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Relationship among the yield contributing characters and with seed yield in pigeonpea (Cajanus cajan (L.) millsp.)

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Abstract

Twenty CGMS-based pigeonpea [*Cajanus cajan* (L.) Millsp.] hybrids were evaluated along with parents and standard check (Maruthi) to study the nature and magnitude of relationship of important agronomic traits with seed yield. The association studies indicated significant positive correlation of seed yield with all characters in parents and crosses except 100-seed weight (g), seed protein (%).

Keywords: correlation coefficient, yield, pigeonpea

Introduction

Pigeonpea [*Cajanus cajan* (L.) Millspaugh] is an important pulse crop of rainfed and semi-arid regions of Asia, Africa and the Caribbean islands. India accounts for over 85% of the global area of 4.6 million hectares. Globally, it is grown under low input environments, primarily as an intercrop with early maturing cereals. Pigeonpea is a major protein supplement for small holding farming families. Pigeonpea is also known for improving soil nutrition by fixing atmospheric nitrogen, releasing soil-bound phosphorus and recycling micro-nutrients. Its extensive root mass and leaf fall are known for improving soil structure and water infiltration in the soil.

Material and Methods

The present study involved five lines, four testers and their 20 F_1 's of pigeonpea obtained by L x T mating design (Kempthorne 1957)^[1] along with a standard check (Maruthi) at ICRISAT, Hyderabad, Telangana during *kharif*, 2016. Crop sown on 25th July, 2016 in a randomized block design with three replications. Each entry was grown in four rows of 4m length in each plot. The spacing adopted between rows to row was 75 cm and 30 cm between plant to plant within a row. Five plants were tagged randomly for recording the observations on 11 characters viz., days to 50% flowering, days to 75% maturity, plant height, primary branches per plant, secondary branches per plant, pods per plant, seeds per pod, 100-seed weight, harvest index, seed protein and seed yield per plant. Genotypic and phenotypic correlation coefficients were calculated using the method given by Jhonson (1955) ^[3].

Results and Discussion

Phenotypic and genotypic correlation coefficients between yield and its components in parents and crosses were presented in Table 1 and Table 2 respectively. The present investigation that highly significant and positive association of pods per plant, days to 50% flowering, days to 75% maturity, plant height, primary branches per plant, pods per plant, seeds per pod, harvest index with seed yield per plant in parents and crosses. Similar results were reported by Asawa *et al.* (1981) ^[5], Natarajan *et al.* (1990) ^[2], Paul and Upadhaya (1991) ^[7], Dhameliya and Pathak (1994) ^[7], Deshmukh *et al.* (2000) ^[9] and Baskaran and Mutiah (2007) ^[6].

Seed protein had non-significant and positive association with seed yield in parents and crosses (Devi *et al.* 2012) ^[4]. The positive and non-significant association of secondary branches per plant with seed yield in parents became positive and significant in crosses.

Non-significant and positive associations observed among the yield components in parents *viz.*, were secondary branches per plant with pods per plant become significant and positive in

Crosses. The positive and significant association in parents between character pairs viz., plant height with primary branches per plant; days to 75% maturity with pods per plant and seeds per pod; days to 75% maturity, plant height, primary branches per plant, seeds per pod with harvest index; seeds per pod with pods per plant, primary branches per plant and seed protein with secondary branches per plant became non-significant in crosses.

Negative and significant association of among character pairs in parents became negative and significant in crosses *viz.*, 100-seed weight with plant height and harvest index. Negative non-significant association of 100-seed weight with primary branches per plant in parents became negative and significant in crosses.

Positive and non-significant association of harvest index with seed protein and seed protein with primary branches per plant in parents became negative and non-significant in crosses. Negative and non-significant association of 100-seed weight with days to 75% maturity in parents became Positive and non-significant in crosses. These changes in correlations between pairs of characters in parents and crosses might have resulted due to the modifications in the linkages as a result of re-combinations in the populations.

