

Journal of Pharmacognosy and Phytochemistry

Available online at www.phytojournal.com



E-ISSN: 2278-4136 P-ISSN: 2349-8234 www.phytojournal.com

JPP 2020; 9(5): 359-364 Received: 01-04-2020 Accepted: 02-05-2020

TJ Bhor

All India Network Project on Jute and Allied Fibres, Mahatma Phule Krishi Vidyapeeth, Rahuri, Maharashtra, India

DP Pacharne

All India Network Project on Jute and Allied Fibres, Mahatma Phule Krishi Vidyapeeth, Rahuri, Maharashtra, India

RS Wagh

All India Network Project on Jute and Allied Fibres, Mahatma Phule Krishi Vidyapeeth, Rahuri, Maharashtra, India

Corresponding Author: TJ Bhor All India Network Project on Jute and Allied Fibres, Mahatma Phule Krishi Vidyapeeth, Rahuri, Maharashtra, India

Genetic variability, correlation and path analysis studies in Jute (*Corchorus olitorius*) germplasm lines

TJ Bhor, DP Pacharne and RS Wagh

Abstract

Fifty two accessions of tossa jute germplasm were evaluated for variability. The highest heritability (h²), genotypic coefficient of variation (GCV) and percent genetic advance over mean were observed for the traits green biomass and fibre weight per plant indicated preponderance of additive gene action. Green biomass weight per plant found to be highest (0.85) genotypic correlation with fibre yield per plant. The traits plant height, base diameter, green biomass weight, days to 50% flowering and fibre weight were found associated with each other. Path coefficient analysis showed that green biomass weight has direct effect on fibre yield per plant. The traits plant height, basal diameter and days to 50% flowering had also indirect effect via green biomass yield on fibre yield per plant. Thus, the result of this investigation suggested that green biomass weight, plant height, basal diameter and days to 50% flowering would be the selection parameters to produce jute varieties with acceptable yield.

Keywords: Heritability, genotypic coefficient of variation, genetic advance, direct effect, improvement, selection

Introduction

Jute (Corchorus sp.) is grown as fibre crop during pre-kharif season and plays an important role in Indian economy. West Bengal alone contributes to 77% of the total Indian Jute Sinha et al. (2004) ^[34]. Jute is one of the most important bast fibre crop of the world being cultivated in Eastern India, Bangladesh, Nepal and some South East Asian countries, and is second only to cotton in terms of production and variety of uses (Islam et al., 2013) ^[15]. Entire life-cycle of jute from cultivation to usage disposal, it is friendly to the environment and produces no toxic material (Sarkar, 2008) [28]. Jute is still the leading cash crop and an important tradable and foreign exchange-earning (4.82%) commodity of Bangladesh. The jute fibre is used, as a raw material, for several products like hessians, sacs, and carpet backings. The two species of the genus Corchorus, which are cultivated as jute crop include C. capsularis (white jute) and C. *olitorius* (tossa jute), each with 2n = 14, are cultivated in our country, but *Corchorus olitorius* is predominantly grown by the farmers. Due to increase the environmental consciousness the demand of natural fibre increases rapidly through the world. To meet this increasing demand there is an urgent need to redesign the ongoing breeding strategies to improve both the yield and quality of jute. Information on the nature and magnitude of genetic variability for the desired characters in the base material and interrelationship among them is useful in breeding for high yield (Kumar et al., 2008) ^[19]. It is therefore, essential to understand the variability, path analysis for prediction of direct and indirect effect of characters on fibre yield in the available germplasm of jute. Increased pressure has been exerted on jute scientists to develop varieties with high yield potential and superior grade fibre that meet the demands of the textile industries. The use of plant genetic resources in breeding research is largely dependent on the available information of their genetic variability. The rates of genetic exploitation of the variation for production of improved cultivars and yield have not received sufficient attention. Rapid genetic improvement of crop depends on the availability of sufficient genetic diversity, which could be selected and combined in various forms to produce reasonable improvement (Denton and Nwangburuka, 2011; Wani et al. 2011) [9, 35]. Estimates of heritability values become more reliable and meaningful when combined with genetic advance (Denton and Nwangburuka, 2011)^[9]. High heritability and genetic advance in a trait indicates the presence of additive genes in such trait and further suggest reliable crop improvement through selection of such a trait (Panse, 1957) [24]. In this study fifty two accessions of tossa jute (Corchorus olitorius) germplasm were used to find out the superior genotypes and presence of variability among them. This study also includes the correlation and path coefficient between yield and its

attributing traits and to find out the extent of direct and indirect effects of fibre yield components.

