

PATH COEFFICIENT ANALYSIS STUDIES ON SEED YIELD AND SEED YIELD CONTRIBUTING CHARACTERS IN GREEN GRAM [*Vigna radiata* (L.) R. Wilczek]

Abstract

The current empirical study on "Path coefficient analysis studies on seed yield and seed yield contributing characters in green gram (*Vigna radiata* (L.) R. Wilczek)" was conducted to assess path analysis in 72 genotypes of green gram. On 11 traits, observations were recorded. The genotypic path coefficient analysis revealed that number of pods per plant, 100-seeds weight and number of seeds per pod expressed positive and higher direct effect on seed yield per plant. The phenotypic path coefficient analysis revealed that number of pods per plant, 100-seeds weight and number of seeds per pod exhibited high and positive direct effects on seed yield per plant. Whereas, negative direct effect on seed yield per plant were contributed through number of clusters per plant and plant height at phenotypic level. The residual effect was found to be 0.0755 and 0.1321 at genotypic and phenotypic path coefficient analysis respectively.

Keywords: Path analysis, Green gram, Residual effect and *Vigna radiata* (L.) R. Wilczek.

1. Introduction

In Indian agriculture, pulse crops play an important role. Quality protein is higher in pulses nearly three times as compared to cereals. Green gram is a legume cultivated for its edible seeds and sprouts across Asia. It has a diploid chromosome number of $2n=2x=22$ and is a member of the fabaceae family and subfamily papilionaceae. Green gram is a native of Central Asia and India, claims Vavilov (1939). Green gram contains about 24 % protein, this is being about 2/3 of the protein content of soybean, twice that of wheat and thrice that of rice. This protein is relatively high in an amino acid and lysine that is lacking in cereal grains. So, green gram and cereal grains diet combining form a balanced amino acid diet. Every 100 g green gram seed contain 132 mg calcium, 2.251 mg niacin, 4.8 mg ascorbic acid, 0.621 mg thiamine, 0.233 mg riboflavin and 114 IU vitamin A (Haytowitz and Matthews, 1986).

In exercising selection programs, the information on path analysis is of

considerable importance. Multiplicative end product of many factors is being a polygenic complex trait and sensitive to environmental variations in grain yield. There are also several component characters of yield. It is necessary to examine whether the effects of different components of yield and other traits on yield are direct or indirect and to what extent? The path coefficient analysis provides an effective mean of finding direct and indirect causes of an association under such circumstances.

2. Materials and methods

72 genotypes of green gram from different origins obtained from the Pulses Research Station, J.A.U., Junagadh were sown in a RBD with 3 replications during *kharif-2021* at Pulses Research Station, J.A.U., Junagadh. Each genotype was arranged in a single row that was 4 metres long with a spacing of 45 cm × 10 cm.

The observations were recorded for 11 traits *viz.*, days to 50 % flowering, days to maturity, plant height (cm), number of primary branches per plant, number of clusters per plant, number of pods per cluster, number of pods per plant, number of seeds per pod, length of pod (cm), 100-seeds weight (g) and grain yield per plant (g) and mean values were used for statistical analysis.

Path coefficient is a standardized partial regression coefficient and measures the direct and indirect effects of one variable upon another and permits the separation of correlation coefficient into the component of direct and indirect effects. The genotypic as well as phenotypic path coefficient analysis was done as per the method suggested by Dewey and Lu (1959).

3. Result and discussion

The path coefficient analysis was done for yield and yield attributes to estimate the direct and indirect effects of various characters on seed yield. Path coefficient analysis considers direct as well as indirect effects of the variables by partitioning the correlation coefficients.

In order to understand these effects, genotypic and phenotypic correlation coefficient of different characters with seed yield was portioned into their direct and indirect effects (Table 1). This facilitates the selection of genotypes on the basis of those traits which will eventually contribute more towards seeds yield.

The genotypic path coefficient analysis (Table 1) revealed that number of pods per plant, 100-seeds weight and number of seeds per pod expressed positive and higher direct effect on seed yield per plant. These results were in conformity with earlier report of Ahmad *et al.* (2013). Goyal *et al.* (2021) for 100-seeds weight and number of pods per plant.

While, negative direct effect on seed yield per plant were contributed through number of clusters per plant, days to 50 % flowering and plant height at genotypic level, indicating that the selection for these traits may have an undesirable impact on seed yield. These results are in accordance with the reports of Prasanna *et al.* (2013) and Mohammed *et al.* (2020) for number of clusters per plant and days to 50 % flowering. Garg *et al.* (2017), Abbas *et al.* (2018) and Asari *et al.* (2019) for plant height.

