## GLOBAL JOURNAL OF BIO-SCIENCE AND BIOTECHNOLOGY

© 2004 - 2017 Society For Science and Nature (SFSN). All rights reserved

www.scienceandnature.org

# VARIABILITY STUDIES IN EARLY GENOTYPES OF PIGEON PEA (Cajanus cajan (L.) Millsp.)

C.P. Bal, S.G. Bhave, B.L. Thaware, S.S, Desai<sup>\*</sup>, S.G. Mahadik and Krishnaja R. Nair Department of Agricultural Botany, College of Agriculture, DBSKKV, Dapoli \*Corresponding author email: <u>desaiss2008@rediffmail.com</u>

## ABSTRACT

The study of general mean, range, genotypic, phenotypic and environmental variance, genotypic, phenotypic and environmental coefficient of variance, heritability and genetic advance were conducted for grain yield and its components among parents, 34 genotypes of the pigeon pea. All the characters studied exhibited significant variability among all population. Higher estimates of genotypic and phenotypic coefficients of variation indicating high level of variability and ample scope for effective improvement. The higher estimates of heritability coupled with high genetic advance as per cent of mean indicated additive gene action for the characters considered.

**KEYWORDS:** pigeon pea, variability, heritability, genetic advance.

### INTRODUCTION

Pigeon pea is also known as 'arhar, tur or redgram' with chromosome number i.e. 2n=2x=22, belongs to family 'leguminosae' act both as a food crop (dried peas, flour or green vegetable peas) and a forage or cover crop. Pigeon pea is important legume crop of rainfed agriculture in the semiarid tropics. Pigeon pea are cultivated in more than 25 tropical and subtropical countries either as sole crop or intermixed with cereals. It's benefits are protein rich seed (21% protein), fuel, fodder and erosion control. It is largely consumed in the form of split pulse as 'dal' while its tender green pods constitute a very favourite vegetable in some parts. The stalks are utilized for various purposes, such as roofing, walling, sides of carts and basket making and burning as fuel. The trait of grain yield is controlled by complex gene action and hence traits contributing to yield must be considered and evaluated. To retrieve culture yielding superiorly with better nutrient value, utilization of diverse parents for introgression is essential. It is therefore, necessary to estimate relative amounts of genetic and non-genetic variability exhibited by different characters using suitable parameters like genotypic coefficient of variability (GCV), Phenotypic coefficient of variability (PCV), heritability estimates (h2) and genetic advance (GA) and genetic association between yield and vield contributing, the information on their aspects will help in the breeders to determine the selection criteria for isolating high yielder genotypes in pigeonpea.

### **MATERIALS & METHODS**

The present study comprised a set of selected 34genotypes. These were sown during Rabi 2014-15 in Randomized Block Design, with a spacing of about 30 x 20 cm in three replication and standard agronomic practices were followed. Five plants selected at random were tagged from each genotype and observations on thirteen quantitative characters (plant height, days to

initiation of flowering, days to 50% flowering, days to maturity, number of primary branches per plant, number of pods per plant, pod length, pod breadth, number of seeds per pod, hundred seed weight, grain yield per plant, straw yield per plant and harvest index) were recorded on these plants.

Genetic variability parameter viz., mean, variance, phenotypic coefficient of variation (PCV) and genotypic coefficient of variation (GCV) (Burton and De Vane, 1953), heritability  $(h^2)$  and Genetic advance (GA) (Johnson *et al.*, 1955) among characters were calculated by following the standard procedures with the help of INDOSTST software's.

### **RESULTS & DISCUSSION**

Wide range of variation was observed among the genotypes for all the characters studied. The characters showing significant mean sum of squares for all the characters indicated the extent of variability existed in the population (table 1). The character plant height and days to initiation of flowering showed high range of variation. Genotype ICPH 2671 (70.43 cm) was the tallest and genotype ICPL 20340 (61.47 cm) was the dwarfest one, whereas genotype ICPL 11249 (57 days) and ICPL 11263 (57 days) were earliest in initiation of flowering. Chandra et al. (1975). Dodake et al. (2009). Vijavalakshmi et al. (2013) and Sharma et al. (2014) also reported wide range of variation for these characters in pigeon pea. Wide genetic variability for plant height was also reported by Aher et al. (1998). Another important character having direct bearing on yield are number of primary branches per plant and number of pods per plant. More the branches more will be the number of pods and ultimately more will be the yield. This result is in conformation with Rao and Rao (2015) and Venketeswarlu (2001) observed maximum variability for number of pods per plant.