Materials and Methods

The experiment was conducted at the All India Network Project on Jute and Allied Fibres, Mahatma Phule Krishi Vidyapeeth, Rahuri during the Kharif 2018. Fifty two accessions of tossa jute (Corchorus olitorius) which collected from AINP jute and allied fibres via Central Research Institute for Jute and Allied Fibres (CRIJAF), Barrackpore, West Bengal, India along with two check variety JRO-524 and JRO-204 were taken for this study. The experiment was laid out in the Randomized Block Design (RBD) with three replications. Seeds were sown in single row of 6m long. Spacing was 30 cm between rows, 5-7cm between plants and 1 m between blocks. Standard production technology from Hand Book on Agricultural Technologies of Jute, Kenaf and Mesta crops (2008) ^[17] was adopted to raise a good crop under optimum management. The recommended agronomic practices were followed to obtain optimum fibre yield. Except days to 50% flowering, which was studied on plot basis, observations on other four quantitative characters namely plant height (cm), base diameter (cm), green weight (g/plant), days to 50% flowering and fibre weight (g/plant) were recorded from ten plants randomly selected from each genotype from each replication. The analysis of variance (ANOVA) for RBD was estimated according to Panse and Sukhtame (1989) ^[25]. The genotypic and phenotypic variances were calculated according to Johnson et al. (1955) [16] and Comstock and Robinson (1952)^[5]. Genotypic coefficient of variation (GCV) and phenotypic coefficient of variation (PCV) were calculated by the method suggested by Singh and Chaudhary (1985) ^[31] whereas heritability in broad sense for yield and its components were worked out by using formula suggested by Hanson et al. (1956) [13]. Genetic advance (GA) was calculated by the method suggested by Johnson et al. (1955) ^[16]. Genotypic and phenotypic correlations were partitioned using the technique outlined by Dewey and Lu $(1959)^{[10]}$.

Results and Discussion

The analysis of variance revealed highly significant differences among genotypes for all the characters indicating presence of considerable variability for the characters under study among the test genotypes (Table 1). This suggests that there was genetic diversity in these characters among the accessions studied and further suggests prospects for meaningful selection for Corchorus improvement. This agree with the reports of Dar and Sharma (2011)^[7], who reported significant variation in quantitative characters in 60 tomatoes genotypes, Kitila et al. (2011) ^[18] reported significant variation in quantitative traits in coffee; Singh et al. (2011) reported significant variations quantitative characters in field pea, Nwangburuka et al. (2012) [23] reported significant variations in nine yield related characters in twenty-nine accessions of okra. Bhandari et al. (2017) [3] reported significant variations in ten yield related characters in 66 genotypes of tomato. Biswas et al. (2018) [4] reported significant variations in seven yield related traits in tossa jute. The mean performance of individual genotype is presented in Table 2. The different genetic parameters like range, mean, phenotypic variance, genotypic variance, phenotypic coefficient of variation, genotypic coefficient of variation, heritability, genetic advance and genetic advance as a percentage of mean for each character are presented in Table 03. The plant height at harvest (113 days) ranged from 190-285cm. The highest score of plant height was recorded in accession JRO-524 (285cm/plant) which was followed by accessions JRO-204 (283cm/plant), OIN-115 (280cm/plant), OIN-124 (275cm/plant) and OIN-110 (272cm/plant). Base diameter was ranged from 1.15 - 1.63 cm. The highest score of base diameter was recorded in accession OIN-186 (1.63cm/plant) which was followed by accessions OIN-124 (1.62 cm/plant), OIN-87, OIN-109, JRO-524 (1.61 cm/plant), JRO-204 (1.59 cm/plant) and accession OIN-110, OIN-111, OIN-115 and OIN-122 (1.58 cm/plant). Green biomass was ranged from 61-260 g. The highest score for green biomass weight was recorded in accession JRO-524 (260 g/plant) which was followed by accessions JRO-204 (235 g/plant), OIN-124 (232g/plant) and OIN-86 OIN-110 (220 g/plant). Days to 50% flowering ranged from 44-70 days. The highest score for days to 50% flowering was recorded in accession JRO-524 (70 days) which was followed by accessions JRO-204 (68 days), OIN-105 (60 days), OIN-134 (58 days) and OIN-109, OIN-115 (54 days). Fibre weight was ranged from 3.73 – 13.76 g/plant. It was highest in accession JRO-524 (13.76 g/plant) which was followed by accessions JRO-204(13.54 g/plant), OIN-115 (12.75 g/plant), OIN-86 (11.59 g/plant accession OIN-110 (10.80 g/plant) and OIN-100 (10.0 g/plant). Based on major yield contributing characters six accessions, viz., JRO-524, JRO-204, OIN-115, OIN-86, OIN-110 and OIN-100 performed relatively better in most of the cases than the studied materials. Hence these accessions may be selected for further study in the next season.

