www.thebioscan.in

# STUDY OF VARIABILITY, CORRELATION AND PATH ANALYSIS IN BRINJAL (SOLANUM MELONGENA L.)

moisture content of fruit (0.9238) and number of flowers per cluster (0.9098).

## KRISHNA PATEL\*, N.B. PATEL, A. I. PATEL, HETAL RATHOD AND DHARMISHTA PATEL

Department of Vegetable Science,

Aspee College Of Horticulture And Forestry, Navsari Agricultural University, Navsari-396 450 e-mail:- Krishnaptl90@Gmail.Com

#### **KEYWORDS** Variability Heritability Geneticadvance correlation Path analysis.

**Received on :** 17.07.2015

Accepted on : 19.10.2015

\*Corresponding author

## **INTRODUCTION**

Solanum is a large and important genus of the family Solanaceae. The egg plant or brinjal or aubergine (Solanum melongena L., 2n = 24) represents the non-tuberous group of Solanum species. Brinjal is the most common popular and widely grown, vegetable crop of both tropic and sub-tropics of the world. It is being grown extensively in India, Bangladesh, Pakistan, China, Philippines, France, Italy and USA. Due to its highest production potential and availability of the produce to consumers, it is also termed as poor man's vegetable (Kumar et al., 2014). India has accumulated wide range of variability in this crop. Further, the crop exhibits rich genetic diversity and scope for improvement for various horticultural traits.

ABSTRACT

The success of any crop improvement programme depends upon the nature and magnitude of genetic variability existing in breeding material with which plant breeder is working, choice of parents for hybridization and selection procedure (Meena and Bahadur, 2013). Genetic variability is essentially the first step of plant breeding for crop improvement which is immediately available for germplasm which is considered as the reservoir of variability for different characters (Vavilov, 1951). Phenotypic and genotypic coefficients of variation are useful in detecting amounts of variability present in germplasm. Heritability and genetic advance help in determining the influence of environment in expression of characters and the extent to which improvement is possible after selection (Robinson *et al.*, 1949). Heritable variation can be effectively studied in conjunction with genetic advance. High heritability

An experiment was conducted on *rabi* 2012-13 for evaluation of 35 genotypes of brinjal for 21 characters revealed that PCV was greater than corresponding GCV for all traits. Maximum PCV (56.26%) and GCV (55.68%) were registered for fruit length: diameter ratio. High estimate of heritability coupled with high genetic advance for most of the yield and its contributing characters which indicating phenotypic selection would be effective for the genetic improvement in these traits. Fruit yield displayed highly significant and positive correlation with number of flowers per cluster (0.49), number of flowers per plant (0.61), number of fruits per cluster (0.41), number of fruits per plant (0.72) and number of branches per plant (0.48) at genotypic level. Path co-efficient study indicated the highest positive direct effect of fruit length : diameter ratio (2.4326) on fruit yield per plant followed by number of fruits per plant (1.2644), fruit diameter (1.2539), dry matter content of fruit (0.9633),

alone is not enough to make efficient selection in segregating generation and needs to be accompanied by a substantial amount of genetic advance (Johanson *et al.*, 1955).

Indirect selection in such a situation is more effective and study of correlation among different economic traits are therefore, essential for an effective selection programme because selection for one or more trait results in correlated response for several other traits and sequence of variation will also be influenced. Hence, the knowledge of genotypic and phenotypic correlation between yield and its contributing characters is very essential.

Correlation studies measure only mutual association between two traits and it does not imply the cause and effect of relationship. Path analysis is a standardized partial regression analysis, which further permits the partitioning of correlation coefficient in to components of direct and indirect effects of independent variable on the dependent variable (Wright, 1921).

## MATERIALS AND METHODS

The present investigation was carried out at Regional Horticultural Research Station, N.A.U., Navsari, during rabi seasons of 2012-13. The experiment was laid out in a randomized block design with three replications. Each genotype consists of three rows with a spacing of 90 x 60 cm and the crop was raised as per package of practices, recommended by of Navsari Agricultural University, Navsari, Gujarat. The crop was maintained properly till last harvest

and observation on growth, yield as well as its contributing characters was noted on five randomly selected plants in each plot at different stages of crop growth. The analysis of variance was done as per Pense and Sukhatme (1978) and genotypic and phenotypic coefficient of variation by Weber and Moorthy (1952). Heritability and genetic advance were calculated according to Johnson *et al.* (1955) and Allard (1960), respectively. Path analysis was done as per the procedure outlined by Wright (1921) and Dewey and Lu (1959).