In general, the yield component traits which showed significant and positive association with seed yield in parents and crosses are of prime importance in determining the seed yield. Such a situation is favourable to a plant breeder as it helps in simultaneous improvement of the characters along with seed yield per se. Hence, the emphasis should be given on days to 50% flowering, days to 75% maturity, plant height, primary branches per plant, pods per plant, seeds per pod and harvest index for obtaining higher yield in pigeonpea.

Table 1: Correlation coefficients for yield and yield components in parents of pigeonpea

		Days 50%	Days to	Plant	Primary	Secondary	Number of	Number	100-seed	Harvest	Seed	Seed yield
Characters		to	75%	height	branches	branches	pods per	of seeds	weight	index	protein	per plant
		flowering	maturity	(cm)	per plant	per plant	plant	per pod	(g)	(%)	(%)	(g)
Days 50% to	Р	1.0000	0.913**	0.788^{**}	0.729**	0.425*	0.832**	0.602**	-0.541**	0.892**	0.268	0.798**
flowering	G	1.0000	0.936**	0.803**	0.844**	0.439*	0.844**	0.682**	-0.689**	0.910**	0.418*	0.811**
Days to 75%	Р		1.0000	0.759**	0.675**	0.363	0.843**	0.461*	-0.313	0.822**	0.140	0.845**
maturity	G		1.0000	0.755**	0.805**	0.377*	0.858**	0.527**	-0.429*	0.839**	0.258	0.859**
Dlant haight (am)	Р			1.0000	0.682**	0.681**	0.700**	0.656**	-0.622**	0.792**	0.312	0.684**
r faitt fiergift (cfif)	G			1.0000	0.801**	0.695**	0.702**	0.705**	-0.804**	0.799**	0.424*	0.685**
Primary branches	Р				1.0000	0.653**	0.496**	0.509**	-0.331	0.462*	0.354	0.494**
per plant	G				1.0000	0.767**	0.585**	0.712**	-0.486**	0.567**	0.445*	0.587**
Secondary	Р					1.0000	0.134	0.376*	-0.369*	0.303	0.379*	0.122
branches per plant	G					1.0000	0.136	0.406*	-0.428*	0.315	0.619**	0.126
Number of pods	Р						1.0000	0.477**	-0.456*	0.857**	0.206	0.987**
per plant	G						1.0000	0.511**	-0.584**	0.861**	0.301	0.987**
Number of seeds	Р							1.0000	-0.451*	0.551**	-0.015	0.428*
per pod	G							1.0000	-0.735**	0.576**	-0.185	0.455*
100-seed weight	Р								1.0000	-0.608**	-0.324	-0.402*
(g)	G								1.0000	-0.809**	-	-0.520**
	D									1.0000	$0./1/^{**}$	0.700**
Harvest index (%)	P C									1.0000	0.260	0.799^{**}
	U D									1.0000	0.384*	0.801***
Seed protein (%)	P C										1.0000	0.197
	G										1.0000	0.283
Seed yield per	P											1.0000
plant (g)	G											1.0000

** Significant at 0.01; * Significant at 0.05

Table 2: Correlation coefficients for yield and yield components in crosses of pigeonpea