The PCV value was found slightly higher than the GCV value for all the characters. This may suggest that slight environmental effect on the phenotype of all the other characters. This report corresponds to the report of Denton and Nwangburuka (2011) ^[29], Nwangburuka *et al.* (2012) ^[23], Mohammed *et al.* (2012) ^[20], Yadav *et al.* (2011) ^[36] and Ayalneh *et al.* (2012) ^[1] who observed slight differences between PCV and GCV in characters studied in *Solanum anguivi*, okra, Ethiopian Duram wheat, Rice and in Tef, respectively.

The PCV ranged from (9.42) for days to 50% flowering to (35.61) for green weight (g)/plant. Similarly, GCV ranged from (9.14) for days to 50% flowering to (33.52) for green weight (g)/plant. The high PCV and GCV values recorded in green weight (g)/plant suggests that this character accounts for most of the variation recorded in C. olitorius. The phenotypic and genotypic variance was the highest in green weight (g)/plant (2594.57 and 2298.89) but the lowest in the basal diameter (cm) (0.0241 and 0.0238) and fibre weight (g)/plant (6.48 and 5.72), respectively. High genotypic variance facilitates selection for the improvement and widens the probability for heritability of traits from parents to offspring (Ayalneh et al. 2012)^[1]. Broad-sense heritability estimates were high for all the charcaters. Dabholkar (1992) ^[6] categorized heritability estimates as low (5-10%), medium (11-30%) and high (>30%). Maximum estimates of broadsense heritability was recorded in basal diameter (cm) (98.90) followed by days to 50% flowering (94.19), green weight (g)/plant (88.60), fibre yield (88.27) and plant height (88.17) This suggests the effect of additive genes in the inheritance of these characters. According to Ghandi et al. (1964) [11] and Ibrahim and Hussein (2006) ^[14], prediction of the response of an individual to selection are more reliable when GCV, estimates of broad-sense heritability and genetic advance is combined instead of relying on the estimates of broad-sense heritability estimates alone. When the GA is high the heritability is mainly due to additive gene effect (Percy and Turcotte, 1991) ^[26]. Hence in present investigation, selection based on characters such as green weight (g)/plant (33.52, 88.60 and 64.99) and fibre weight (g)/plant (33.60, 88.27 and 64.46) combining high GCV, heritability and GA, respectively are under additive gene effect and will be effective in accurate prediction of yield. This agree with the reports of Bello *et al.* (2006) ^[2] in Okra and Mohammed *et al.* (2012) ^[20] in their work with wheat; Yadav *et al.* (2011) ^[36] in their report on rice and Biswas *et al.* (2018) ^[4] in tossa jute. High heritability and genetic advance is an indication of how much selection to improve a character can be based of phenotypic performance (Johnson *et al.*, 1955) ^[16].

Estimates of phenotypic and genotypic correlation coefficients between each pair of characters are given in Table 4. The results showed that the magnitude of genotypic correlation is higher than the phenotypic correlation indicating that elimination of environmental effects led to strengthen genetic association. The correlation analysis revealed that there was significant and positive association of green biomass weight, plant height, basal diameter and days to 50% flowering with fibre yield per plant The traits plant height, basal diameter, green weight, days to 50% flowering were also associated with each other both at genotypic and phenotypic level.. Similar result was obtained by Singh *et al.* (2013) ^[33], Nayak and Baisakh (2008) ^[22] and Mostofa *et al.* (2002) ^[21].

