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Research Article

Studies on Variability, Character Association and Path Analysis on Groundnut (Arachis hypogaea L.)

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ABSTRACT

In the present investigation 18 Spanish bunch groundnut genotypes were used for variability, character association and path analysis studies in 19 plant characters. In general higher GCV and heritability estimates coupled with high genetic advance as per cent of mean recorded for LLS severity, reducing sugar and non-reducing sugar indicated lesser influence of environment in expression of these characters and these characters are controlled by additive gene effect, hence, amenable for simple selection. Pod yield per plant exhibited positive significant association with number of pods per plant, total sugar, kernel yield, non reducing sugar, test weight, SCMR, harvest index, oil content and shelling per cent, whereas, LLS severity, reducing sugar, kernel yield, stomata length, LLS severity, test weight, SCMR, days to maturity and oil content exerted the positive direct effect on pod yield, whereas, non-reducing sugar, stomata frequency, shelling per cent and harvest index had maximum indirect direct effects on pod yield per plant. Thus, due emphasis should be placed on these characters while selecting genotypes for high yield with LLS tolerance in groundnut

Key words: Genetic variability, Heritability, Character association and Path analysis

INTRODUCTION

Groundnut (*Arachis hypogaea* L.) is an important annual legume oilseed crop grown in tropical and sub-tropical regions of the world. It is a valuable source of dietary protein, oil and fodder for livestock. Its kernel contains 48-50% oil and 26-28% protein and a rich source of dietary fibre, minerals and vitamins. The low productivity and production of groundnut is mainly due to several biotic

and abiotic stresses affecting the crop at various growth stages. Among the biotic stresses, late leaf spot (*Phaeoisariopsis personata* [(Berk. and Curt.) Deighton]) and rust (*Puccinia arachidicola* Speg.) are most important. These two diseases often occur together and causes up to 50-70% of yield losses in the crop²⁹. Use of fungicides to control leaf spots usually increases production costs by 10%.

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Therefore, development of cultivars resistant/tolerant to these diseases could be effective in reducing the production costs and the detrimental effects of chemicals on our ecosystem and improving quality production. Genetic variability is the basic requirement for crop improvement as this provides wider scope for selection. Thus, effectiveness of selection is dependent upon the nature, extent and magnitude of genetic variability present in the population and the extent to which it is heritable. Understanding the relationship between yield and its components is of the paramount importance for making the best use of the relationships in selection. The data obtained from correlation coefficient can be augmented by path analysis. Path coefficient analysis splits the genotypic correlation coefficient into the measure of direct and indirect effects. Hence, in present investigation an attempt was made to assess the variability of important pod attributes and LLS disease resistance components along with the indices of variability i.e. genotypic coefficient of variation (GCV), phenotypic coefficient of variation (PCV), heritability in broad sense (h2 bs), genetic advance (Gs) and genetic advance as percentage of mean (GAM), the magnitude of relationship of individual yield components on yield, interrelationships among themselves and to measure their relative importance in the different genotypes of groundnut.

MATERIAL AND METHODS

The field experiment was carried out in randomized block design with three replications at Oilseed Research Station, Latur (MS) during Kharif, 2017. The experimental material comprised of 18 groundnut genotypes including three checks viz., JL-24, LGN-1 and LGN -123. The sowing was carried out by dibbling at the spacing of 30 x 10 cm between the rows and plant, respectively. The data was collected on five randomly chosen plants from each treatment and observations were recorded on pod yield plant per plant (g), number of pod plant per plant, kernel yield plant per plant(g), days to maturity, shelling per cent, test weight (g), harvest index (%), oil content (%), LLS

disease severity (%), stomata frequency, stomata size (µm), SPAD chlorophyll meter reading, reducing sugar (mg/g), non-reducing sugar (mg/g) and total sugar(mg/g). Late leaf spot severity was scored on the 1-9 point scale as described by Subrahmanyam et al.²⁹ and then the score was transformed to percentage using arc-sine arc-sine transformation formula. Analysis of variance was carried out as per the method suggested by Panse and Sukhatme²⁴. Phenotypic and genotypic coefficients of variations were computed as per Burton⁴, heritability (broad sense) and genetic advance as followed as per Allard¹. The genotypic and phenotypic coefficients of correlations were calculated using the method given by Johnson et al.¹². Path coefficient analysis was carried out by using phenotypic and genotypic correlations coefficients as per the method suggested by Dewey and Lu⁶.

