

Original Research Article

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Improvement in Seed Yield and Related Traits of Linseed Genotypes (*Linum usitatissimum* L.) through Various Selection Parameters in Mid-Hills of North-West Himalayas

Satish Paul, Nimit Kumar* and Pankaj Chopra

Department of Crop Improvement, CSK Himachal Pradesh Krishi Vishvavidyalaya,
Palampur-176062, India

*Corresponding author

ABSTRACT

Keywords

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Forty five linseed genotypes were subjected to study the interrelationship among the traits at the Experimental Farm of the Department of Crop Improvement, CSK Himachal Pradesh Krishi Vishvavidyalaya, Palampur, during *rabi* 2015-2016. The genotypic and phenotypic correlation coefficient obtained between different traits was similar in direction, while in magnitude, genotypic correlation was higher than the corresponding phenotypic correlations for most of the traits. Seed yield per plant had maximum significant and positive genotypic and phenotypic association with biological yield per plant followed by harvest index and seeds per capsule. Harvest index plant contributed indirectly through plant height followed by technical height and biological yield per plant has maximum indirect effects through capsules per plant on seed yield per plant. Principal component analysis (PCA) showed that seed yield per plant, harvest index, biological yield per plant seeds per capsule and capsules per plant were found in same group.

Introduction

Lineed (*Linum usitatissimum* L.; n=15) is an important oilseed crop which is the only species in Linaceae family with economic values (Tadesse *et al.*, 2009). It has nutrients and pharmaceutical uses and used for edible and lightening purposes and also in animal fat and poultry diets (Khan *et al.*, 2010). Linseed contain 30-45% oil, making it an important industrial crop. It has high unsaturated fatty acids, especially Linolenic acid (Khan *et al.*, 2010). Due to less impression of direct selection for yield, more efforts should be over indirect selection for yield components. Proper understanding of association of

different traits, provide more reliable selection criterion to achieve a high seed yield (Akbar *et al.*, 2001). Correlation coefficients may not evolve satisfactory results in uncovering the real interrelationships among the traits. Nevertheless, selection for yield via highly correlated traits becomes easy if the contribution of different characters to yield is quantified using path coefficient analysis (Dewey & Lu, 1959). Multivariate statistical techniques which simultaneously analyze multiple measurements on each individual under investigation are widely used in analysis of interrelationships. Among the

Multivariate techniques, principal component analysis (PCA) had been shown to be very useful in selecting genotypes for breeding program that meet the objective of a plant breeder (Mohammadi and Prasanna, 2003). PCA may be used to reveal patterns and eliminate redundancy in data sets (Adams, 1995) as morphological and physiological variations routinely occur in crop species. The present investigation was carried out to study the associations among yield and yield related traits in linseed.

Materials and Methods

Interrelationships for various traits was studied in 45 local collection (Table 1) of linseed during *rabi* 2015-16 at Experimental Farm of the Department of Crop Improvement, CSK Himachal Pradesh Krishi Vishvavidyalaya, Palampur, India (32°8' N, 76°3' E) represents humid sub-temperate climate zone with annual rainfall of 2500mm and acidic soil with pH of 5.0 to 5.6. The experiment was conducted using randomized complete block design with three replications. Each replication consisted of three rows of each genotype. Row to row distance was 30 cm with row length of 3 meter and plant to plant distance was 10 cm. Data was recorded on five randomly selected plants in each replication for plant height, technical height, primary branches per plant, secondary branches per plant, capsules per plant, biological yield per plant, seeds per capsule, seed yield per plant and harvest index calculated as:-

$$\text{Harvest index (\%)} = \frac{\text{Seed yield per plant}}{\text{Biological yield per plant}} \times 100$$

Statistical analysis: The recorded data was subjected to analysis of genotypic and phenotypic correlation coefficients as suggested by Al-Jibouri *et al.*, (1958) and the path coefficient analysis was conducted as suggested by Dewey and Lu (1959).

PCA: Principal component analysis (PCA) analysis was performed using XLSTAT software to determine the best relationships among characters.