### **RESULTS AND DISCUSSION**

The analysis of variance revealed the highly significant differences among the genotypes for all the 21 traits studied indicating there is substantial genetic variability for these traits. The range, general mean, genotypic and phenotypic coefficient of variation, heritability and genetic advance in per cent of mean for all traits are presented in Table 1. The results revealed that phenotypic coefficient of variation (PCV) was higher than its respective genotypic coefficient of variation (GCV) for all the characters studied but minimal differences between them. The GCV helps in comparison and measurement of genetic variability among different characters. High magnitude of GCV was recorded for number of fruits per plant. (Lohakare et al., 2008; Sherly and Santhi, 2008; Naik et al., 2010; Muniappan et al., 2010; Kumar et al., 2011; Sabeena et al., 2011: Dhaka and Soni, 2012 and Kumar et al., 2013) and moderate for fruit yield per plant (Lohakare et al., 2008 and Dhaka and Soni, 2012), for plant height (Sherly and Santhy, 2008 and Tripathi et al., 2009) which indicated that there is considerable scope for improving these characters in desirable direction through a selection programme.

High estimate of heritability coupled with high genetic advance was observed in plant height (Sherly and Santhy, 2008, Tripathi *et al.*, 2009 and Meena and Bahadur, 2014), number of flowers per cluster (Sabeena *et al.*, 2011), number of flowers

per plant, number of fruits per cluster (Lohakare *et al.*, 2008 and Naik *et al.*, 2010), number of fruits per plant (Sabeena *et al.*, 2011; Dhaka and Soni, 2012; Kumar *et al.*, 2013 and Meena and Bahadur, 2014), fruit yield per plant (Kumar *et al.*, 2011; Dhaka and Soni, 2012 and Kumar *et al.*, 2013), average fruit weight, number of branches per plant, length of peduncle, fruit length, fruit diameter, fruit length: diameter ratio, fresh weight of fruit and phenol content (Table 1).

Prior to any breeding programme for the improvement in crops, it is imperative to obtain information regarding the interrelationship of different plant characters with yield and among themselves since it facilitates a quick assessment of high yielding genotypes in selection programme. The real or true association could be known only through genotypic correlation which eliminates the environmental influence.

The estimation of genotypic correlations were higher in magnitude than the corresponding phenotypic values for all characters which indicated that there was a high degree of inter-relationship between two variables at genotypic level (Table 2). Fruit yield showed highly significant and positive association with number of fruits per plant. Similar observation obtained (Nalini et al., 2009; Muniappan et al., 2010; Dahatonde et al., 2010 and Shinde et al., 2012). Fruit yield per plant had also positive and significant correlation with number of branches per plant (Nalini et al., 2009 and Shinde et al., 2012), number of flowers per cluster, number of flowers per plant and number of fruits per cluster which indicating that there was simultaneous selection for these characters might bring an improvement in fruit yield plant. Highly significant at genotypic levels with days to 1st flowering, days to 50 per cent flowering and significant with number of days to first picking. Similar results was reported by Muniappan et al. (2010).

The path coefficient analysis permits the separation of direct and indirect effects through related traits by partitioning the genotypic correlation coefficients (Table 3) The path analysis

	<b>D</b>			•		
Table 1.	Range mean	and components o	t variance t	or various	traits in	hrinial
rubic ii	Runge, mean	unu components o	i variance i	or various	traits in	orinjur