RESULTS AND DISCUSSION

In the present study the analysis of variance for 19 characters revealed that significant differences were observed for all the characters. The variations of different traits under this study revealed that the phenotypic coefficient of variations (PCV) were higher than genotypic coefficient of variations (GCV) for all the characters studied indicating the role of environmental variance in the total variance (Table 1). Lesser differences were observed between PCV and GCV in all cases indicated greater role of genetic components and less influence bv environment. Similar observations were also reported by Vasanthi et al.,³² and Ashish et al.,². Higher GCV estimates were observed for LLS severity, reducing sugar and non-reducing sugar. Hence, these characters can be relied upon and simple selection can be practiced for further improvement. These findings are in accordance with the earlier reports of Dolma et al.,⁷, Padmaja et al.,²³ and Ashish et al.,² for LLS severity; Chari⁵ for non-reducing sugar, Giri *et al.*,⁸ and Sawargaonkar *et al.*,²⁷ for reducing sugar and LLS severity; Kahate and Toprope¹⁵ for LLS disease severity, reducing sugar and non- reducing sugar. Moderate GCV 1382

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were recorded for stomata breadth, total sugar, stomata length, number of pod per plant, test weight kernel yield and pod yield per plant. The results are in accordance with Vasanthi et al.,³² and Kumar et al.,¹⁶ for test weight and kernel yield; Rao et al.,²⁶ for number of pods per plant, kernel yield and pod yield; Shukla and Rai³⁰ for kernel yield and pod yield. The lowest GCV values were recorded for oil content, shelling per cent, days to maturity, stomata frequency, harvest index and SCMR. The similar estimates were recorded for shelling per cent and oil content by Wani et $al.^{33}$ and for stomatal frequency per mm² by Kahate and Toprope¹⁵

High heritability coupled with high genetic advance as per cent of mean recorded for LLS severity, reducing sugar and nonreducing sugar indicated lesser influence of environment in expression of these characters and these characters are controlled by additive gene effect, hence, amenable for simple selection. These results are in accordance with earlier reports of Vasanthi et al.,³², Padmaja et al.,²³ and Ashish et al.,² for LLS severity; Giri et al.,⁸ and Sawargaonkar et al., ²⁷ for LLS severity and reducing sugar; Kahate and Toprope¹⁵ for LLS severity, reducing sugar non- reducing. High to moderate and heritability coupled with moderate genetic advance as per cent of mean has been noticed in stomata breadth, total sugar, test weight, SCMR, stomata length, kernel yield, pod yield and number of pods per plant. Similar result found by Nindini et al.,²² for pod yield and kernel vield; Patil et al.,²⁵ for test wt. and kernel wt; Kahate *et al.*,¹⁴ for test wt., stomatal length and stomatal breadth. High to moderate heritability and low genetic advance as per cent of mean was noticed for stomata frequency, oil content, and days to maturity and shelling indicated the influence of nonadditive gene action. This suggests limited scope for further improvement of these characters. The results are in accordance with earlier work of Sunneta et al.,³¹ for oil content and Kahate and Toprope¹⁵ for stomata frequency and oil content.

In the present study, genotypic correlations were higher than phenotypic correlations for most of the characters. These indicate that the strong inherent association between the characters governed largely by genetic causes and reduced by environmental forces. The environment and genotype x environment interaction played a major role in determining these associations between the characters. The results pertaining to correlation studies are presented in table 2. The pod yield per plant exhibited highest, positive and significant association with number of pod per plant followed by total sugar, kernel yield, nonreducing sugar, test weight, SCMR, harvest index, oil content and shelling per cent. The similar kinds of associations earlier reported by Sharma and Dashora²⁸ for number of pods per plant and kernel yield, Gouda Patil et al.,¹⁰ for number of pods per plant and shelling per cent, Azad and Hamid³ and Rao et al.,²⁶ for number of pods per plant, kernel yield and test weight, Kadam et al.,¹³ for number of pods per plant, harvest index, test weight and oil content, Kahate et al.,¹⁴ for kernel yield, harvest index, non- reducing sugar and test weight and John and Reddy¹¹ for number of pod per plant, kernel yield per plant, test weight and shelling per cent.

The pod yield also exhibited negative and significant association with stomata frequency, stomata size (length and breadth), LLS severity and reducing sugar. The similar kind of findings were reported by Gopal et al.9 for LLS severity, Giri et al.,8 for LLS severity and reducing sugar, Kahate et al.,¹⁴ for stomata size, stomata frequency, LLS disease severity and reducing sugar.

The positive and highly significant interrelationships were observed among yield contributing characters like number of pod per plant, kernel yield, shelling and test weight and morpho-biochemical traits like LLS severity, reducing sugar, stomata frequency and size. The results are in accordance with earlier reports of Lakshimidevamma et al.,¹⁹ for kernel yield with test wt., Mahalakshmi et *al.*,²¹ for kernel yield with test wt. and shelling; Kaur et al.,¹⁷ for LLS severity with stomatal

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Rathod and Toprope Int. J. Pure App. Bio frequency; Li Dun²⁰ for LLS severity with reducing sugar. Kahate *et al.*,¹⁴ for kernel yield with harvest index, non-reducing sugar and test wt.

The interrelationships were also negative and highly significant among yield contributing characters like number of pods per plant, kernel yield, test weight with morpho-biochemical traits like LLS severity, reducing sugar, stomata frequency and size. The similar result reported by Giri *et al.*,⁸ and Kahate *et al.*,¹⁴.

