4 and PS-1469
Best SCA Crosses	P9-2-2 x Him Soya, PK 472 x Him Soya, Shivalik x Hara Soya and DS-1213 x Him Soya	P169-3 x Hara Soya, SL-679 x Him Soya, PK 472 x Him Soya, Shivalik x Hara Soya, SL-795 x Him Soya and Bragg x Hara Soya

The *per se* performance of most of the parent was moderate to high for protein and oil content. Among the hybrids, positive significant SCA effects for protein content were exhibited by P9-2-2 x Him Soya, PK-472 x Hara Soya and DS-1213 x Him Soya. On the other hand many crosses viz., P169-3 x Hara Soya, SL-679 x Him Soya, PK 472 x Him Soya, SL-795 x Him Soya and Bragg x Hara Soya showed significantly high SCA effects for oil content (Table 4). Among these hybrids most of them had high *per se* and involving good general combiner parents, hence, could be used effectively in quality improvement of soybean. Therefore on the basis of combining ability effects, cross PK 472 x Him Soya appeared promising for both protein and oil content. While P9-2-2 x Him Soya and P169-3 x Hara Soya were good specific combiner for protein and oil content, respectively. Similar to present findings Mc Kendry *et al.*, (1985) and Gadag *et al.*, (1999) also reported importance of non-additive gene effects for protein content. On the contrary, Zhu *et al.*, (1994) and Sabbouh *et al.*, (1998) noted significant GCA and additive effects for protein content in soybean. In case of oil content, as per present study Sharma *et al.*, (1997) and Sabbouh *et al.*, (1998) recorded partial dominance and non-additive gene

action and Raut *et al.*, (2000) reported dominance gene effects in 2 crosses for seed oil content in soybean. Contrary to this, significant additive gene action was reported by Zhu *et al.*, (1994) for oil content.

Based on various estimates viz., mean performance, heterosis and combining ability the best soybean parents and crosses for quality traits were identified (Table 5). For both protein and oil content, high heterosis and positive SCA effects were recorded in cross Shivalik x Hara Soya and could be further exploited. This cross involved geographically diverse parents and at least one parent of high GCA effects in combination of high x high, high x average and high x low. Majority of soybean breeding work is based on pedigree method, which exploits the fixable additive type of gene effects. However, this alone may be inadequate in view of the large non-additive component for both quality traits. Further, high heterosis for oil and protein along with positive SCA effects was observed in these crosses, which might be due to dominance or epistasis. Although heterosis breeding makes maximal use of the non-additive genetic effects appears to be difficult for improving soybean in view of the non-availability of

mass pollination systems needed for hybrid seed production. Accordingly, the feasible alternative is to consider simultaneous exploitation of both additive and non-additive gene action by adopting recurrent selection procedures. Under similar situation Harer and Deshmukh (1991) recommended biparental mating in soybean followed by recurrent selection. Since crossing is tedious in soybean, genetic male sterility and single seed or single pod descent method could be employed to facilitate recurrent selection schemes (Wilcox, 1998). Therefore, these crosses could throw transgressive segregants in still advance generations. Further, the results of present study suggested that heterosis coupled with high SCA effects might be considered as criteria for selecting cross for further improvement with respect to quality traits.

To conclude, combining ability mean squares estimated revealed that both the quality traits displayed preponderance of non-additive gene effects. Often the unfavorable association between protein and yield generally met with are due to tight linkages, which could be broken by raising a large population. In the present study H-330 x Him Soya revealed superiority for protein content, oil content as well as yield might be considered a new favorable combination resulted due to breaking of tight linkages. Through the use of recurrent selection suggested that such negative relationship may be caused by linkage and improvements can be made for such traits by enforcing simultaneous selection in large segregation population.

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