

## Genotypic Correlations and Path Coefficient Analysis in Blackgram *Vigna mungo* (L) Hepper\*\*

A. R. MUTHIAH<sup>1</sup> and V. SIVASUBRAMANIAN<sup>2</sup>

Yield (kg/ha)

Grain yield

SW NE

1685 1633

1440 1537

1154 1149

69 76

147 160

1282 1454

1574 1430

1524 1434

69 76

147 N.S.

1502 1456

1402 1433

1375 1431

24 11

49 22

Correlation Coefficients and path coefficients were computed for yield components in 50 types of blackgram. The results indicated that all the characters studied showed significant positive association with seed yield at genotypic and phenotypic levels. Pod number and cluster number not only showed high positive association among themselves but also showed positive correlation with other characters with the exception of hundred seed weight. Besides pod yield, pod number was the most important character in deciding the grain yield per plant. Pod length had a negative direct effect. The indirect effect of all the yield components via pod length were always negative. The residual effect was of a low magnitude.

A study was undertaken on 50 Indian types of blackgram to assess 1) the phenotypic, genotypic and environmental correlations between yield and its components as well as between different components themselves and 2) the direct and indirect influence of the various components on the yield.

### MATERIAL AND METHODS

A set of 50 types of blackgram of indigenous origin were selected for the present study. These types were raised in randomised blocks replicated three times at the experimental area of Tamil Nadu Agricultural University, Coimbatore. Each genotype was represented by a single row of three metre length. The rows were spaced 60 cm apart with intervals of 15 cm between the plants. Five plants were selected at random from each treatment in every replication

and observations were recorded for nine quantitative characters viz., plant height, number of branches, number of clusters per plant, number of pods per plant, length of pod, number of seeds per pod, hundred seed weight, pod yield per plant and seed yield per plant. The mean values of these five plants were used for the statistical analysis. The path coefficient analysis developed by Wright 1921 was applied to the data and the estimates were obtained by the method adopted by Dewey and Lu (1959).

### RESULTS AND DISCUSSION:

Genotypic, phenotypic and environmental correlations between the different pairs of characters were estimated and presented in Table. 1. All the characters studied showed significant positive association with seed yield at genotypic and phenotypic levels. The

\*\* Forms a Part of M. Sc. (Ag) thesis of the first author submitted to the Tamil Nadu Agri. University, Coimbatore-641 003.

1, 2 - Dept. Agri. Botany TNAU, Coimbatore-3.



enviromental correlation coefficients were in almost all cases lower than the genotypic and phenotypic correlations indicating that the traits selected for the study are less influenced by the environment. The high degree of positive correlation between cluster number and pod number as well as those between these characters and yield indicate the utility of these characters for the development of high yielding varieties. Similar significant postive association of pod number with seed yield was also observed by Verma and Dubey (1970) and Singh *et al.* (1972) in black gram and also in red gram by Joshi (1973). Significant positive correlation of cluster number with seed yield of blackgram was also indicated by Singh *et al.* (1975). Singh and Malhotra (1973) also reported that the number of clusters and pods showed significant positive association with seed yield of red gram. In the present study cluster number and pod number also showed positive genotypic correlations with other characters except 100 seed weight. However, Singh *et al.* (1975) in black gram reported that while cluster number was highly and positively associated with plant height and primary branches no association with hundred seed weight was observed.

The results of a path analysis of the data is presented in Table 2. The analysis indicated that other than the pot yield the pod number and seeds per pod had significant direct effect on seed yield with 44.65 per cent and 26.31 per cent contribution respectively. Singh *et al.* (1972) in blackgram also reported the direct positive effect of pod number.

Very high positive effect of pod number on seed yield of soybean was reported by Lal and Md. Fazulul Haque (1972). In the present study the high correlation of cluster number with seed yield was mainly due to the indirect effects via pod number (39.05%). A negative indirect effect of cluster number via pod number was however observed in Soybean by Sengupta and Kataria (1971).

The indirect influence of plant height and branches via pod number was also considerable. This type of relationship has been observed in Soybean by Malhotra *et al.* (1972). Considering the genetic correlations together the results of path analysis, pod number and seeds per pod appear to be the most important criteria determining yield. Pod length had a negative direct effect. The indirect effect of all the yield components viz pod length were always negative. The result was in accordance with the result of Chandel *et al.* (1973) in green gram.

In the present study, the residual effect was of low magnitude (0.2691) suggesting that most of the important contributing to seed yield have been utilised.

#### REFERENCES

- CHANDEL, K. P. S., B. S. JOSHI and K. C. PANT 1973. Yield of mung beans and its components *Indian J. Genet.*, 33: 271-76
- DEWEY, D. R. and K. H. LU. 1959. A correlation and path coefficient analysis of components of crested wheat grass seed production *Agron J.*, 51: 515-18
- JOSHI, S. N. 1973. Variability and Correlation studies in pigeon pea (*Cajanus cajan* L) *Madras Agric. J.* 60 412-14



February 1981]

# PATH ANALYSIS IN BLACKGRAM

LAL, V. S. and Md., FAZULUL HAQUE 1971. Path analysis of yield components in soybean. *Indian J. Genet.*, 31: 357-62

MALHOTRA, R. S., K. B. SINGH and H. S. DHALIWAL. 1972. Correlation and path coefficient analysis in Soybean (*Glycine max* (L) Merr.) *Indian J. Agril. Sci.* 42: 26-29

SENGUPTA K and K. S. KATARIA, 1971. Path coefficient analysis for some characters in Soybean *Indian J. Genet.* 31: 290-95

SINGH, K. B. and R. S. MALHOTRA 1973. Yield components in Pigeon pea *Madras Agric J.* 60: 364-66.