The path co-efficient studies (Table 3) indicated that total sugar, kernel yield , LLS severity , test weight , SCMR , days to maturity , stomata length and oil content exerted positive direct effect on pod yield. Hence, a direct selection criterion should be followed for these traits to improve the pod yield. Similar results were earlier reported by Giri *et al.*,⁸ for kernel yield per plant , Khan *et al.*,¹⁸ for test weight , *Zaman et al.*,³⁴ for days to maturity , Azad and Hamid³ for kernel yield and test weight and Kadam *et al.*,¹³ for oil content.

Negative direct effects on pod yield were also exhibited by some characters *viz.*, nonreducing sugar, stomatal frequency, shelling, harvest index and reducing sugar. The similar kinds of results earlier reported by Kahate *et al.*,¹⁴ for stomata frequency and shelling and Lakshmidevamma *et al.*,¹⁹ for shelling percent.

Thus for development high yielding disease resistance varieties in groundnut due emphasis should be given to LLS severity, reducing sugar, non reducing sugar, kernel vield plant⁻¹, harvest index, stomata size and frequency. All these characters had high GCV and PVC and heritability. Positive association of pod yield with number of pod per plant followed by total sugar, kernel yield, nonreducing sugar, test weight, SCMR, harvest index, oil content and shelling per cent and negative and significant association with stomata frequency, stomata size (length and breadth), LLS severity and reducing sugar indicated that these characters can be improved through direct selection.

Sr. No.	Characters	Range	Mean	GCV (%)	PCV (%)	h ² (%) (Broad sense)	Genetic advance	GA as percent of mean	
1	No. of pod plant per plan (g)	14.40-24.80	20.85	13.15	14.87	78.1	4.99	23.94	
2	Kernel yield plant per plant(g)	8.16-13.95	11.53	12.22	15.89	59.1	2.23	19.35	
3	Harvest index (%)	33.47-47.62	39.02	7.33	13.00	31.8	3.32	8.51	
4	Days to maturity	93.66-110.00	99.46	3.85	4.01	91.9	7.56	7.60	
5	Shelling (%)	60.67-69.33	65.52	3.49	4.07	73.6	4.05	6.18	
6	LLS severity (%)	2.33-63.33	21.75	79.04	79.23	99.5	35.34	162.44	
7	Test weight (g)	35.31-58.24	47.00	12.32	12.40	98.7	11.85	25.22	
8	SCMR	33.23-48.40	44.38	8.63	9.19	88.1	7.41	16.69	
9	Non reducing sugar (mg/ g)	6.10-14.40	10.40	29.56	29.81	98.4	6.28	60.41	
10	Reducing sugar (mg/ g)	0.63-3.80	1.61	50.06	50.32	98.9	1.65	102.58	
11	Total sugar(mg /)	8.80-15.03	12.01	18.94	19.14	97.9	4.64	38.62	
12	Oil content (%)	46.90-48.00	47.42	0.80	0.92	74.7	0.67	1.42	
13	Stomata frequency (Adaxial)	110.46-126.80	117.78	3.88	4.06	91.6	9.02	7.66	
14	Stomata frequency (Abaxial)	109.20-125.26	116.30	3.95	4.09	93.1	9.14	7.86	
15	Stomata length (µm) (Adaxial)	13.87-21.55	17.92	11.07	11.67	89.9	3.87	21.63	
16	Stomata length (µm) (Abaxial)	11.39-21.25	16.92	14.13	14.46	95.5	4.81	28.45	
17	Stomata breadth (µm) (Adaxial)	5.88-13.20	10.42	16.76	18.02	86.5	3.34	32.11	
18	Stomata breadth (µm) (Abaxial)	5.42-12.14	9.39	20.85	21.68	92.5	3.88	41.31	
19	Pod yield plant per plant (g)	13.44-20.56	17.57	10.69	14.36	55.4	2.88	16.39	

Table 1: Estimates of variability, heritability and genetic advance in Groundnut

Table 2: Genotypic (G) and phenotypic (P) coefficients among yield, yield contributing and morpho-biochemical characters in groundnut