The results of path coefficient analysis are given in Table 5. Green weight (0.88) had highest positive direct effect (genotypic) on fibre yield per plant followed by days to 50% flowering (0.12) and plant height (0.04). This findings is in agreement with Senapati *et al.* (2006) ^[30]. Therefore direct selection based on these characters would be feasible. Base

diameter per plant exhibited negative direct effects towards fibre yield, but its significant positive correlations with fibre yield per plant indicated that the indirect selection through green weight, plant height and days to 50% flowering could be made for high yielding jute genotypes as they having positive indirect effects. Similar findings were reported by Pervin et al. (2012) [27]. The residual effect (R) was 0.48, indicating there was also some other characters which although not studied but influenced the yield of fibre per plant. The green weight per plant of the present study had higher values of direct effects even than their respective correlation coefficients indicating their prime importance in fibre yield. Sawarkar et al. (2014) ^[29] also suggested that green weight per plant and plant height would be an important parameter in predicting fibre yield in jute. The traits plant height, basal diameter and days to 50% flowering had also indirect effect via green biomass yield on fibre yield per plant. Thus, the result of this investigation suggested that green biomass weight, plant height, basal diameter and days to 50% flowering would be the selection parameters to produce jute varieties with acceptable yield.

Declaration

The authors declare no conflict of interest.

Acknowledgment

The authors are thankful to the Network Co-ordinator, All India Network Project on Jute and Allied Fibres, Kolkatta for providing the germplasm lines of jute for evaluation and the financial assistance to conduct the experiment. The financial assistance received from Mahatma Phule Krishi Vidyapeeth, Rahuri and Government of Maharashtra, for the study is gratefully acknowledged.

Table 1: Results of analysis of variance (ANOVA) for Jute traits

Sauraa of		Mean sum of squares						
variation DF		Plant height (cm)	Basal diameter (cm)	Green biomass plant ⁻¹ (g)	Days for 50% flowering	Fibre yield plant ¹ (g)		
Replication	2	278.56	0.001	244.55	73.90	0.55		
Treatments	51	2097.69**	0.072**	7192.37**	66.91**	17.93**		
Errors	102	89.77	0.0003	295.68	1.35	0.76		

Sr.	Constras	Plant height	Basal diameter	Green biomass plant ⁻	Days for 50%	Fibre yield plant ⁻
No.	Genotypes	(cm)	(cm)	$^{1}(g)$	flowering	¹ (g)
1.	OIN 86	269	1.63	220	51	11.59
2.	OIN 87	268	1.61	204	51	8.40
3.	OIN 88	249	1.44	144	53	6.39
4.	OIN 89	200	1.21	86	46	4.08
5.	OIN 90	243	1.36	145	45	5.79
6.	OIN 91	234	1.28	120	44	6.40
7.	OIN 92	190	1.17	61	50	6.91
8.	OIN 93	226	1.24	112	50	6.01
9.	OIN 94	194	1.21	88	51	7.21
10.	OIN 95	203	1.15	73	52	3.63
11.	OIN 96	200	1.17	79	49	3.73
12.	OIN 97	210	1.18	84	50	7.80
13.	OIN 98	210	1.22	104	52	5.62
14.	OIN 99	209	1.18	84	53	4.82
15.	OIN 100	252	1.56	192	53	10.00
16.	OIN 101	218	1.27	136	45	8.00
17.	OIN 102	196	1.22	88	45	4.80
18.	OIN 103	209	1.30	108	52	5.22
19.	OIN 104	266	1.55	194	52	9.32
20.	OIN 105	218	1.24	104	60	5.22
21.	OIN 106	206	1.24	92	48	3.77