Sr. No	Character	Range	GCV%	PCV%	Heritability (Broadsense%)	Genetic advance	Geneticadvance % of mean
1.	Days to 1 <sup>st</sup> flowering	44.33-9.67	6.08	8.36	53.00	4.90	9.13
2.	Days to 50 per cent flowering	54.00-68.33	4.55	6.90	43.60	3.92	6.20
3.	Days to first picking	65.00-86.67	7.45	8.72	73.01	9.78	13.11
4.	Days to last picking	139.33-160.67	3.09	4.33	51.19	6.91	4.56
5.	Plant height (cm)	71.74-117.54	12.04	13.08	84.74	22.08	22.84
6.	Number of flowers per cluster	1.00-7.33	37.74	38.32	96.98	3.04	76.56
7.	Number of flowers per plant	26.73-112.13	27.98	28.67	95.27	41.72	56.27
8.	Number of fruits per cluster	1.00-4.00	39.97	40.76	96.17	1.73	80.75
9.	Number of fruits per plant	12.27-44.13	29.19	30.75	90.12	15.88	57.10
10.	Fruit yield per plant(kg/plant)	1.37-3.27	18.46	20.34	82.34	0.82	34.50
11.	Average fruit weight (g)	57.61-179.29	26.63	27.15	96.14	44.84	53.79
12.	Number of branches per plant	6.27-13.27	16.35	19.60	69.61	2.77	28.11
13.	Length of peduncle (cm)	4.53-9.21	21.06	22.23	89.79	2.32	41.11
14.	Fruit length(cm)	7.93-26.23	31.20	31.77	96.43	9.03	63.12
15.	Fruit diamater (cm)	1.94-5.65	31.44	32.02	96.40	2.26	63.59
16.	Fruit length:diameter ratio	1.71-10.18	55.68	56.26	97.95	5.39	113.52
17.	Fresh weigh of fruit(g)	58.52-165.52	24.65	24.67	99.84	40.85	50.74
18.	Moisture content of fruit (%)	90.09-91.22	0.27	0.29	88.23	0.48	0.53
19.	Dry matter content of fruit (%)	8.85-9.99	2.80	2.87	95.64	0.53	5.65
20.	Total phenols (mg/100g)	0.63-1.91	32.72	32.73	99.92	0.87	67.37
21.	Vitamin C (mg/100g)	9.93-16.27	12.56	12.66	98.49	3.17	25.69

		-							• ··	-	
Charact	ers	1F	50%F	FP	LP	РН	NFC	NFP	FC	FP	AFW
Yield	rg	-0.47**	-0.53**	-0.39*	-0.19	0.09	0.49**	0.61**	0.41**	0.72**	0.09
	rp	-0.29	-0.3	0.32	-0.12	0.08	0.44	0.51	0.39	0.61	0.09
1F	rg		1.2	0.78**	0.06	0.27	-0.53**	-0.43**	-0.45**	-0.49**	0.23
	rp		0.63**	0.51**	0.05	0.09	-0.37**	-0.27**	-0.34**	-0.39**	0.16
50%F	rg			0.57**	0.1	0.3	-0.62**	-0.49**	-0.57**	-0.64**	0.27
	rp			0.66**	0.06	0.14	-0.40**	-0.36**	-0.36**	-0.39**	0.16
FP	rg				0	-0.13	-0.35	-0.31	-0.44**	-0.42**	0.09
	rp				0.02	-0.13	-0.30**	-0.28**	-0.37**	-0.34**	0.07
LP	rg					0.18	-0.18	-0.40*	-0.43**	-0.55**	0.41**
	rp					0.12	-0.14	-0.28**	-0.29**	-0.36**	0.26**
PH	rg						-0.26	-0.08	-0.12	-0.1	0.27
	rp						-0.26**	-0.09	-0.08	-0.1	0.25**
NFC	rg							0.79**	0.71**	0.76**	-0.46**
	rp							0.77**	0.68**	0.70**	-0.44**
NFP	rg								0.68**	0.91**	-0.54**
	rp								0.64**	0.83**	-0.51**
FC	rg									0.78**	-0.48**
	rp									0.74**	-0.46**
FP	rg										-0.55**
4 5347	rp										-0.53**
AFW	rg										
	rp										
INDE	ig										
DI	ra										
r L	rp										
FI	ra										
11	rn										
ED	ra										
	rn										
	rø										
2.0	rn										
FW	rg										
	rp										
мс	rg										
_	rp										
DC	rg										
	rp										
TP	rg										
	rp										
1	•										

Table 2: Genotypic and phenotypic correlation of fruit yield per plant with other characters in various genotypes of brinjal