		171	н	514	SH			SCMR	N.R. sugar	R. Sugar	Total sugar	Oil content	Stomata frequency		Stomata length		Stomata breath		
Characters		KY		DM		LLS Severity	TW						Adaxial	abaxial	Adaxial	abaxial	Adaxial	abaxial	РҮ
No. of pod plant per plan (g)	G P	0.9352** 0.8968**	-0.0461 0.2538	-0.1264 -0.0880	0.2921* 0.2670	-0.9468** -0.8624**	0.9745** 0.8999**	0.6583** 0.5402**	0.9821** 0.9076**	-0.9819** -0.7555**	0.9779** 0.9053**	0.3504** 0.2768*	-0.9812** -0.9447**	-0.9637** -0.9308**	-0.9571** -0.9150**	-0.9586** -0.9108**	-0.9107** -0.8733**	-0.9574** -0.9243**	0.9914** 0.9266**
Kernel yield plant per plant(g)	G P	1.0000 1.0000	-0.275* 0.2417	-0.1938 -0.1186	0.5931** 0.5070**	-0.9982** -0.8040**	0.8685** 0.8400**	0.6636** 0.4652**	0.9359** 0.7849**	-0.9162** -0.7555**	0.9388** 0.7909**	0.4342** 0.2970*	-0.9494** -0.8574**	-0.9372** -0.8370**	-0.9795** -0.8617**	-0.9840** -0.8505**	-0.9679** -0.8515**	-0.9512** -0.8446**	0.9682** 0.9714**
Harvest index (%)	G P		1.0000 1.0000	0.6775** 0.3749**	-0.633** -0.2332	0.1273 0.0397	0.0203 0.0681	-0.5677** -0.3095*	0.2492 0.2012	-0.1980 -0.165	0.2663 0.2126	0.1112 0.0120	0.0099 -0.1587	0.0044 -0.1417	-0.0273 -0.1798	-0.1053 -0.1804	-0.1744 -0.2493	-0.3309* -0.3351*	0.4421** 0.3161*
Days to maturity	G P			1.0000 1.0000	-0.2872* -0.2156	0.2332 0.2173	-0.2248 -0.2041	-0.3857** -0.3248*	-0.0879 -0.0806	0.0752 0.0723	-0.0920 -0.0831	0.0310 0.0070	0.2384 0.2080	0.2530 0.2231	0.1939 0.1678	0.1478 0.1251	0.1007 0.0486	0.0299 0.0268	-0.1337 -0.0718
Shelling (%)	G P				1.0000 1.0000	-0.4481** -0.3897**	0.4133** 0.3646**	0.7761** 0.3501**	-0.8514** 0.1209	-0.1209 -0.1159	0.1317 0.1220	0.2996* 0.1847	-0.2795* -0.2618	-0.3024* -0.2699*	-0.3919** -0.3302*	-0.3926** -0.3536**	-0.4175** -0.3299*	-0.2352 -0.2235	0.3751** 0.2914*
LLS severity (%)	G P					1.0000 1.0000	-0.9482** -0.9463**	-0.7761** -0.7263**	-0.8514** -0.8492**	0.8333** 0.8321**	-0.8540** -0.8505**	-0.3600** -0.3121*	0.9523** 0.9256**	0.9498** 0.9295**	0.9305** 0.8942**	0.9045** 0.8928**	0.8345** 0.7922**	0.8338** 0.8133**	-0.9150** -0.7971**
Test weight (g)	G P						1.0000 1.0000	0.6937** 0.6505**	0.9059** 0.9051**	-0.8938** -0.8919**	0.9062** 0.9048**	0.3499** 0.3096*	-0.9809** -0.9571**	-0.9595** -0.9455**	-0.9725** -0.9388**	-0.9836** -0.9731**	-0.9589** -0.9181**	-0.9335** -0.9156**	0.9447** 0.8382**
SCMR	G P							1.0000 1.0000	0.5180** 0.4844**	-0.5520** -0.5113**	0.5036** 0.4722**	0.2539 0.2078	-0.6487** -0.5692**	-0.6370** -0.5717**	-0.5761** -0.5071**	-0.6161** -0.5618**	-0.5337** -0.4565**	-0.4378** -0.3900**	0.6401** 0.4412**
Non reducing sugar (mg/ g)	G P								1.0000 1.0000	-0.9909** -0.9883**	0.9989** 0.9985**	0.3319* 0.2962*	-0.9377** -0.9175**	-0.9116** -0.8979**	-0.9105** -0.8820**	-0.9043** -0.8955**	-0.8878** -0.8506**	-0.9508** -0.9340**	0.9520** 0.8404**
Reducing sugar (mg/ g)	G P									1.0000 1.0000	-0.9833** -0.9786**	-0.3020* -0.2651*	0.9237** 0.9030**	0.8927** 0.8757**	0.8902** 0.8607**	0.8884** 0.8790**	0.8729** 0.8284**	0.9322** 0.9113**	-0.8623** -0.8110**
Total sugar(mg /)	GP										1.0000 1.0000	0.3411* 0.3056*	-0.9385** -0.9175**	-0.9144** -0.9007**	-0.9137** -0.8845**	-0.9060** -0.8963**	-0.8892** -0.8537**	-0.9532** -0.9227**	0.9761** 0.8461**
Oil content (%)	G P											1.0000 1.0000	-0.2550 -0.2027	-0.2258 -0.2182	-0.3063* -0.2500	-0.3254* -0.2797*	-0.3527** -0.3023*	-0.3585** -0.3140*	0.4168** 0.2846*
Stomata frequency (Adaxial)	G P												1.0000 1.0000	0.9978** 0.9804**	0.9824** 0.9569**	0.9634** 0.9503**	0.9095** 0.8807**	0.9347** 0.9227**	-0.9301** -0.8844**
Stomata frequency (Abaxial)	G P													1.0000 1.0000	0.9786** 0.9552**	0.9498** 0.9385**	0.8840** 0.8632**	0.9137** 0.9075**	-0.9850** -0.8556**
Stomata length (µm) (Adaxial)	G P														1.0000 1.0000	0.9872** 0.9583**	0.9456** 0.9017**	0.9433** 0.9270**	-0.8760** -0.8675**
Stomata length (µm) (Abaxial)	G P															1.0000 1.0000	0.9918** 0.9480**	0.9550** 0.9445**	-0.9381** -0.8675**
Stomata breadth (μm) (Adaxial)	G P																1.0000 1.0000	0.9733** 0.9316**	-0.9753** -0.8531**
Stomata breadth (μm) (Abaxial)	GP																	1.0000 1.0000	-0.9434** -0.8720**