Results and Discussion

The possible increase the seed yield through yield related traits, as primary target of crop improvement, requires understanding the inter relationship between various yield contributing traits or in fact, yield components. The correlation coefficients between different characters are given in Table 2. The results of experiment revealed that the traits plant height, secondary branches per plant, capsules per plant, biological yield per plant, seeds per capsule and harvest index had the strong positive association with seed yield per plant at both genotypic and phenotypic levels. Whereas, technical height and primary branches per plant exhibited positive but non-significant associations with seed yield per plant. The genotypic and phenotypic correlation coefficient was similar in directions, while in magnitude, genotypic correlations were mostly higher than corresponding phenotypic correlations. Similar findings were reported by Sohan *et al.*, (2004), Joshi (2004). Thus the low phenotypic correlation could results due to the masking and modifying effect of environment on the association of characters at genotypic level. On the basis of present studies, it can be concluded that the selection based on traits *viz.*, plant height, secondary branches per plant, capsules per plant, biological yield per plant, seeds per capsule and harvest can provide better result for improvement of seed yield per plant in linseed, as earlier reported by Tariq *et al.*, (2014) plant height, number of capsules per plant, number of seeds per capsule, Yadav (2001) for number of capsules per plant and number of seeds per capsule, Muhammad *et al.*, (2003) for number of branches per plant.