## Table 2: Cont.....

Characters	NBP	PL	FL	FD	L:D	FW	МС	DC	ТР	Vt.C
Yield	0.48**	-0.18	0.02	0.09	0.02	0.12	0.1	-0.12	0.01	-0.21
	0.31	-0.15	0.01	0.09	0.01	0.11	0.08	-0.09	0.01	-0.18
1F	-0.3	-0.03	0.31	-0.09	0.21	0.22	0.12	-0.12	-0.1	-0.21
	-0.20*	-0.02	0.23*	-0.05	0.14	0.16	0.12	-0.08	-0.08	-0.15
50%F	-0.26	-0.17	0.06	0.1	-0.03	0.27	0.15	-0.12	-0.07	-0.16
	-0.20*	-0.09	0.04	0.09	-0.01	0.18	0.12	-0.08	-0.04	-0.09
FP	0.3	0.03	0.25	0.14	0.17	0.12	-0.02	0.03	0.18	-0.28
	-0.24*	0.05	0.20*	-0.1	0.14	0.1	-0.01	0.02	0.15	-0.23
LP	0.09	0.05	-0.12	0.39*	-0.26	0.26	-0.03	-0.01	0.11	-0.02
	0.01	0.05	-0.07	0.29**	-0.19*	0.18	-0.03	0.01	0.08	-0.01
PH	0.22	-0.04	0.04	0.21	-0.05	0.22	-0.11	0.22	0.01	-0.11
	0.16	-0.07	0.01	0.18	-0.06	0.20*	-0.12	0.21*	0.01	-0.09
NFC	0.37*	-0.01	0.19	-0.19	0.18	-0.5	-0.13	0.11	0.06	-0.06
	0.34**	0.01	0.19*	-0.18	0.18	-0.49**	-0.12	0.11	0.06	-0.06
NFP	0.50**	-0.09	0.3	-0.36*	0.38*	-0.52**	-0.07	0.14	-0.13	-0.16
	0.45**	-0.06	0.29**	-0.35**	0.37**	-0.51**	-0.06	0.13	-0.13	-0.16
FC	0.29	-0.05	0.3	-0.51*	0.42*	-0.39*	-0.06	0.09	0.02	0.02
	0.25**	-0.04	0.29**	-0.49**	0.40**	-0.38**	-0.05	0.07	0.02	0.02
FP	0.51**	-0.18	0.2	-0.32	0.31	-0.50**	0.01	0.01	-0.06	-0.09
	0.39**	-0.14	0.20*	-0.30**	0.31**	-0.47**	-0.01	0.01	-0.05	-0.09

KRISHNA PATEL et al.,

#### Table 2: Cont.....

Characters	NBP	PL	FL	FD	L:D	FW	MC	DC	TP	Vt.C
AFW	-0.23	-0.03	-0.15	0.43**	-0.3	0.92**	0.1	-0.09	0.09	-0.15
	-0.16	-0.03	-0.15	0.41**	-0.29**	0.90**	0.1	-0.1	0.08	-0.15
NBP		-0.06	-0.13	0.11	-0.14	-0.22	-0.18	0.28	0.16	0.04
		-0.04	-0.09	0.07	-0.12	-0.18	-0.13	0.19	0.13	0.03
PL			0.24	-0.06	0.21	-0.22	-0.18	0.28	0.16	0.04
			0.23*	-0.05	0.19*	-0.07	-0.13	0.11	0.04	0.11
FL				-0.73**	0.92**	-0.13	-0.01	-0.03	0.01	-0.2
				-0.69**	0.90**	-0.13	0	-0.02	0.01	-0.20*
FD					-0.89**	0.34*	0.13	-0.13	0.05	0.01
					-0.86**	0.34**	0.12	-0.12	0.04	0.01
L:D						-0.24	-0.06	0.02	-0.03	-0.1
						-0.24*	-0.05	0.02	-0.03	-0.1
FW							0.13	-0.15	0.08	-0.21
							0.12	-0.15	0.08	-0.20*
MC								-0.95**	-0.09	-0.28
								-0.89**	-0.08	-0.25**
DC									0.1	0.22
									0.1	0.21*
TP										-0.1
										-0.1
* Significant a	tP = 0.05	** Significa	pt at P = 0.01	(0.05 = 0.196, 0.0)	(1 = 0.257)					

Table 2. Diversional indiversion from the former of the second			and the set of hereing
Lable 3: Direct and indirectellect of twent	v čausai variadies on truit v	leid der diant in various	genotypes of printal
Tuble of Billeet and man cetencet of them	, caaba ranaoico on nanc j	ieia pei piane in ranous	genet, pee et stinga