 Table 2: Mean performance of fibre yield and yield contributing characters

22.	OIN 107	227	1.31	136	50	6.41
23.	OIN 108	219	1.29	124	50	5.60
24.	OIN 109	268	1.61	144	54	6.61
25.	OIN 110	272	1.58	220	53	10.80
26.	OIN 111	267	1.58	202	51	9.60
27.	OIN 112	220	1.31	108	45	9.02
28.	OIN 113	240	1.42	132	50	6.42
29.	OIN 114	242	1.54	164	50	6.97
30.	OIN 115	280	1.58	215	54	12.75
31.	OIN 116	264	1.55	192	51	8.41
32.	OIN 117	221	1.37	140	50	5.81
33.	OIN 118	208	1.21	84	47	3.62
34.	OIN 119	235	1.25	136	45	6.41
35.	OIN 120	266	1.50	184	49	9.99
36.	OIN 121	252	1.39	176	51	8.01
37.	OIN 122	266	1.58	205	51	9.29
38.	OIN 123	223	1.25	114	51	6.50
39.	OIN 124	275	1.62	232	53	9.28
40.	OIN 125	210	1.31	113	51	4.83
41.	OIN 126	225	1.31	128	50	5.20
42.	OIN 127	228	1.34	136	52	5.45
43.	OIN 128	244	1.36	140	50	6.40
44.	OIN 129	261	1.55	184	53	8.41
45.	OIN 130	240	1.40	156	52	8.00
46.	OIN 131	230	1.39	144	50	6.39
47.	OIN 132	245	1.42	165	50	7.62
48.	OIN 133	235	1.30	136	51	6.81
49.	OIN 134	230	1.28	128	58	6.01
50.	OIN 135	207	1.18	86	48	4.88
51.	JRO 524©	285	1.61	260	70	13.76
52.	JRO 204©	283	1.59	235	68	13.54
	Mean	235	1.37	143	51	7.18
	S.E. + at 5%	5.47	0.009	9.94	0.67	0.50
	C.D. + at 5%	15.35	0.026	27.87	1.88	1.41
	C.V.	4.03	1.18	12.03	2.27	12.14

Table 3: Range, variability, heritability and other genetic parameters

Characters	Mean	Range	GV	PV	GCV	PCV	H^2	GA	%GA
Plant height (cm)	235.0	190-285	669.31	759.08	10.99	11.71	88.17	50.04	21.26
Basal diameter (cm)	1.37	1.15-1.63	0.0238	0.0241	11.24	11.30	98.90	0.32	23.02
Green biomass plant ⁻¹ (g)	143.0	61-260	2298.89	2594.57	33.52	35.61	88.60	92.97	64.99
Days for 50% flowering	51.0	44-70	21.85	23.20	9.14	9.42	94.19	9.34	18.29
Fibre yield plant ⁻¹ (g)	7.18	3.73-13.76	5.72	6.48	33.30	35.45	88.27	4.63	64.46

GV-Genotypic Variance, PV- Phenotypic Variance, GCV- Genotypic Coefficient of Variation, PCV- Phenotypic Coefficient of Variation, H²-Heritability, GA- Genetic Advance, % GA- Genetic Advance over Percentage of Mean.

Table 4: Phenotypic (P)	and Genotypic (G)	correlation coefficient	among different yield component
-------------------------	-------------------	-------------------------	---------------------------------

Characters	Correlation coefficient	Basal diameter (cm)	Green biomass plant ⁻¹ (g)	Days for 50% flowering	Fibre yield plant ⁻ ¹ (g)
Dlanthaisht (ana)	Р	0.89**	0.91**	0.44**	0.77**
Plant height (cm)	G	0.96**	0.97**	0.48**	0.83**
Basal diameter (cm)	Р		0.90**	0.40**	0.75**
	G		0.95**	0.42**	0.79**
Green biomass plant-1	Р			0.47**	0.87**
(g)	G			0.49**	0.85**
Days for 50%	Р				0.49**
flowering	G				0.52**

*, ** significant at 5% and 1% respectively.

Table 5: Path coefficient (Genoty)	bic) analysis showing direct (Bo	old) and indirect effects of component traits in jut	Э
------------------------------------	----------------------------------	--	---

Characters	Plant height (cm)	Basal diameter (cm)	Green weight plant ⁻ ¹ (g)	Days for 50% flowering	Correlation with fibre yield
Plant height (cm)	0.04	-0.14	0.87	0.06	0.83**
Basal diameter (cm)	0.04	-0.14	0.85	0.05	0.79**
Green weight plant ⁻¹ (g)	0.04	-0.14	0.88	0.06	0.85**
Days for 50% flowering	0.02	-0.06	0.44	0.12	0.52**

Residual effect = 0.50

Table 6: Path coefficient (Phenotypic) analysis showing direct (Bold) and indirect effects of component traits in jute.