Table.1 Details of material used in present study

S. No.	Collection No.	Crop Name	Botanical Name	Site of collection		State
				Village	Disst.	
1	KLSA-1	Linseed	<i>Linum usitatissimum</i>	Utrala	Kangra	H.P.
2	KLSA-2	Linseed	<i>Linum usitatissimum</i>	Harer Baijnath	Kangra	H.P.
3	KLSA-3	Linseed	<i>Linum usitatissimum</i>	Drugh Nala Baijnath	-	H.P.
4	KLSA-4	Linseed	<i>Linum usitatissimum</i>	Harer Baijnath	Kangra	H.P.
5	KLSA-5	Linseed	<i>Linum usitatissimum</i>	Balh Harer Baijnath	Kangra	H.P.
6	KLSA-6	Linseed	<i>Linum usitatissimum</i>	Balh Harer Baijnath	Kangra	H.P.
7	KLSA-7	Linseed	<i>Linum usitatissimum</i>	Dramlu Harer Baijnath	Kangra	H.P.
8	KLSA-8	Linseed	<i>Linum usitatissimum</i>	Dramlu Harer Baijnath	Kangra	H.P.
9	KLSA-9	Linseed	<i>Linum usitatissimum</i>	Kholi Deol Baijnath	Kangra	H.P.
10	KLSA-10	Linseed	<i>Linum usitatissimum</i>	Phatar	Kangra	H.P.
11	KLSA-11	Linseed	<i>Linum usitatissimum</i>	Bhattu	-	H.P.
12	KLSA-12	Linseed	<i>Linum usitatissimum</i>	Tramal	-	H.P.
13	KLSA-13	Linseed	<i>Linum usitatissimum</i>	Dak Bangra Chauotra	Mandi	H.P.
14	KLSA-14	Linseed	<i>Linum usitatissimum</i>	Chauotra Joginder Nagar	Mandi	H.P.
15	KLSA-15	Linseed	<i>Linum usitatissimum</i>	Hara Bagh Joginder Nagar	Mandi	H.P.
16	KLSA-16	Linseed	<i>Linum usitatissimum</i>	Hara Bagh Joginder Nagar	Mandi	H.P.
17	KLSB-1	Linseed	<i>Linum usitatissimum</i>	Jia	Kangra	H.P.
18	KLSB-2	Linseed	<i>Linum usitatissimum</i>	Jia	Kangra	H.P.
19	KLSB-3	Linseed	<i>Linum usitatissimum</i>	Jia	Kangra	H.P.
20	KLSB-4	Linseed	<i>Linum usitatissimum</i>	Jagehar	Kangra	H.P.
21	KLSB-5	Linseed	<i>Linum usitatissimum</i>	Chamotu Jia	Kangra	H.P.
22	KLSB-6	Linseed	<i>Linum usitatissimum</i>	Chamotu Jia	Kangra	H.P.
23	KLSB-7	Linseed	<i>Linum usitatissimum</i>	Chamotu Jia	Kangra	H.P.
24	KLC-3	Linseed	<i>Linum usitatissimum</i>	Nagehar	-	H.P.
25	KLC-4	Linseed	<i>Linum usitatissimum</i>	Nagehar	-	H.P.
26	KLC-9	Linseed	<i>Linum usitatissimum</i>	Baijnath	Kangra	H.P.
27	KLC-10	Linseed	<i>Linum usitatissimum</i>	Arki	Solan	H.P.
28	KLC-11	Linseed	<i>Linum usitatissimum</i>	Ahju Baijnath	Kangra	H.P.
29	KLC-12	Linseed	<i>Linum usitatissimum</i>	Kandi ,Palampur	Kangra	H.P.
30	KLC-13	Linseed	<i>Linum usitatissimum</i>	Trehal ,Baijnath	Kangra	H.P.
31	KLC-14	Linseed	<i>Linum usitatissimum</i>	Utrala Baijnath	Kangra	H.P.
32	KLD-1	Linseed	<i>Linum usitatissimum</i>	Patti Panchrukhi	Kangra	H.P.
33	KLD-2	Linseed	<i>Linum usitatissimum</i>	Patti Panchrukhi	Kangra	H.P.
34	KLD-3	Linseed	<i>Linum usitatissimum</i>	Chandropa Patti Panchrukhi	Kangra	H.P.
35	KLD-4	Linseed	<i>Linum usitatissimum</i>	Palah Patti Panchrukhi	Kangra	H.P.
36	KLD-5	Linseed	<i>Linum usitatissimum</i>	Patti Panchrukhi	Kangra	H.P.
37	KLD-6	Linseed	<i>Linum usitatissimum</i>	Chandropa Patti Panchrukhi	Kangra	H.P.
38	KLD-7	Linseed	<i>Linum usitatissimum</i>	Palah Patti Panchrukhi	Kangra	H.P.
39	KLD-8	Linseed	<i>Linum usitatissimum</i>	Palah Patti Panchrukhi	Kangra	H.P.
40	KLD-10	Linseed	<i>Linum usitatissimum</i>	Chandropa Patti Panchrukhi	Kangra	H.P.
41	T-397	Linseed	<i>Linum usitatissimum</i>			
42	Baner	Linseed	<i>Linum usitatissimum</i>			
43	Nagarkot	Linseed	<i>Linum usitatissimum</i>			
44	Him Alsi-2	Linseed	<i>Linum usitatissimum</i>			
45	Himani	Linseed	<i>Linum usitatissimum</i>			

Table.2 Estimates of genotypic (G) and phenotypic (P) correlation coefficients among different traits of linseed

		Technical height (cm)	Primary branches per plant	Secondary branches per plant	Capsules per plant	Biological yield per plant (gm)	Seeds per capsule	Harvest index (%)	Seed yield per plant (gm)
Plant Height (cm)	P	.4987**	.1835*	.2110*	.0880	.0502	.0907	.2134*	.1682*
	G	.6970**	.2043*	.2035*	.1138	.0560	.2237**	.2862**	.1686*
Technical height (cm)	P		.0571	.0749	-.0954	-.0981	.1687*	.1948*	.0292
	G		.0765	.1092	-.1084	-.1229	.3203**	.2315**	.0208
Primary branches per plant	P			.8811**	.6562**	.0931	.3201**	-.0200	.1042
	G			.9741**	.7206**	.1031	.6052**	-.0340	.0986
Secondary branches per plant	P				.6153**	.1257	.2676**	.0090	.1715*
	G				.6978**	.1410	.6234**	.0141	.1941*
Capsules per plant	P					.2089*	.1700*	-.0802	.1827*
	G					.2117*	.2544**	-.0995	.1825*
Biological yield per plant (gm)	P						-.0943	-.3133**	.5534**
	G						-.1790	-.3263**	.5701**
Seeds per capsule	P							-.0455	.2051*
	G							-.1233	.2180*
Harvest index (%)	P								.5180**
	G								.5377**