Characters	Direct	Indirect ef	fect on yiel	d							
	effect on yield	1F	50%F	FP	LP	PH	NFC	NFP	FC	FP	AFW
1F	0.0964		0.11	0.07	0.006	0.02	-0.05	-0.04	-0.04	-0.04	0.02
50%F	-0.3046	-0.36		-0.17	-0.03	-0.09	0.18	0.15	0.17	0.19	-0.08
FP	-0.0186	-0.01	0.01		0.001	0.002	0.006	0.005	0.008	0.008	-0.001
LP	-0.0992	-0.006	-0.01	0.0009		-0.01	0.01	0.04	0.04	0.05	-0.04
PH	-0.0251	-0.007	-0.0076	0.003	0.004		0.006	0.002	0.003	0.002	-0.006
NFC	0.9098	-0.48	-0.56	-0.32	-0.16	0.23		0.72	0.64	0.69	-0.42
NFP	-1.1919	0.51	0.59	0.37	0.48	0.1	-0.94		-0.81	-1.09	0.65
FC	-0.6108	0.27	0.35	0.27	0.26	0.07	-0.43	-0.41		-0.48	0.29
FP	1.2644	-0.62	-0.82	-0.54	-0.7	-0.13	0.96	1.16	0.99		-0.69
AFW	-0.1538	-0.03	-0.04	-0.01	-0.06	-0.04	0.07	0.08	0.07	0.08	
NBP	0.3375	-0.1	-0.08	-0.1	0.03	0.07	0.12	0.17	0.09	0.17	-0.08
PL	-0.1963	0.006	0.03	-0.07	-0.01	0.008	0.001	0.01	0.01	0.03	0.006
FL	-1.0124	-0.32	-0.06	-0.25	0.12	-0.04	-0.19	-0.3	-0.3	-0.2	0.15
FD	1.2539	-0.11	0.13	-0.18	0.5	0.26	-0.24	-0.45	-0.64	-0.41	0.54
L:D	2.4326	0.52	-0.07	0.42	-0.65	-0.13	0.44	0.94	1.03	0.76	-0.74
FW	0.7106	0.15	0.19	0.08	0.18	0.15	-0.35	-0.37	0.27	-0.35	0.65
MC	0.9238	0.11	0.13	-0.02	-0.02	-0.1	0.12	-0.07	-0.06	0.009	0.01
DC	0.9633	-0.12	-0.12	0.03	-0.005	0.22	0.11	0.13	0.08	0.01	-0.09
TP	-0.1956	0.02	0.01	-0.03	-0.02	-0.0006	-0.01	0.02	-0.004	0.01	-0.017
Vt. C	-0.0412	0.008	0.006	0.01	0.001	0.004	0.002	0.006	-0.001	0.004	0.006

revealed that positive direct effect on yield per plant was recorded for days to 1<sup>st</sup> flowering, number of flowers per cluster, number of fruits per cluster, number of branches per plant, fruit diameter, fruit length : diameter ratio, fresh weight of fruit, moisture content of fruit and dry matter content of fruit. Similar results were also found for days to 1<sup>st</sup> flowering (Prabhu *et al.*, 2008), flowers per cluster (Nalini *et al.*, 2009), for number of fruits per plant Sharma and Swaroopan, 2000; Prabhu *et al.*, 2008; Nalini *et al.*, 2009; Muniappan *et al.*, 2010; Dahatonde *et al.*, 2010; Shinde *et al.*, 2012 and Thangamani and Jansirani, 2012) number of branches per plant (Prabhu *et al.*, 2008, Kumar *et al.*, 2011 and Thangamani and Jansirani, 2012), fruit diameter (Sharma and Swaroopan, 2000 and Thangamani and Jansirani, 2012) and dry matter content of fruit (Thangamani and Jansirani, 2012). This

indicated that these characters were directly selected for fruit yield improvement programme.

Direct negative effect on fruit yield per plant was found with days to 50 per cent flowering, days to first picking, days to last picking, plant height, number of flowers per plant, number of fruits per cluster, average fruit weight, length of peduncle, fruit length, total phenols and vitamin C. Similar results were also noted by (Sharma and Swaroopan, 2000 and Shinde et *al.*, 2012) for 50 per cent flowering (Praneetha, 2006; Naliyadhara et *al.*, 2007; Prabhu et *al.*, 2008; Kumar et *al.*, 2011; Dahatonde et *al.*, 2010; Muniappan et *al.*, 2010 and Thangamani and Jansirani, 2012), for plant height (Sharma and Swaroopan, 2000 and Kumar et *al.*, 2011), for fruit length (Thangamani and Jansirani, 2012) for total phenol content and for vitamin C (Praneetha, 2006). Therefore, indirect selection practiced on