Characters	Plant height (cm)	Basal diameter (cm)	Green weight (qt./ha)	Days for 50% flowering	Correlation with fibre yield
Plant height (cm)	0.007	-0.13	0.86	0.05	0.77**
Basal diameter (cm)	0.006	-0.15	0.85	0.04	0.74**
Green weight (qt./ha)	0.006	-0.14	0.95	0.05	0.86**
Days for 50% flowering	0.002	-0.06	0.44	0.10	0.49**

Residual effect = 0.48



Fig 1: Genotypical path Diagram for Fiber yield g/plant

References

- Ayalneh T, Habtamu Z, Amsalu A. Genetic variability, heritability and genetic advance in tef (Eragrostis tef (Zucc.) Trotter) lines at sinana and adaba. Intern. J of Pl. Breed. & Genet. 2012; 6:40-46.
- Bello D, Sajo AA, Chubado D, Jellason JJ. Variability and correlation studies in okra (*Abelmoschus esculentus* L. Moench.). J. of Sustainable Develop.in Agric. and Environ. 2006; 2:120-126.
- 3. Bhandari HR, Kartikeya S, Eswar RG. Genetic variability, heritability and genetic advance for yield traits in tomato (*Solanum lycopersicum* L.). Intern. J. of Curr. Microbi. and Appl. Sci. 2017; 6(7):4131-4138.
- Biswas SK, Islam SN, Sarker DH, Moniruzzaman, Tareq Z. Genetic variability, heritability and genetic advance for yield related characters of tossa jute (*Corchorus olitorius*) genotypes. J. Biosci. Agric. Res. 2018; 17(01):1416-1421
- 5. Comstock RE, Rabinson HF. Genetic parameter, their estimation and significance. In: in proceed. on 6th Intern.Grassland Congress. 1952, 284-291.
- Dabholkar AR. Elements of biometrical genetics. Concept Publishing Company, New Delhi, India. 1992, 138-140.
- Dar RA, Sharma JP. Genetic variability studies of yield and quality traits in tomato (*Solanum lycopersicum* L.). Intern. J. of Pl. Breed. & Genet. 2011; 5:168-174.
- 8. Das A, Kumar D. Genetic divergence and character association for yield and quality attributing characters in

tossa jute (*Corchorus olitorius* L.). Electron.J. of Pl. Breed. 2016; 7(3):529-537.

- Denton OA, Nwangburuka CC. Heritability, genetic advance and character association in six yield related characters of *Solanum anguivi*. Asian J. of Agric. Res. 2011; 5:201-207.
- 10. Dewey DR, Lu KH. A correlation and path coefficient analysis of components of crested wheat grass seed production. Agron. J. 1959; 51:515-518.
- 11. Ghandi SM, Sanghai AK, Nathawat KS, Bhatnagar MP. Genotypic variability and correlation coefficient to grain yield and a few other quantitative characters in Indian wheat. Indian J. of Pl. Breed. & Genet. 1964; 24:1-8.
- 12. Hand Book on agricultural technologies of jute, kenaf and mesta crops Bangladesh Jute Research Institute. Manik Mia Avenue, Dhaka-1207, 2008.
- 13. Hanson CH, Robinson HF, Comstock RE. Biometrical studies of yield in segregating population of Korean lespedeza. Agron. J. 1956; 48:268-272.
- 14. Ibrahim MM, Hussein RM. Variability, heritability and genetic advance in some genotypes of roselle (*Hibiscus sabdariffa* L.) World J. of Agric.Sci. 2006; 2:340-245.
- 15. Islam S, Azam MS, Sazia S, Abu AS, Maksudul A, Shamim R, *et al.* Improved salt tolerance of jute plants expressing the kat E gene from *Escherichia coli*. Turkish J. of Biology. 2013; 37:206-211.
- Johnson HW, Robinson HF, Comstock RE. Estimates of genetic and environmental variability in soybean. Agron. J. 1955; 47:314-318.