Table.4 Eigenvectors and eigen values of 4 principal components for 9 characters of 45 linseed genotypes

	Plant Height (cm)	Technical height (cm)	Primary branches per plant	Secondary branches per plant	Capsules per plant	Biological yield per plant (gm)	Seeds per capsule	Harvest index (%)	Seed yield per plant (gm)	Eigenvalue	Variability (%)	Cumulative (%)
PC1	0.230	0.110	0.524	0.529	0.453	0.200	0.256	0.109	0.242	2.956	32.844	32.844
PC2	0.005	-0.163	-0.185	-0.147	0.010	0.527	-0.399	0.328	0.610	2.067	22.963	55.807
PC3	0.615	0.640	-0.147	-0.118	-0.231	-0.113	0.034	0.329	0.040	1.674	18.600	74.407
PC4	-0.064	-0.229	0.080	0.082	0.243	-0.533	-0.342	0.670	-0.144	0.978	10.866	85.273

Table.3 Estimates of genotypic (G) and phenotypic (P) direct and indirect effects of different traits on seed yield in linseed

		Plant Height (cm)	Technical height (cm)	Primary branches per plant	Secondary branches per plant	Capsules per plant	Biological yield per plant (gm)	Seeds per capsule	Harvest index (%)	Seed yield per plant (gm)
Plant Height (cm)	P	-.1041	.0004	-.0182	.0244	.0072	.0398	.0451	.1736	.1682*
	G	-.1320	.0131	-.1173	.1113	.0157	.0447	.0041	.2290	.1686*
Technical height (cm)	P	-.0519	.0009	-.0057	.0087	-.0079	-.0779	.0046	.1584	.0292
	G	-.0920	.0188	-.0439	.0597	-.0149	-.0980	.0058	.1852	.0208
Primary branches per plant	P	-.0191	.0000	-.0994	.1020	.0541	.0739	.0089	-.0162	.1042
	G	-.0270	.0014	-.5738	.5328	.0992	.0822	.0110	-.0272	.0986
Secondary branches per plant	P	-.0220	.0001	-.0876	.1158	.0507	.0998	.0075	.0073	.1715*
	G	-.0269	.0021	-.5590	.5470	.0961	.1124	.0113	.0113	.1941*
Capsules per plant	P	-.0092	-.0001	-.0652	.0713	.0824	.1658	.0031	-.0653	.1827*
	G	-.0150	-.0020	-.4135	.3817	.1376	.1687	.0046	-.0796	.1825*
Biological yield per plant (gm)	P	-.0052	-.0001	-.0093	.0146	.0172	.7937	-.0026	-.2548	.5534**
	G	-.0074	-.0023	-.0592	.0771	.0291	.7971	-.0032	-.2611	.5701**
Seeds per capsule	P	-.0094	.0001	-.0318	.0310	.0091	-.0749	.3180	-.0370	.2051*
	G	-.0295	.0060	-.3473	.3409	.0350	-.1427	.4542	-.0986	.2180*
Harvest index (%)	P	-.0222	.0002	.0020	.0010	-.0066	-.2487	-.0013	.8002	.5180**
	G	-.0378	.0043	.0195	.0077	-.0137	-.2601	-.0022	.8133	.5377**

Fig.1 Scree plot of 45 genotypes of linseed on principal components 1-9 (F1-F9)