Table	3:	Cont	•
-------	----	------	---

Characters	, NBP	PL	FL	FD	L:D	FW	мс	DC	TP	Vt. C	Correlation Coefficient
10	0.02	0.002	0.02	0.000	0.02	0.02	0.01	0.01	0.01	0.02	0.47**
	-0.02	-0.003	0.03	-0.009	0.02	0.02	0.01	-0.01	-0.01	-0.02	-0.47
50 %I	0.05	-0.02	-0.03	0.009	0.02	-0.00	-0.04	0.03	0.02	0.05	-0.33
	0.005	-0.01	-0.004	0.002	-0.003	-0.002	0.01	-0.0007	-0.003	0.005	-0.39*
	-0.009	-0.005	0.01	-0.03	0.02	-0.02	0.003	0.0006	-0.01	0.002	-0.19
РН	-0.005	0.001	-0.001	0.005	0.001	-0.005	0.002	-0.005	-0.0001	0.003	0.09
NFC	0.34	-0.007	0.17	-0.176	0.16	-0.45	-0.12	0.1	0.05	-0.059	0.49**
NFP	-0.6	0.1	-0.36	0.43	-0.46	0.63	0.09	-0.17	0.16	0.19	0.61**
FC	-0.17	0.03	-0.18	0.31	-0.26	0.24	0.04	-0.05	-0.01	-0.01	0.41**
FP	0.65	-0.23	0.26	-0.41	0.39	-0.63	0.01	0.02	-0.07	-0.12	0.72**
AFW	0.03	0.004	0.02	-0.06	0.04	-0.14	-0.01	0.01	-0.01	0.02	0.09
NBP		-0.02	-0.04	0.03	-0.04	-0.07	-0.06	0.09	0.05	0.01	0.48**
PI	0.01		-0.04	0.01	-0.04	0.01	0.02	-0.02	-0.009	-0.02	-0.18
FI	0.13	-0.24		0.73	-0.93	0.13	0.01	0.03	-0.008	0.2	0.02
FD	0.14	-0.07	-0.91	011 0	-1.12	0.43	0.16	-0.16	0.06	0.004	0.09
1.0	-0.34	0.51	2 25	-2.17	=	-0.59	-0.15	0.05	-0.09	-0.25	0.02
E.D	0.16	0.05	0.09	0.24	0.17	0.55	0.15	0.05	0.05	0.15	0.12
MC	-0.10	-0.03	-0.09	0.24	-0.17	0.10	0.09	-0.11	0.00	-0.15	0.12
MC DC	-0.17	-0.13	-0.01	0.12	-0.05	0.12	0.00	-0.00	-0.06	-0.26	0.1
	0.27	0.09	-0.03	-0.12	0.02	-0.15	-0.92	0.00	0.1	0.21	-0.12
	-0.03	-0.009	-0.001	-0.009	0.007	-0.01	0.017	0.02		0.02	0.01
Vt. C	-0.002	-0.005	0.008	-0.0002	0.004	0.008	0.01	-0.009	0.004		-0.21

Residual effect = 0.9447

these characters will result in the improvement of respective characters and ultimately fruit yield.

#### REFERENCES

Allard, R. W. 1960. "Principles of Plant Breeding". J. Willey and Sons, Inc. New York. pp. 224-232.

Dahatonde, K., Dod, V. N., Nagre, P. K. and Wag, A. P. 2010. Correlation and path analysis studies in purple fruited brinjal. *Asian J. Hort.* 5(2): 428-430.

Dhaka, S. K. and Soni, A. K. 2012. Genetic variability in brinjal (Solanum melongena L.). Asian J. Hort. 7(2): 537-540.

Dewey, D. R. and Lu, K. H. 1959. A correlation and path coefficient analysis of components of crested wheat grass seed production. *Agronomy J.* 51: 515-518.

Johnson, H. W., Robinson, H. F. and Comstock, R. E. 1955. Estimates of genetic and environmental variability in soybean. *Agron. J.* 47: 314-318.

Kumar, R., Anjali, K., Singh, A. K. and Maurya, S. 2014. Screening of bacterial wilt resistant accessions ofbrinjal for jharkhand region of India.*The Ecoscan.* 8(1&2): 67-70.

Kumar, R. S., Arumugam, T., Anandkumar, C. R. and Premalakshmi, V. 2013. Genetic variability for quantitative and qualitative characters in brinjal (*Solanum melongena* L.). *Academic J.* 8(39): 4956-4959.