Table 3: Path coefficients for yield contributing and mo	orpho-biochemical characters in Groundnut
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Characters	No. of pod/ plant	Kernel yield (g)	Harvest	Days to maturity	Shelling %	LLS Severity	Test weight (g)	SCMR	N.R. sugar (mg/g)	R. Sugar (mg/g)	Total Sugar (mg/g)	Oil content (%)	SF/mm ² Ad (%)	SF/mm ² Ab	Stomata length (µm)		Stomata Breadth (μm)		Correlation with pod
			index (%)	maturity	/0	(%)								()	Ad	Ab	Ad	Ab	yield
No. of pod plant	G-0.1602	0.0850	0.0535	0.1466	-0.3389	0.0984	0.1305	-0.7637	-0.1394	0.1392	-0.1345	-0.4066	0.1384	0.1181	0.1104	0.1122	0.0566	0.1108	0.9914**
per plan (g)	P 0.0423	0.0379	0.0107	-0.0037	0.0113	-0.0365	0.0380	0.0228	0.0384	-0.0380	0.0383	0.0117	-0.0399	-0.0393	-0.0387	-0.0385	-0.0369	-0.0391	0.9266**
Kernel yield plant	G0.5156	0.7207	-0.4464	-0.3141	0.9612	-0.6177	0.6506	0.0755	0.5168	-0.4849	0.5215	0.7038	-0.9487	-0.9190	-0.9874	-0.9948	-0.9686	-0.5416	0.9682**
per plant(g)	P 0.9786	0.0912	0.2637	-0.1294	0.5533	-0.8774	0.9166	0.5076	0.8565	-0.5076	0.8630	0.3241	-0.9357	-0.9133	-0.9403	-0.9281	-0.9292	-0.9216	0.9714**
Harvest index (%)	G0.0073	0.0437	-0.1587	-0.1075	0.1005	-0.0202	-0.0032	0.0901	-0.0396	0.0314	-0.0423	-0.0177	-0.0016	-0.0007	0.0043	0.0167	0.0277	0.0525	0.4421**
That Yest Index (70)	P -0.0005	0.0004	-0.0018	-0.0007	0.0004	-0.0001	-0.0001	0.0006	-0.0004	0.0003	-0.0004	0.0000	0.0003	0.0003	0.0003	0.0003	0.0005	0.0006	0.3161*
Days to maturity	G-0.0481	-0.0737	0.2576	0.3802	-0.1092	0.0887	-0.0855	-0.1467	-0.0334	0.0286	-0.0350	0.0118	0.0907	0.0962	0.0737	0.0562	0.0383	0.0114	-0.1337
Duys to maturity	P -0.0015	-0.0020	0.0063	0.0168	-0.0036	0.0036	-0.0034	-0.0054	-0.0014	0.0012	-0.0014	0.0001	0.0035	0.0037	0.0028	0.0021	0.0008	0.0004	-0.0718
Shelling (%)	G-0.0779	-0.1581	0.1689	0.0765	-0.2665	0.1194	-0.1102	-0.1253	-0.0345	0.0322	-0.0351	-0.0799	0.0745	0.0806	0.1045	0.1046	0.1113	0.0627	0.3751**
Bliefining (70)	P -0.0728	-0.1382	0.0636	0.0588	-0.2726	0.1062	-0.0994	-0.0954	-0.0330	0.0316	-0.0333	-0.0504	0.0714	0.0736	0.0900	0.0964	0.0899	0.0609	0.2914*
LLS severity (%)	G-0.4256	-0.5030	0.1917	0.3511	-0.6747	0.5057	-0.4277	-0.1687	-0.2819	0.2547	-0.2860	-0.5421	0.4339	0.4302	0.4011	0.3620	0.2565	0.2555	-0.9150**
EED severing (ve)	P -0.0068	-0.0063	0.0003	0.0017	-0.0031	0.0078	-0.0074	-0.0057	-0.0067	0.0065	-0.0067	-0.0024	0.0072	0.0073	0.0070	0.0070	0.0062	0.0064	-0.7971**
Test weight (g)	G0.4810	0.5027	0.0100	-0.1109	0.2040	-0.4680	0.4936	0.3424	0.4471	-0.4412	0.4473	0.1727	-0.4842	-0.4736	-0.4800	-0.4855	-0.4733	-0.4608	0.9447**
rest weight (g)	P 0.0638	0.0595	0.0048	-0.0145	0.0258	-0.0670	0.0709	0.0461	0.0641	-0.0632	0.0641	0.0219	-0.0678	-0.0670	-0.0665	-0.0689	-0.0651	-0.0649	0.8382**
SCMR	G0.2226	0.2244	-0.1919	-0.1304	0.1590	-0.2624	0.2346	0.3381	0.1751	-0.1866	0.1703	0.0858	-0.2193	-0.2154	-0.1948	-0.2083	-0.1805	-0.1480	0.6401**
SCIVIK	P 0.0025	0.0021	-0.0014	-0.0015	0.0016	-0.0033	0.0030	0.0046	0.0022	-0.0024	0.0022	0.0010	-0.0026	-0.0026	-0.0023	-0.0026	-0.0021	-0.0018	0.4412**
Non reducing sugar	G-0.9169	0.8737	-0.2327	0.0820	-0.1207	0.7948	-0.8457	-0.4836	-0.9336	0.9251	-0.9325	-0.3099	0.8754	0.8511	0.8500	0.8442	0.8288	0.8876	0.9520**