The phenotypic path coefficient analysis (Table 1) revealed that highest positive direct effect on seed yield were exerted through number of pods per plant followed by 100-seeds weight and number of seeds per pod. So, emphasis should be given to these traits in selection program for improvement of seed yield in green gram. Which coincides earlier reported by Parihar *et al.* (2018) for 100-seeds weight and number of seeds per pod; Ahmad *et al.* (2013) and Prasanna *et al.* (2013) for number of seeds per pod, 100-seeds weight and number of pods per plant; Tabasum *et al.* (2010) for number of pods per plant and 100-seed weight.

However, negative direct effect on seed yield per plant were contributed through number of clusters per plant and plant height at phenotypic level, indicating that the selection for these traits may have an undesirable impact on seed yield. Similar outcome were also reported by Abhisheka and Mogali (2020) and Mohammed *etal.* (2020) for number of clusters per plant; Goyal *etal.* (2021) and Abbas *etal.* (2018) for plant height.

It was apparent from the phenotypic path analysis that higher direct effects as well as appreciable indirect influences were exerted by number of pods per plant, 100-seeds weight and number of seeds per pod towards seed yield per plant. These three characters also exhibited highly significant and positive association with seed yield per plant. Hence, these may be considered as most important yield contributing

characters and due emphasis should be placed on this component while breeding for high seed yield in green gram.

The residual effect was found to be 0.0755 at genotypic path coefficient analysis, while it was 0.1321 at phenotypic path coefficient analysis. The results of residual effect indicated that the majority of the yield attributes have been included in the study of path analysis. It can also be concluded that the characters which are most important for correlation studies are also important for path analysis. Thus, it can be suggested that correlation and path analysis study should be considered together for rapid gain of final genetic improvement for seed yield in green gram. These results are in accordance with the reports of Manivelan *et al.* (2019), Kumar *et al.* (2018) and Garg *et al.* (2017).

4. Conclusion

From the present investigation, it can be suggested that number of pods per plant, 100-seeds weight and number of seeds per pod expressed positive and higher direct effect on seed yield per plant at genotypic path analysis. The phenotypic path analysis revealed that number of pods per plant, 100-seeds weight and number of seeds per pod had high and positive direct effect on seed yield per plant and that was found to be the most important yield components. The residual

effect was of low magnitude suggesting that the majority of the yield attributes have been included in the study of path analysis.

Table 1: Genotypic and Phenotypic path coefficient analysis showing direct (diagonal and bold) and indirect effects of different characters on seed yield perplant in 72 genotypes of green gram

Characters		Days to 50% flowering	Days to maturity	Plant height (cm)	Number of primary branches per plant	Number of clusters per plant	Number of pods per cluster	Number of pods per plant	Number of seeds per pod	Length of pod (cm)	100- seeds weight (g)
Days to 50% flowering	G	-0.0380	-0.0339	-0.0069	0.0006	0.0173	0.0215	0.0191	0.0065	0.0023	-0.0110
	P	0.0018	0.0011	0.0003	0.0001	-0.0006	-0.0008	-0.0007	-0.0002	-0.0001	0.0004
Days to maturity	G	0.0674	0.0754	0.0280	0.0078	-0.0507	-0.0537	-0.0629	0.0198	0.0060	0.0406
	P	0.0006	0.0010	0.0003	0.0001	-0.0005	-0.0006	-0.0006	0.0002	0.0001	0.0004
Plant height (cm)	G	-0.0040	-0.0082	-0.0221	-0.0042	0.0020	0.0025	0.0028	-0.0067	-0.0074	-0.0036
	P	-0.0014	-0.0026	-0.0095	-0.0015	0.0008	0.0010	0.0012	-0.0024	-0.0027	-0.0015
Number of primary branches per plant	G	-0.0001	0.0005	0.0010	0.0053	0.0017	0.0012	0.0016	0.0006	0.0009	0.0011
	P	0.0005	0.0009	0.0023	0.0147	0.0045	0.0030	0.0038	0.0015	0.0020	0.0029
Number of clusters per plant	G	0.0329	0.0486	0.0067	-0.0233	-0.0723	-0.0477	-0.0591	0.0100	-0.0097	0.0183
	P	0.0184	0.0280	0.0046	-0.0166	-0.0544	-0.0336	-0.0408	0.0066	-0.0055	0.0126
Number of pods per cluster	G	-0.0436	-0.0548	0.0085	0.0168	0.0508	0.0770	0.0622	-0.0016	0.0049	-0.0224
	P	-0.0300	-0.0382	-0.0074	0.0140	0.0429	0.0696	0.0513	-0.0015	0.0042	-0.0190
Number of pods per plant	G	-0.4492	-0.7468	-0.1136	0.2720	0.7320	0.7234	0.8948	-0.0834	0.0732	-0.3031
	P	-0.3284	-0.4914	-0.1014	0.2161	0.6216	0.6114	0.8290	-0.0696	0.0693	-0.2649
Number of seeds per pod	G	-0.0754	0.1157	0.1330	0.0520	-0.0610	-0.0089	-0.0410	0.4402	0.1501	0.0510
	P	-0.0418	0.0843	0.1246	0.0488	-0.0584	-0.0103	-0.0406	0.4837	0.1419	0.0480
Length of pod (cm)	G	-0.0028	0.0037	0.0155	0.0076	0.0062	0.0029	0.0038	0.0157	0.0460	-0.0012
	P	-0.0011	0.0027	0.0098	0.0046	0.0034	0.0021	0.0029	0.0101	0.0343	-0.0005
100- seeds weight (g)	G	0.1787	0.3324	0.1017	0.1238	-0.1565	-0.1793	-0.2091	0.0715	-0.0158	0.6174
	P	0.1327	0.2226	0.0931	0.1146	-0.1352	-0.1596	-0.1867	0.0579	-0.0083	0.5842
Genotypic correlation with Seed yield per plant (g)	G	-0.334**	-0.267*	0.135	0.458**	0.469**	0.539**	0.612**	0.473**	0.250*	0.387**
Phenotypic correlation with Seed yield per plant (g)	P	-0.249*	-0.192	0.117	0.395**	0.424**	0.482**	0.619**	0.486**	0.235*	0.363**