		vest index	nd X <sub>13</sub> =Harv	er plant(g) ai	raw yield p	$t(g), X_{12} = st$	ld per plan	$\zeta_{11} = \text{grain yie}$	ight >	fred seed we	pod, $X_{10}$ = hunc	of seeds per	X <sub>9</sub> = number	readth,	h(cm), X <sub>8</sub> = pod b	ngt
$X_{7}=$	ls per plant,	number of pod	lant, $X_6 = 1$	anches per p	primary bra	number of	urity, $X_5 =$	days to matu	ering, X <sub>4</sub> =	to 50% flow	$x_3 = days$	n of flowerir	/s to initiatio	$\zeta_2 = day$	lant height(cm), λ	ĨL.
ı	315.951	173.980	64.910	23.722	17.163	1.186	16.509	417.328	1.745	806.020	790.314	1306.804	58.897	66	Error	
	2592.834*	1294.350*	846.118*	334.148*	32.006*	65.162*	28.110*	60779.21*	139.28*	7214.627*	8893.333*	9695.578*	702.909*	33	Genotypes	
	88.376	1350.580	12.650	1.998	1.602	4.247	3.526	3534.512	31.809	64.647	57.020	152.529	734.030	2	Replication	
1	<b>∧</b> 13	<b>∧</b> 12	<b>V</b> <sup>11</sup>	<b>∧</b> 10	<b>v</b> 9	v <sup>8</sup>	7	$\mathbf{v}^{6}$	$\Lambda_5$	24	$\Lambda_3$	$\mathbf{A}_2$	2	£	variation	
	V	V	V	V	v	V	v	v	v	V	V	V	V	Чf	Source of	
1					aracters	ititative cha	teen quar	iance for thi	ysis of vari	LE 1. Anal	TAB					

<u>e</u> × pod

13	12	11	10	9	8	7	6	S	4	ω	2	1	Sr. No.	Γ
Harvest index (%)	Straw yield per plant(g)	Grain yield per plant(g)	100 seed weight	Number of seeds per pod	Pod breadth	Pod length	Number of pods per plant	Number of primary branches per plant	Days to maturity	Days to 50% flowering	Days to initiation of flowering	Plant height (cm)	Characters	<b>ABLE 2.</b> Estimation of genetic variab
30.816	10.418	37.97	21.641	18.099	12.694	14.027	36.686	20.578	7.669	12.956	15.727	4.207	PCV (%)	oility parame
28.194	9.447	35.883	20.537	12.492	12.522	9.357	36.498	20.386	7.068	12.137	14.257	3.955	GCV (%)	ters for fourt
83.707	82.226	89.312	90.057	47.64	97.318	44.500	98.977	98.144	84.926	87.758	82.183	88.403	$h^2$	teen charac
53.138	17.647	69.859	40.148	17.763	25.448	12.859	74.800	41.604	13.418	23.422	26.626	7.662	GAM	ters

The total variability in each of the eleven characters could be partitioned into three components viz., phenotypic, genotypic and environmental variation. . In present investigation genotypic variances were relatively smaller Maximum phenotypic than phenotypic variances. variance for thirty-four genotypes was found in number of pods per plant (618.14) and Pod length (0.45) indicated lower magnitude of phenotypic variance and similar results in genotypic variance also. PCV was found to be higher than genotypic coefficient of variation for all the traits studied. Grain yield per plant (37.97%) exhibited maximum PCV. Highest value of genotypic coefficient of variation was registered for number of pods per plant (36.498%) whereas, the character plant height (3.955%) exhibited minimum genotypic coefficient of variation. These results are in agreement with Rathanaswamy et al. (1973), Shoram (1983), Balyan and Sudhakar (1985), Khapre et al. (1993) and Rao and Rao (2015) in pigeon pea.