(mg/ g)	P -0.8407	0.7271	-0.1864	0.0746	-0.1120	0.7867	-0.8385	-0.4487	-0.9263	0.9155	-0.9250	-0.2744	0.8499	0.8317	0.8170	0.8295	0.7879	0.8652	0.8404**
Reducing sugar	G0.1531	-0.1428	0.0309	-0.0117	0.0188	-0.1299	0.1393	0.0860	0.1545	-0.1559	0.1533	0.0471	-0.1440	-0.1391	-0.1388	-0.1385	-0.1361	-0.1453	-0.8623**
(mg/ g)	P 0.2245	-0.1886	0.0414	-0.0180	0.0289	-0.2077	0.2226	0.1276	0.2467	-0.2496	0.2443	0.0662	-0.2254	-0.2186	-0.2148	-0.2194	-0.2068	-0.2275	-0.8110**
Total maar(ma /)	G 0.2433	0.1936	0.3386	-0.1169	0.1675	-0.6859	0.1522	0.6403	0.8700	-0.2502	0.9715	0.4337	-0.8933	-0.9626	-0.18617	-0.7519	-0.6306	-0.7120	0.9761**
Total sugar(mg /)	P 0.6217	0.5431	0.1460	-0.0570	0.0838	-0.5840	0.6213	0.3242	0.6857	-0.6720	0.6867	0.2099	-0.6300	-0.6185	-0.6074	-0.6154	-0.5862	-0.6433	0.8461**
O^{1} - (0)	G 0.0783	0.0970	0.0249	0.0069	0.0669	-0.0804	0.0782	0.0567	0.0742	-0.0675	0.0762	0.2234	-0.0570	-0.0504	-0.0684	-0.0727	-0.0788	-0.0801	0.4168**
Oil content (%)	P 0.0025	0.0027	0.0001	0.0001	0.0017	-0.0029	0.0028	0.0019	0.0027	-0.0024	0.0028	0.0091	-0.0019	-0.0020	-0.0023	-0.0026	-0.0028	-0.0029	0.2846*
Stomata frequency	G 0.4948	-0.4787	-0.0050	-0.1202	0.1409	-0.4802	0.4946	0.3271	0.4728	-0.4658	0.4732	0.1286	-0.5042	-0.5031	-0.4954	-0.4858	-0.4586	-0.4713	-0.9301**
(Adaxial)	P 0.0411	-0.0373	0.0069	-0.0090	0.0114	-0.0402	0.0416	0.0247	0.0399	-0.0393	0.0399	0.0088	-0.0435	-0.0426	-0.0416	-0.0413	-0.0383	-0.0401	-0.8844**
Stomata frequency	G 0.6927	-0.6462	-0.0077	-0.4444	0.5312	-0.6682	0.6853	0.1188	0.6012	-0.5679	0.6060	0.3965	-0.7526	-0.7564	-0.7189	-0.6682	-0.5526	-0.6048	-0.9850**
(Abaxial)	P -0.0250	-0.0225	-0.0038	0.0060	-0.0073	0.0250	-0.0254	-0.0154	-0.0241	0.0235	-0.0242	-0.0059	0.0264	0.0269	0.0257	0.0252	0.0232	0.0244	-0.8586**
Stomata length	G -0.3748	-0.3836	-0.0107	0.0759	-0.1534	0.3644	-0.3808	-0.2256	-0.3565	0.3486	-0.35780	-0.1200	0.3847	0.3832	0.3916	0.3866	0.3703	0.3694	-0.8760**
(µm) (Adaxial)	P 0.0174	-0.0164	0.0034	-0.0032	0.0063	-0.0170	0.0178	0.0096	0.0168	-0.0164	0.0168	0.0048	-0.0182	-0.0182	-0.0190	-0.0182	-0.0171	-0.0176	-0.8675**
Stomata length	G -0.6620	-0.6795	-0.0727	0.1020	-0.2711	0.6246	-0.6792	-0.4254	-0.6244	0.6135	-0.6256	-0.2247	0.6653	0.6558	0.6817	0.6905	0.6849	0.6595	-0.9381**
(µm) (Abaxial)	P -0.0357	-0.0333	-0.0071	0.0049	-0.0139	0.0350	-0.0381	-0.0220	-0.0351	0.0344	-0.0351	-0.0110	0.0372	0.0368	0.0376	0.0392	0.0372	0.0370	-0.8486**
Stomata breadth	G 0.0778	-0.0827	0.0149	-0.0086	0.0357	-0.0713	0.0819	0.0456	0.0758	-0.0745	0.0759	0.0301	-0.0777	-0.0755	-0.0808	-0.0847	-0.0854	-0.0831	-0.9753**
(µm) (Adaxial)	P -0.0014	-0.0013	-0.0004	0.0001	-0.0005	0.0012	-0.0014	-0.0007	-0.0013	0.0013	-0.00013	-0.0005	0.0014	0.0014	0.0014	0.0015	0.0016	0.0015	-0.8531**
Stomata breadth	G -0.3098	-0.3078	-0.1071	0.0097	-0.0761	0.2698	-0.3021	-0.1416	-0.3076	0.3016	-0.3084	-0.1160	0.3024	0.2956	0.3052	0.3090	0.3149	0.3236	-0.9434**
(µm) (Abaxial)	P -0.8340	-0.0762	-0.0302	0.0024	-0.0202	0.0734	-0.0826	-0.0352	-0.0843	0.0823	-0.0845	-0.0283	0.0833	0.0819	0.0837	0.0852	0.0841	0.0903	-0.8720**

