

SABRAO Journal of Breeding and Genetics 53 (2) 157-171, 2021

COMPARATIVE PERFORMANCE AND GENETIC ATTRIBUTES OF UPLAND COTTON GENOTYPES FOR YIELD-RELATED TRAITS UNDER OPTIMAL AND DEFICIT IRRIGATION CONDITIONS

J. SHAVKIEV^{1*}, A. AZIMOV¹, S. NABIEV¹, S. KHAMDULLAEV¹, B. AMANOV², M. KHOLIKOVA², H. MATNIYAZOVA¹ and U. YULDASHOV¹

¹Institute of Genetics and Plants Experimental Biology, Academy of Sciences of Uzbekistan, Tashkent Region, Kibray district, Yukori-Yuz, Uzbekistan

²Department of Biology, Chirchik State Pedagogical Institute, Tashkent Region, Chirchik City, Amir Temur, Uzbekistan

*Corresponding author email: jaloliddinshavkiev1992@gmail.com Email addresses of coauthors: azimov.abdulahat@bk.ru, m.saydigani@mail.ru, shuxratxamdullayev@mail.ru, amanov.81@bk.ru, holiqova19811@mail.ru, matniyazova@mail.ru, utkirbekyuldashov_1991@mail.ru

SUMMARY

Water scarcity is a stress factor that drastically affects and reduces crop productivity. The objective of this study was to explore the influence of water deficiency on seed cotton yield and its associated traits and to study the coefficient of variances, heritability, and GG for various traits in upland cotton (*Gossypium hirsutum* L). Four cotton cultivars, i.e., Ishonch, Navbakhor-2, C-6524, and Tashkent-6, and their F1 diallel hybrids were grown in a randomized complete block design with factorial arrangement and four replications under optimal and deficit irrigation conditions during 2018-2019 in the Tashkent region of Uzbekistan. Significant ($P \le 0.01$) differences were observed among the parental genotypes and their F1 hybrids for boll weight, 100-seed weight, seed cotton yield, and lint%. The genotypes exhibited significant ($P \le 0.05$) and nonsignificant differences for bolls per plant under water deficit and optimal irrigation conditions, respectively. Overall, deficit irrigation conditions negatively affected the yield-contributing traits and eventually the seed cotton yield. The parental cultivars Ishonch and Navbakhor-2 and their F1 diallel hybrids were found to be more stable and performed better than the other genotypes under both water regimes. On average, the genotypic and phenotypic variances for various traits were greater under water deficit conditions than under the optimal irrigation regime. Broad-sense heritability estimates were the highest for boll weight, 100-seed weight, seed cotton yield, and lint% but were low for bolls per plant. Seed cotton yield revealed highly significant positive associations with all traits under optimal irrigation and water deficit conditions but not with 100-seed weight under deficit conditions. The parental genotypes and their hybrids can be used as source materials for developing genotypes with drought tolerance.

Keywords: Genetic variability, coefficient of variances, heritability, genetic gain, correlation coefficient, upland cotton (*Gossypium hirsutum* L.)

Key findings: Overall, the deficit irrigation condition negatively affected yield-contributing traits and eventually seed cotton yield. Two parental cultivars, viz., Ishonch and Navbakhor-

2, and their F1 diallel hybrids were found to be more stable and performed better for various traits than other genotypes under optimal and deficit irrigation conditions.

Manuscript received: December 10, 2020; Decision on manuscript: January 25, 2021; Accepted: February 12, 2021. © Society for the Advancement of Breeding Research in Asia and Oceania (SABRAO) 2021

Communicating Editor: Dr. Naqib Ullah Khan

INTRODUCTION

Cotton (*Gossypium hirsutum* L.) is the most important economic crop grown for textile fibers in more than 70 countries worldwide. Cotton fibers and their by-products have dominated the economy of developing and developed countries (Khokhar *et al.*, 2017). Promising cotton genotypes show only enhanced seed cotton yield and lint (%), and improving fiber quality will eventually improve the cotton trade worldwide (Amanov *et al.*, 2020).

Water deficit conditions affect and decrease the total plant stature by up to 35% and leaf area index by 8% due to the interception of solar radiation (Pettigrew, 2004). Saranga et al. (1991) observed that the change in water potential accounts for 87% of the change in plant height. The combined phenomenon of abiotic stress and growth reduction affects seed cotton yield and fiber quality (Pettigrew, 2004; Babar et al., 2009; Shavkiev et al., 2019a, Marguba et al., 2020). In cotton, boll formation and fiber development are the most critical stages that are mainly affected by drought stress (Radin et al., 1992; Tohir et al., 2018). Krieg (2000) revealed that the flowering period is the most critical period to drought stress. In cotton, drought tolerance is genetically controlled through various physiological processes (Ahmad et al., 2011; Shavkiev et al., 2019b). Cotton cultivars display varied genetic responses as reflected by their morphophysiological and biochemical parameters under water deficit environments (Hinze et al., 2012; Patil et al., 2017; Jaloliddin et al., 2020).