Heritability estimates was highest for number of pods per plant (98.977%) followed by number of primary branches per plant (98.144%), pod breadth (97.318%), hundred seed weight (90.057%) and grain yield per plant (89.312%). These results are in agreement with Rathanaswamy et al. (1973), Patil et al. (1989) and Rao and Rao (2015) in pigeon pea. High heritability coupled with high genetic advance reveals the presence of lesser environmental influence and prevalence of additive gene action in their expression. . Sharma et al. (2014), Rao and Rao (2015) also reported high heritability along with high genetic advance for number of pods per plant. High heritability estimates with low genetic advance as percent of mean were indicated by the characters pod breadth, straw yield per pant, days to maturity and plant height suggested the influence of non-additive gene action. Therefore, improvement of the traits which are having high heritability along with high genetic advance would be more effective if the selection pressure in the present material could be rigorously applied (table 2).

#### **CONCLUSION & RECOMMENDATION**

The analysis of variance revealed significant variation among the genotypes for all the characters studied. The estimates of mean sum of squares showed comparatively wide range of variation for the characters number of pods per plant and days to initiation of flowering, while the lowest variation was recorded for pod length.

Among the genotypes, ICPL 20340 was the dwarfest, while genotype ICPH 2671 was the tallest. The genotype ICP 7035 recorded maximum number of pods per plant. The genotype ICPL 20325 showed maximum value for pod length. The genotype ICPL 7035 showed maximum value for pod breadth whereas maximum number of seeds per pod was recorded in ICPL 11285. The genotype ICPL 20329 (17.17 g) recorded maximum grain yield per plant. The genotype ICP 7035 recorded maximum hundred seed weight. The genotype ICPL 11339 recorded maximum straw yield per plant whereas maximum harvest index was recorded in ICPL 20329. Phenotypic variances were higher than genotypic variances for all the characters studied. Most of the characters showed comparatively higher estimates of environmental variance indicating the influence of environment on those characters. The

genotypes that showed better performance for yield and yield contributing characters can be evaluated further for selection.

#### REFERENCES

Aher, R.P., Thombre, B.B. and Dahat, D.V. (1998) Genetic and character association in pigeon pea. *Legume Res.*, **21** (1): 41-44.

Balyan, H.S. and Sudhakar, M.V. (1985) Variability, character association and path coefficient studies on genotypes of early maturing group in pigeon pea. *Madras Agric. J.*, **72** (3): 168-172.

Burton, G.W and E.H. De Vane (1953) Estimating heritability in tall fescue (*Festuca arundinacea*) from replicated clonal material. *Agron. J.*, **45**: 478-481.

Chandra, T., Tripathi, B.K. and Katiyar, R.P. (1975) Genetic variability, heritability and genetic advance of yield and it's components in arhar *J. Maharastra Agril. Univ.*, **6**: 95-99.

Dodake, S.S., Patil, B.B., Gare, B.N. and Buril, A.V. (2009) Genetic variability and correlation studies in pigeon pea under sub-montane zone of Maharastra. *J. Maharatra agril.* Univ., **34**(2): 144-146.

Khapre, P.R., Pawar D.T., Misal, V.M. and Chavan, K.M. (1993) Genetic variability, correlation and path coefficient analysis in newly developed strains of pigeon pea. *J. Maharastra agric. Univ.*, **18**(3): 494-496.

Patil, H.S., Narkhede, B.N. and Deokar, A.B. (1989) Genetic parameters, character association and path analysis in pigeon pea. *J. Maharastra agric. Univ.*, **14**(1): 54-56.

Rao Jagan Mohan and Rao Thirumalala. (2015) Genetic analysis for yield and its components in pigeon pea. *Intl. J. of applied biology and pharmaceutical technology.*, **6**(2): 189-190.

Rathanaswamy, R., Veeraswamy, R., Regupathy, A. and Palaniswamy, G.A. (1973) Studies on genetic variability of certain quantitative characters in red gram. *Madras Agric. J.*, **63**(3): 204-206.

Sharma Ritesh, Gangwar Raveesh Kumar, Yadav Vivek (2014) A study on genetic variability and correlation in pigeon pea. *Intl. J. of Sci. and Res.*, **3**(9): 826-828.

Shoram, J. (1983) Studies on genetic variability for some quantitative characters in pigeon pea. *Madras Agric. J.*, **70**(1): 146-148.

Venkateswarlu, O. (2001) genetic variability in pigeon pea. *Legume Res.*, **24**(3): 205-206.

Vijayalakshmi, P., Pavankumar, D., Sreelaxmi, A., Anuradha, G., Anuradha, C. (2013) Correlation, variability and heritability in pigeon pea. *Advances in Bio Res.*, **4**(2):129 134.