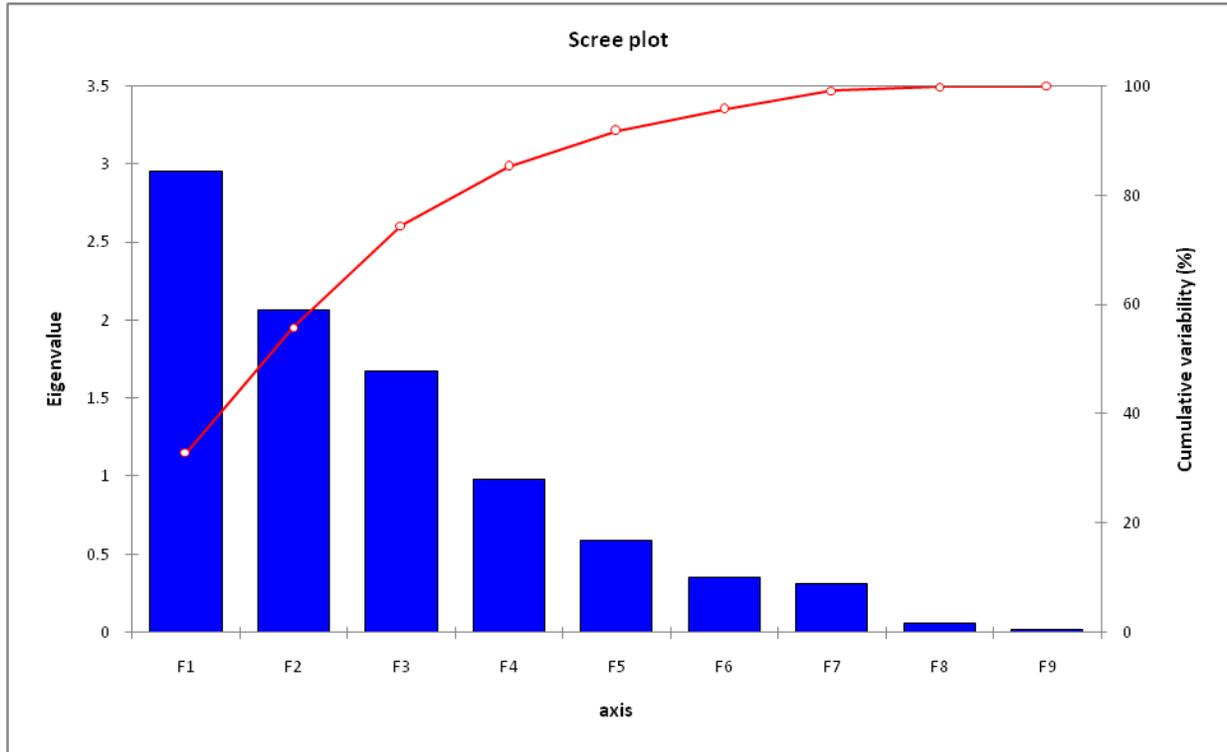
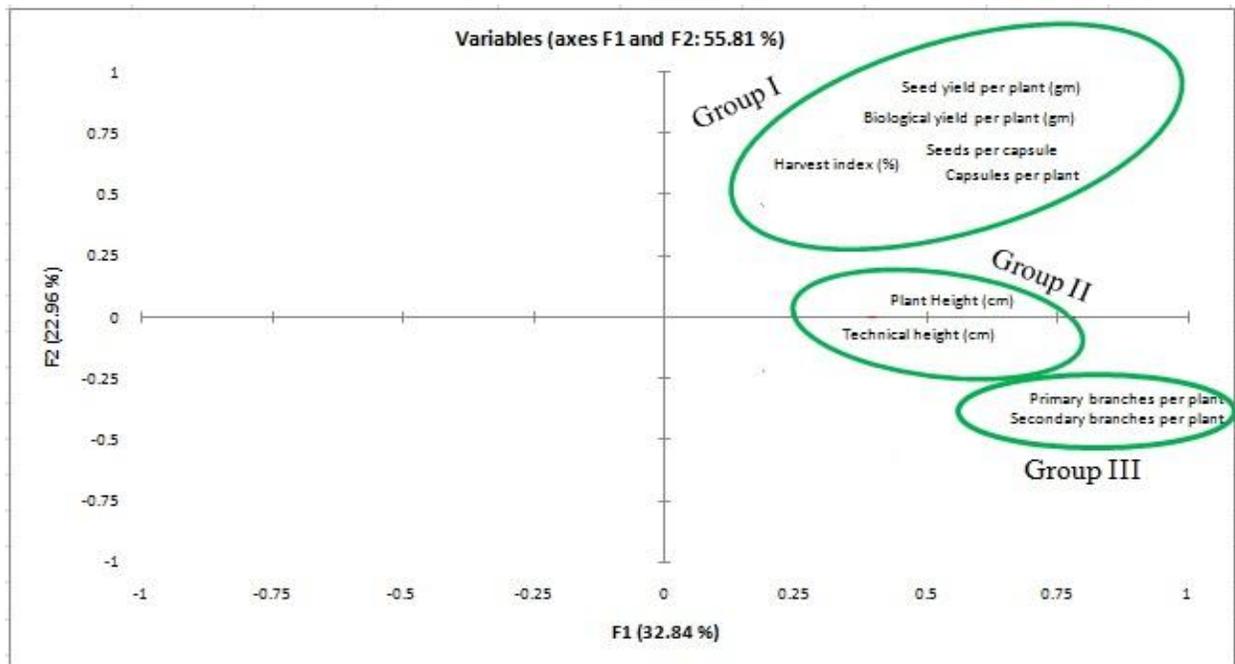