		Days 50%	Days to	Plant	Primary	Secondary	Number of	Number	100-seed	Harvest	Seed	Seed yield
Characters		to	75%	height	branches	branches	pods per	of seeds	woight (g)	indox (%)	protein	per plant
		flowering	maturity	(cm)	per plant	per plant	plant	per pod	weight (g)	muex (70)	(%)	(g)
Days 50% to	Р	1.0000	0.417**	0.436**	0.508**	0.397**	0.448 * *	0.486**	-0.360**	0.369**	-0.070	0.543**
flowering	G	1.0000	0.484**	0.473**	0.653**	0.422**	0.467**	0.563**	-0.491**	0.391**	-0.076	0.569**
Days to 75%	Р		1.0000	0.401**	0.279*	0.072	0.153	0.073	0.002	0.170	0.072	0.264*
maturity	G		1.0000	0.438**	0.396**	0.097	0.165	0.089	-0.048	0.190	0.005	0.282*
Plant height (cm)	Р			1.0000	0.228	0.335**	0.450**	0.434**	-0.252	0.234	0.107	0.521**
	G			1.0000	0.250	0.343**	0.457**	0.456**	-0.358**	0.238	0.180	0.528**
Primary branches	Р				1.0000	0.597**	0.419**	0.172	-0.475**	0.240	-0.060	0.430**
per plant	G				1.0000	0.677**	0.503**	0.210	-0.739**	0.280*	-0.002	0.519**
Secondary branches	Р					1.0000	0.627**	0.261*	-0.605**	0.181	0.241	0.608**
per plant	G					1.0000	0.645**	0.286*	-0.766**	0.188	0.537**	0.627**
Number of pods per	Ρ						1.0000	0.193	-0.509**	0.491**	0.241	0.933**
plant	G						1.0000	0.205	-0.682**	0.495**	0.461**	0.933**
Number of seeds	Р							1.0000	-0.255*	0.194	-0.075	0.274*
per pod	G							1.0000	-0.367**	0.209	-0.101	0.294*
100-seed weight (g)	Р								1.0000	-0.194	-0.037	-0.524**
	G								1.0000	-0.257*	-0.553**	-0.702**
Harvest index (%)	Р									1.0000	-0.147	0.710**
	G									1.0000	-0.303*	0.714**

Seed protein (%) P G				1.0000 1.0000	0.163 0.299*
Seed yield per plant P					1.0000
(g) G					1.0000

** Significant at 0.01; * Significant at 0.05

Conclusion

Character association studies revealed that selection based on all the characters except secondary branches per plant, seed protein, 100-seed weight in parents and seed protein, 100-seed weight in hybrids will be effective for bringing improvement in seed yield as they had significant and positive correlation with seed yield per plant.

References

- 1. Kemp thorne O. An introduction to genetic statistics. John Wiley and Sons, Inc., New York. 1957; 222-240.
- Natarajan C, Thyagarajan K and Ayymperumal A. Genetic variability, correlation and path analysis in pigeonpea. Madras Agricultural Journal. 1990; 77: 378-381.
- 3. Johnson HW, Robinson HF and Comstock RE. Estimates of genetic and environmental variability in soybean. Agric. J. 1955; 47:314-318.
- 4. Devi SR, Prasanthi L, Reddy KHP Reddy BVB. Studies on interrelationships of yield and its attributes and path analysis in pigeonpea (*Cajanus cajan* (L.) Millsp.). Legume Research. 2012; 35(3):207-213.
- 5. Asawa BM, Chandra RK and Pandey RL. Character correlations and divergence in pigeonpea. Indian Journal of Agricultural Sciences. 1981; 51:12-17.
- 6. Baskaran K and Muthiah AR. Associations between yield and yield attributes in pigeonpea (*Cajanus cajan* (L.) Millsp). Legume Research. 2007; 30(1):64-66.
- 7. Paul SK and Upadhaya LP. International between yield and yield contributing characters in pigeonpea (*Cajanus cajan* (L.) Millsp). International Journal of Tropical Agriculture. 1991; (2):135-140.
- 8. Dhameliya HR and Pathak AR and Zaveri PP. Genetic analysis of heterogeneous population in pigeonpea (*Cajanus cajan* (L.) Millsp). Gujarat Agricultural University Research Journal. 1994; 20(1):46-51.
- Deshmukh RB, Rodge RG, Patil JV Sahane DV. Genetic variability and character association in pigeonpea under different cropping systems. Journal of Maharashtra Agricultural Universities. 2000; 25(2):176-178.