- 17. Kenaf, mesta crops Hand Book on agricultural technologies of jute, Bangladesh Jute Research Institute. Manik Mia Avenue, Dhaka-1207. 2008.
- 18. Kitila O, Alamerew S, Kufa T, Garedew W. Variability of qualitative traits in limmu coffee (*Coffea arabica* L.) in Ethiopia. Intern. J. of Agric. Res. 2011; 6:482-493.
- 19. Kumar D, Agarwal RC, Begum T. Analysis for identification of distinct and uniform extant jute (*Corchorus olitorius* L. and *C. capsularis* L.) varieties. Seed Res. 2008; 36:121-134.
- Mohammed A, Geremew B, Amsalu A. Variation and associations of quality parameters in Ethiopian duram wheat (*Triticum turgidum* L. var. duram) genotypes. Intern. J. of Pl. Breed. & Genet. 2012; 6:17-31.
- 21. Mostofa MG, Islam MR, Morshed Alam ATM, Mahbub Ali SM, Mollah MAF. Genetic variability, heritability correlation studies in Kenaf (*Hibiscus cannabinus* L). Onl. J. Biol. Sci. 2002; 2(6):422-424.
- 22. Nayak BK, Baisakh B. Character association and path analysis in tossa jute (*Corchorus olitorius* L.). Environ Ecol. 2008; 26(1A):361-363.
- 23. Nwangburuka CC, Denton OA, Kehinde OB, Ojo DK, Popoola AR. Genetic variability and heritability studies in cultivated okra (*Abelmoschus esculentus* L. Moench.), *Span.* Intern. J. of Agric. Res. 2012; 10:123-129.
- 24. Panse VG. Genetics of qualitative characters in relation to plant breeding. Indian J. of Genet. and Pl. Breed. 1957; 17:318-328.
- 25. Panse VG, Sukhatme PV. Statistical Methods for Agricultural Workers. Indian Council of Agricultural Research, New Delhi, India, 1989.
- 26. Percy RG, Turcotte EL. Inheritance of male-sterile mutant ms13 in American Pima cotton. Crop Sci. 1991; 31:1520-1521.
- 27. Pervin N, Haque GKMN. Path coefficient analysis for fibre yield related traits in deshi jute (*Corchorus capsularis* L.). IRJALS. 2012; 1(3):72-77.
- Sarkar S. Good practices for jute and allied fibre crops. In: Souvenir, International Symposium on jute and allied fibre Production, Utilization and Marketing, Organized by Indian Fibre Society (Eastern Region). Kolkata, 9-12 January. 2008, 1-3.
- 29. Sawarkar A, Yumnam S, Patil SG, Mukherjee S. Correlation and path coefficient analysis of yield and its attributing traits in tossa jute (*Corchorus olitorius* L.). The Bioscan, 2014; 9(2):883-887.
- 30. Senapati S, Nasim Ali MD, Sasmal BG. Genetic variability, heritability and genetic advance in *Corchorus* sp. Environ. Ecol. 2006; 24S(1):1-3.
- Singh RK, Chaudhury BD. Biometrical methods of quantitative genetic analysis. Kalyani Publishers, New Delhi, India, 1985.
- 32. Singh A, Singh S, Babu JDP. Heritability, character association and path analysis studies in early segregating population of field pea (*Pisum sativum* L. var. arvense). Intern. J. of Pl. Breed. and Genet. 2011; 5:86-92.
- 33. Singh CM, Babu GS, Binod K, Mehandi S. Analysis of quantitative variation and selection criteria for yield improvement in exotic germplasm of upland rice (*Oryza sativa* L.). Bioscan. 2013; 8(2):485-492.
- Sinha MK, Sengupta D, Sen HS, Ghosh T. Jute and jutelike fibres: current situation. Sci. Cult. 2004; 70(1, 2):32-37.
- 35. Wani BA, Ram M, Yasin A, Singh E. Physiological traits in integration with yield and yield components in wheat

http://www.phytojournal.com

(*Triticum aestivum* L.) study of their genetic variability and correlation. Asian J. of Agric. Res. 2011; 5:194-200.

36. Yadav SK, Pandey P, Kumar B, Suresh BG. Genetic architecture, inter-relationship and selection criteria for yield improvement in rice (*Oryza sativa* L.). Pakistan J. of Biolo. Sci. 2011; 14:540-545