Fig.2 Biplot of 45 genotypes of linseed on Principal Component (F1 and F2) axis I and II



Path coefficient analysis permits the separation of the correlation coefficients into components of direct and indirect effects. Such information may be useful in predicting correlated responses of different characters towards directional selection. Keeping seed yield per plant as resultant variable and other traits as causal variables, the following results were obtained. The direct and indirect effects of genotypic path coefficients were mostly higher in magnitude than the corresponding phenotypic path coefficients (Table 3). Similar finding with respect to path coefficients have been reported by Gauraha and Rao (2011) and Reddy *et al.*, (2013). Harvest index exerted maximum direct effect on seed yield per plant followed by biological yield per plant and seeds per capsule while plant height and primary branches per plant showed negative direct effects on seed yield per plant at both (genotypic and phenotypic) levels. Also we see that harvest index exerts maximum indirect effect via plant height followed by technical height on seed yield per plant and biological yield per plant has maximum indirect effects via capsules per plant on seed yield per plant. Seed yield had maximum genotypic and phenotypic correlation with biological yield per plant followed by harvest index and seeds per capsule so direct selection of plants based on these three traits i.e., biological yield per plant, harvest index and seeds per capsule would be effective to increase seed yield.

As path coefficient analysis determines the effect of individual traits on overall yield, principal component was also performed to determine the performance of individual advance lines and their effect on different variables.

The principal component analysis (PCA) was performed for traits (Table 4) which revealed four most informative principal components with eigen values of 2.956, 2.067, 1.674 and

0.978 respectively, which together accounted 85.273% of the total variance (Fig. 1.) for all the traits viz., plant height, technical height, primary branches per plant, secondary branches per plant, capsules per plant, biological yield per plant, seeds per capsule, seed yield per plant and harvest index. As earlier reported by Kumar and Paul (2016) The three groups were found for the traits studied (Fig. 2). The group I had 5 traits in a group (seed yield per plant, biological yield per plant, harvest index, seeds per capsule and capsules per plant) and group II had 2 traits (plant height and technical height) and in group III (primary branches per plant and secondary branches per plant) had 2 traits.

The figure shows that traits within the group are closely associated like seed yield per plant, biological yield per plant, harvest index, seeds per capsule and capsules per plant all fall under the same group it shows the traits within the group are more associated. The results showed that group I traits could be used for selecting high yielding lines. In the present investigation it is observed that there were positive significant associations between seed yield per plant with plant height, secondary branches per plant, capsules per plant, biological yield per plant, seeds per capsule and harvest index. The positive associations of such traits indicated especially that such characters can be applied for direct and indirect selection in breeding programmes. Additionally, the PCA showed that the traits fall in three groups. Group I had 5 traits including seed yield per plant so group I traits could be used for selecting high yielding lines.

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