SINGH, U. P., U. SINGH and P. SINGH. 1975. Estimates of variability heritability and correlation for yield and its components in *Urd*, *Madras Agric. J.* 62: 71-72

SINGH, K. B. G. S. BHULLAR, R. S. MALHOTRA and J. K. SINGH 1972. Estimates of genetic variability, correlation and path coefficients in *Urd* and their importance in selection *J. Res. PAU*, 9: 410-16

VERMA, S. N. P. and C. S. DUBEY 1970. Correlation studies in blackgram (*Phaseolus mungo* L.) *Allahabad Farmer* 44: 419.

WRIGHT S. 1921. Correlation and causation. *J. Agric. Res.* 20: 557-85



TABLE-1 Genotypic, Phenotypic and environmental correlations between different pairs of characters

	Plant height	Branches	Cluster number	Pod length	Seeds per pod	100 seed weight	Pod yield	Seed yield
Plant height		+0.5973** +0.5978** +0.1960**	+0.7937** +0.7685** +0.2652**	+0.7302** +0.6827** +0.0489	+0.7056** +0.6262** +0.0028	+0.4450** +0.4270** -0.0048	+0.8241** +0.8019** +0.2246**	+0.7899** +0.7674** +0.2229**
Branches			+0.7435** +0.7154** +0.2447**	+0.5602** +0.5050** -0.0297	+0.4459** +0.3852** -0.0646	+0.3956** +0.3753** -0.0142	+0.6757** +0.6641** +0.2137**	+0.6685** +0.6461** +0.2130**
Cluster number				+0.5627** +0.5262** +0.1208**	+0.5546** +0.4907** +0.0515	+0.2792** +0.2667** +0.0566	+0.8758** +0.8699** +0.7731**	+0.8505** +0.8434** +0.7738**
				+0.3367** +0.3185** +0.0841*	+0.3677** +0.3296** +0.0430	+0.0749 +0.0721 -0.0035	+0.8776** +0.8782** +0.8976**	+0.8981** +0.8967** +0.8916**
Pod length					+0.9213** +0.9002** +0.8160**	+0.6446** +0.5967** +0.0092**	+0.5993** +0.5554** +0.0809*	+0.5581** +0.5208** +0.0814*
Seeds per pod						+0.4585** +0.4030** -0.0100	+0.5989** +0.5316** +0.0259	+0.5717** +0.5035** +0.0287
100 seed weight							+0.0259** +0.2754** +0.0276	+0.02831** +0.2700** +0.0122
Pod yield								+0.9511** +0.9499** +0.9626**

Note: \*\* Significant at 1% level ( $P=0.01$ )  
 \* Significant at 5% level ( $P=0.05$ )



February 1980]

PATH ANALYSIS IN BLACKGRAM

Note: \*\* Significant at 1% level (P=0.01)  
\* Significant at 5% level (P=0.05)

TABLE 2 Path analysis of correlation coefficients

	Direct effect	Indirect effect via variables					Total genotypic			
		Plant height	Branches	Cluster number	Pod number	Pod length	Seeds per pod	100 seed weight	Pod yield	correlation coefficient
Seed yield and plant height	0.0957 (12.11)	--	0.0513 (6.49)	-0.0978 (-12.38)	0.2553 (32.32)	-0.0975 (-12.34)	0.1061 (13.43)	0.0326 (4.13)	0.4442 (56.23)	0.7899
Seed yield and branches	0.0859 (12.85)	0.0572 (8.56)	--	-0.0916 (-13.70)	0.2302 (34.43)	-0.0735 (-10.99)	0.0670 (10.02)	0.0290 (4.34)	0.3643 (54.49)	0.6685
Seed yield end cluster number	-0.1232 (-14.49)	0.0759 (9.92)	0.0638 (7.50)	--	0.3321 (39.05)	-0.0751 (-8.83)	0.0834 (9.81)	0.0205 (2.41)	0.4731 (55.63)	0.8505
Seed yield and pod number	0.4010 (44.65)	0.0609 (6.78)	0.0493 (5.49)	-0.1020 (-11.36)	--	-0.0450 (-5.01)	0.0553 (6.16)	0.0055 (0.61)	0.4731 (52.68)	0.8981
Seed yield and pod length	-0.1336 (-23.94)	0.0699 (12.52)	0.0473 (8.47)	-0.0683 (12.42)	0.1350 (24.19)	--	0.1385 (24.82)	0.0472 (8.46)	0.3231 (57.89)	0.5581
Seed yield and seeds per pod	0.1504 (26.31)	0.0675 (11.81)	0.383 (6.70)	-0.0683 (-11.95)	0.1474 (25.78)	-0.1230 (-21.51)	--	0.0366 (6.40)	0.3228 (56.46)	0.5717
Seed Yield and 100 seed weight	0.0733 (25.89)	0.0426 (15.05)	0.0340 (12.01)	-0.0344 (-12.15)	0.0302 (10.67)	-0.0861 (-30.41)	0.0689 (24.34)	--	0.5546 (54.61)	0.2831
Seed yield and pod yield	0.5391 (56.68)	0.0789 (8.30)	0.0580 (6.10)	-0.1079 (-11.34)	0.3520 (37.01)	-0.0800 (-8.41)	0.0900 (9.46)	0.0210 (2.21)	--	0.9511

Residual value = 0.2691

Note : Figures in parantheses denote percentages

v. r.