Kumar, S., Sharma, J. P. and Chopra, S. 2011. Studied on Variability, heritability and genetic advance for morphological and yield traits in brinjal (*Solanum melongena* L.). *Mysore J. Agric. Sci.* **45(1):** 63-66

Lohakare, A. S., Dod, V. N. and Peshattiwar, P. D. 2008. Genetic variability in green fruited brinjal. *Asian J. Hort.* 3(1): 114-116.

Meena, O. P. and Bahadur, V. 2014. Assessment of genetic variability, heritability and genetic advance among tomato (*Solanum lycopersicum* L.) germplasm. *The Bioscan.* 9(4): 1619-1623.

**Meena, O. P. and Bahadur, V. 2013.** Assessment of breeding potential of tomato (*Lycopersicon esculentum* Mill.) germplasm using D<sup>2</sup> analysis. *The Bioscan.* **8(4):** 1145-1148.

Muniappan, S., Saravannan, K. and Ramya, B. 2010. Studies on Genetic divergence and variability for certain economic characters in egg plant (Solanum melongena L.). Electronic J. Plant Breeding. 1(4): 426-465.

Naik, K., Sreenivasulu, G. B., Prashanth, S. J., Jayaprakashnarayan, R. P., Madalageri, M. B. and Mulge, R. 2010. Studies on genetic variability and its importance in brinjal (*Solanum melongena* L.). *Asian J. Hort.* **4(2)**: 380-382.

Nalini, A. D., Salimath, P. M. and Patil, S. A. 2009. Association and path coefficient analysis in elite germplasm lines of brinjal (*Solanum melongena* L.). *Karnataka J. Agric Sci.* **22(5):** 965-966.

Naliyadhara, M. V., Golani, I. J., Mehta, D. R. and Purohit, V. L. 2007. Genetic variability, correlation coefficient and path analysis in brinjal. *Orissa J. Hort.* 35(2): 92-96.

Panse, V. G. and Shukhatme, P. V. 1978. "Statistical Methods For Agricultural Workers" (3<sup>rd</sup> end.).*Indian Council of Agricultural Research, New Delhi, India*. pp. 152-161.

**Prabhu, M.; Natarajan, S. and Veeraragavathatham, D. 2008.** Correlation and path analysis in egg plant (*Solanum melongena* L.). *Indian J. Agri. Res.* **42(3):** 232-234.

**Praneetha, S. 2006.** Path analysis studies in brinjal (*Solanum melongena* L.). *Indian J. Hort.* **63(3):** 335-337.

Sabeena, F. A., Mehta, N., Ansari, S. and Gavel, J. P. 2011. Variability studies in brinjal (*Solanum melongena* L.) in Chhattisgarh plains. *Electronic J. Plant Breeding*. 2(2): 275-281.

Sharma, T. V. R. S. and Swaroop, K. 2000. Genetic variability and character association in brinjal (*Solanum melongena* L.). *Indian J. Hort.* **57(1):** 59-65.

Sherly, J. and Shanthi, A. 2008. Variability, Heritability and Genetic advance in brinjal (*Solanum melongena* L.). Orissa J. Hort. **36(2)**: 24-28.

Shinde, K. G., Birajdar, U. M., Bhalekar, M. N. and Patil, B. T. 2012. Correlation and path analysis in egg plant (*Solanum melongena* L.). *Veg. Sci.* **39(1):** 108-110.

Thangamani, C. and Jansirani, P. 2012. Correlation and path analysis studies on yield and attributing characters in brinjal (*Solanum melongenaL.*). *Electronic J. Plant Breeding*. **3(3):** 939-944.

Tripathi, M. K., Singh, A. K., Singh, B. K. and Rai, V. K. 2009.

Genetic variability, heritability, genetic advance among different quantitative characters of brinjal (*Solanum melongena* L.). *Haryana J. Hort. Sci.* **38(3&4):** 334-335.

Vavilov, N. I. 1951. The origin variation immunity and breeding of cultivated plant. *Soil Science*. 72: 482.

**Weber, C. R. and Moorthy, B. R. 1952.** Heritable and non heritable relationship and variability of oil content and agronomic characters in the F<sub>2</sub> generation of soybean crosses. *Agron. J.* **44:** 202-209.

Wright, S. 1921. Correlation and causation. J. Agric. Res. 20: 557-585.