REFERENCES

1. Allard, R. W., Principles of plant breeding. John Wiley and Sons, Inc. New York. PP: 485 (1960).

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- Ashish, J., Nadaf, H. L. and Gangadhara, K., Genetic analysis of rust and late leaf spot in advanced generation recombinant inbreed lines of groundnut (*Arachis hypogaea* L.) *Int. J. Genet. Engg. And Biotech*, 5(2): 109 – 114 (2014).
- Azad, M. A. K. and Hamid, M. A., Genetic variability, character association and path analysis in groundnut (*ArachishypogaeaL.*). *Thailand J. Agric. Sci.*, **33(3-4):** 153-157 (2000).
- Burton, G. W., Quantitative inheritance in grasses. *Proc.6th Intl. Grassld. Cong.* 1: 227-283 (1952).
- Chari, S. R., Genetic analysis of morphological and biochemical traits in groundnut. M.Sc. (Agri.) Thesis Submitted to Acharya N.G. Ranga Agril. University, Hyderabad (2005).
- Dewey, D. R. and Lu, K. H. A., correlation and path analysis of components of crested wheat grass seed production. *Agronomy Journal*, **51**: 515-518 (1959).
- Dolma, T., Sekhar, M. R. and Reddy, K. R., Genetic variability, correlation and path analysis for yield its components in late leaf spot resistance in groundnut (*Arachis hypogaea* L.). J. Oilseeds Res., 27(2): 154-157 (2010).
- Giri, R. R., Toprope, V. N. and Jagtap, P. K., Genetic variability, character association and path analysis for yield, its component traits and late leaf spot, *Phaeoisariopsis personata* (Berk and curt), in groundnut. *Int. J. Plant Sci.* 4(2): 551 – 555 (2009).
- Gopal, K., Shaik, K. A. and Prasad, B. G., Estimation of losses due to tikka leaf spot in groundnut (*ArachishypogaeaL.*).*Legume res.*, 29(4): 289-291 (2006).
- Gouda Patil, K., Kenchanagoudar, P. V., Parameshwarappa K. G. and Salimath, P. M. A., study of correlation and path analysis in groundnut Karnataka *J. Agric. Sci.*, **19(2)**: 272-277 (2006).
- 11. John, K. and Reddy, P., Character

Association and Path Analysis Studies for Pod Yield and Its Components in Early Segregating Population of Groundnut (*Arachis hypogaea* L.). *Int. J. Curr. Res. Biosci. Plant Biol.* **2(7):** 149-157 (2015).