*, ** Significant at 5 and 1 % levels, respectively

Genotypic residual effect = 0.0755 Phenotypic residual effect = 0.1321

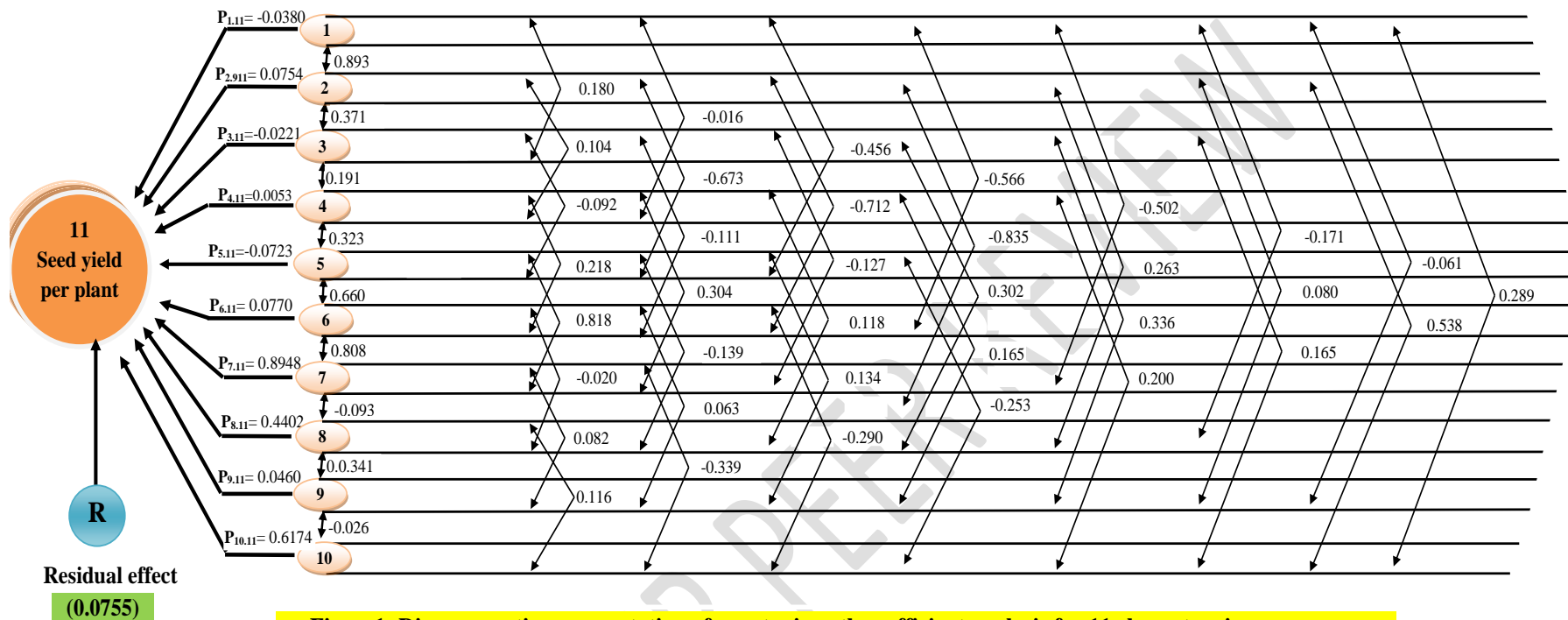


Figure1: Diagrammatic representation of genotypic path coefficient analysis for 11 characters in green gram