- Johnson, H. W., Robinson, H. I. and. Comstock, R. E., Estimates of genetic variability in soybean. *Agronomy Journal*, 47: 314-318 (1955).
- Kadam, P. S., Desai, D. T., Chinchane, V. N. and Sharma, V., Correlation and path analysis in groundnut, (*ArachishypogaeaL.*) J. Oilseed Res., 26: 63-65 (2009).
- Kahate, N. S., Toprope, V. N. and Gadakh, S. S., Correlation and path analysis for yield, morphology and biochemical traits in groundnut (*Arachis hypogea* L.) *BIOINFOLET* 11(3B): 868 870 (2014).
- Kahate, N. S. and Toprope, V. N., Genetic variability and character association in groundnut. *J. Agric. Res. Technol.*, **39(2):** 343 346 (2014).
- Kumar, C. V. S., Kumar, V. I. and Rajmani, S., Genetic variability, correlation and path analysis in groundnut. *Prog. Agric.*, 4: 69-70 (2008).
- Kaur, S., Dhillon, M. and Sokhi, S. S., Stomata characteristics of groundnut genotypes resistant susceptible to late leaf spot. *Plant Dis. Res.*, 4(4): 15-18 (1989).
- Khan, A. M. R., Khan, M. I. and Tahir, Genetic variability and criterion for the selection of high yielding peanut genotypes. *Pakistan J. Agric. Res.*, 16(1): 9-12 (2000).
- Lashmidevamma, T. N., Byre Gowada, M. and Mahadevu, P., Character association and path analysis in groundnut (*ArachishypogaeaL.*). *Mysore J. Agric. Sci.*, 38(2): 221-226 (2004).
- Li Dun, Influence of multiple biochemical compounds on resistance of peanut plant to rust. J. South China Agric. Univ., 17(2): 44-49 (1996).
- Mahalakshmi, P., Manivannan, N. and Murlidharan, V., Variability and correlation studies in groundnut (*Arachishypogaea L.*). *Legume res.*, 28(3): 194-197 (2005).
- 22. Nandini, C., Savithramma, D. L. and Naresh

Rathod and Toprope

Babu, N., Genetic variability analysis for water use efficiency in F8 recombinant inbreed lines of groundnut (*Arachis hypogaea* L.) *CurrentBiotica*, **5(3):** 282 – 288 (2011).

- 23. Padmaja, D., Rao, B., Eswari, K. B. and Reddy, S. M., Genetic variability, heritability for late leaf spot tolerance and productivity traits in a recombinant inbreed line population of groundnut (*Arachis hypogaea* L.). *J. Agric. And Veter. Sci.*, 5(1): 36–41 (2013).
- Panse, V. G. and Sukhtame, P. V., Statistical methods of *agricultural research workers New Delhi ICAR Publication* (2nd Ed.). PP: 361 (1961).
- 25. Patil, A. S., Punewar, A. A., Nandwar, H. R and. Shah, K. P., Estimation of variability parameters for yield and its component traits in groundnut (Arachis hypogaea L.).*The bioscan.* **9(2)**: 749-754 (2014).
- Rao, V. T., Venkanna, V., Bhadru, D. and Bharathi, D., Studies on variability, character association and path analysis on Groundnut (*Arachishypogaea* L.) *Inter. J. Pure App.Biosci*, 2 (2): 194-197 (2014).
- 27. Sawargaonkar, S. L., Giri, R. R. and Hudge, B. V., Character association and path analysis of yield component traits and late leaf spot disease traits in groundnut (Arachis hypogaea L.). *Agricultural Science Digest.* 30(2): 115 119 (2010).
- 28. Sharma, H. and Dashora, A., Character association and path analysis in groundnut, *Arachishypogaea*L. *J. Oilseed Res.*, **26**: (2009).
- 29. Subrahmanyam, P., Moss, J. P., Mcdonald,

D., Rao, P. V. S. and Rao, V. R., Resistance to leaf spot caused by *Cercosporidium personatum* in wild *Arachis* species. *Plant Disease.* **69:** 951-954. (1985).

- Shukla, A. K. and Rai, P. K., Evaluation of Groundnut Genotypes for Yield and Quality Traits. *Annals Plant and Soil Res.* 16(1): 41-44 (2014).
- 31. Suneetha, K., Dasarada Rami Reddy, D. R. C. and Ramana, J. V., Genetic variability character association groundnut. National Symposium: Enhancing Productivity of Groundnut for Sustaining *Food and Nutritional Security. NRCG, Junagadh.* Oct. 11-12. (2004).
- Vasanthi, R. P., Vishnuvardhan, K. M., Reddy, K. H. P. and Reddy, B. V. B., Genetic variability studies for yield attributes and resistance to foliar diseases in groundnut (*Arachis hypogaea* L.)*Inter. J. Appli. Biology and Pharma. Tech.*, 3(1): 390 – 394 (2012).
- 33. Wani, S. C., Deshmukh, S. N. and Satpute, G. N., Variability in advance generation lines of groundnut (Arachis hypogaea L). National Enhancing Productivity Symposium: of Groundnut for *Sustaining* Food and Nutritional Security. NRCG. Junagadh October 11-12 (2004).
- 34. Zaman, M. A., Tuhina-Khatun, M., Ullah, M. Z., Moniruzzamn, M. and Alam, K. H., Genetic variability and path analysis of groundnut (*Arachishypogaea L.*). A Sci. J. Krishi Found., 9(1&2): 29-36 (2011).