Characters

- | | |
|-----------------------------------------|------------------------------|
| 1. Days to 50% flowering | 7. Number of pods per plant |
| 2. Days to maturity | 8. Number of seeds per pod |
| 3. Plant height (cm) | 9. Length of pod (cm) |
| 4. Number of primary branches per plant | 10. 100-seeds weight (g) |
| 5. Number of clusters per plant | 11. Seed yield per plant (g) |

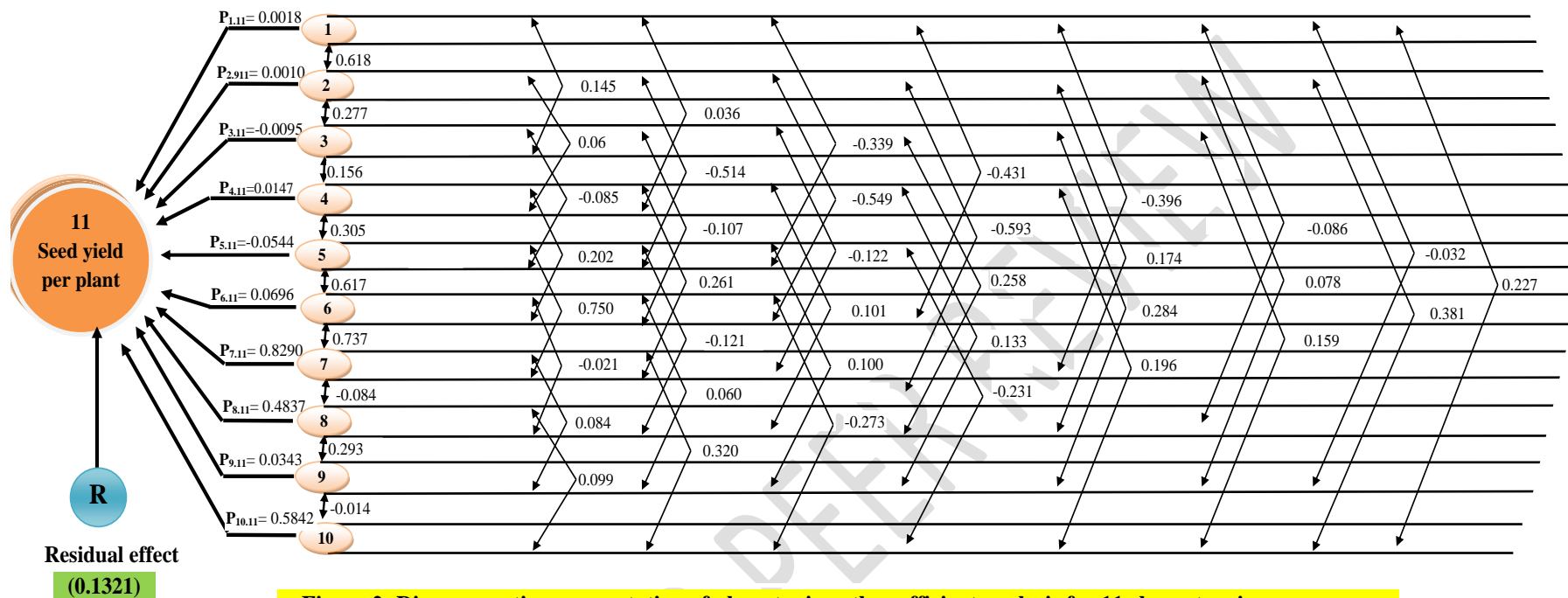


Figure 2: Diagrammatic representation of phenotypic path coefficient analysis for 11 characters in green gram

- | | |
|------------------------------------------------|-------------------------------------|
| 1. Days to 50% flowering | 7. Number of pods per plant |
| 2. Days to maturity | 8. Number of seeds per pod |
| 3. Plant height (cm) | 9. Length of pod (cm) |
| 4. Number of primary branches per plant | 10. 100-seeds weight (g) |
| 5. Number of clusters per plant | 11. Seed yield per plant (g) |

Characters

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