Locally adapted germplasm can be used for the development of new genotypes through different breeding programs. Population structures are of great importance for the utilization of genetic diversity in the development of superior cultivars that combine the positive traits conditioned by diverse cotton germplasm. The majority of cotton cultivars have been developed under irrigation conditions and have experienced intensive selection to increase seed cotton yield and desirable fiber quality (Rosenow et al., 1983). Xamidov (2019) reported that cotton crops need an adequate amount of water for normal growth and development. Under irrigation water shortage, the crop experiences severe water deficiency, thereby exhibiting reductions in crop yield (Nabiev et al., may become 2020). This situation increasing critical due to the excessive withdrawal of ground water, which will definitely affect irrigation water resources in the near future (Ullah et al., 2008; Bozorov et al., 2016).

Cotton breeders have continued their efforts to create high-yielding cotton cultivars with improved fiber quality by usina existina cotton germplasm. Peohlman and Selper (1995) clarified that yield-contributing and fiber-quality traits are quantitatively heritable. Thus, seed cotton vield and its contributing components and fiber quality can be utilizing improved by new cross combinations developed through appropriate breeding programs. Breeders are mostly interested in possessing sufficient knowledge on genetic components, such as genetic variability, coefficient of variation, heritability, and genetic gain, to devise breeding plans that are appropriate for achieving specific breeding goals (Ali and Khan, 2007; Dhamayanathi et al., 2010). Cotton breeders have always encouraged genetic variability in breeding populations and suggested that screening breeding materials for tolerance to different biotic and abiotic stresses is an initial requirement. Great genetic variability, along with genotypic, phenotypic, and environmental coefficients of variation, has been reported among various upland cotton populations for quantitative and qualitative traits (Dhivya et al., 2014; Dahiphale et al., 2015; Shao et al., 2016; Adsare et al., 2017).

Heritability is an effective statistical tool that helps cotton breeders assess environmental impacts on various traits in a breeding nursery. It is an effective indicator for determining the level at which parental traits are passed down from generation to generation. Thus, heritability and GG can be a powerful tool for plant breeders to select appropriate breeding schemes (Chandio et al., 2003; Baloch et al., 2004). Abbas et al. (2013) provided information on high heritability coupled with GG for yield and yield component traits in upland cotton. Moderate to high heritability has been reported for bolls per plant, boll weight, seed cotton yield, lint percentage, fiber length, and strength in upland cotton genotypes (Aziz et al., 2014; Srinivas et al., 2014; Ahsan et al., 2015; Wagar et al., 2016; Nizamani et al., 2017).

Therefore, cotton improvement via selection largely depends on the discovery of genetic variability. Often, selection is based solely on phenotypic expression, which is often misleading because of environmental influence. Thus, data on genotypic, phenotypic, and environmental variances are of great importance for However, effective selection. the genotypic coefficient of variation does not provide an exact idea of the total heritable variation. Magadum et al. (2012) stated that the amount of genetic variation can be related by heredity. Heritability has to be estimated to estimate the proportion of phenotypic variance attributable to genetic variance. Such an estimation is vital because it also provides the basis for effective selection. Magadum et al. (2012) also pointed out that genetic variability,

along with the heritability of a character, will indicate the possibility and extent to which improvement is feasible through selection on a phenotypic basis. The heritability value alone may not provide clear predictability of the breeding value (Mishra et al., 2015). Hence, combination with GG over means is more effective and reliable in predicting the resultant effect of selection (Patil et al., 1996; Ramanjinappa et al., 2011). In presence of the highest magnitude of genetic variability, knowledge on heritability and GG helps the breeder select the desired characters achieve objectives. Therefore, to а thorough knowledge of genetic variability, heritability, and genetic development is necessary to improve the valuable economic traits of crop plants (Burton, 1952; Svarup and Chaugle, 1962).

The development of high-yielding cotton cultivars requires a comprehensive study of the relationship among growth resistance to adverse traits, environmental factors, and productivity in cotton. Heredity and physiological processes in crop plants depend on biological characteristics and environmental conditions. In other words, probability of inheritance the is determined by the degree of the main environmental factors (Bozorov et al., 2016, Nabiev et al., 2020). However, identifying the most precise indexes for evaluating plant drought tolerance remains а requirement. Therefore, identifying drought-tolerant traits and potential indexes and inducing them through different breeding schemes for crop improvement are necessary.

view the Keeping in above discussion, the importance of genetic studies, and the relationships among various quantitative and qualitative traits, the present study was performed to study the genetic variability, coefficient of variation, heritability, and GG in four parental genotypes (Ishonch, Navbakhor-2, C-6524, and Tashkent-6) and their F1 diallel hybrids for seed cotton yield and its component traits under optimal and deficit irrigation conditions.

MATERIALS AND METHODS

Genetic material, experimental site, and irrigation conditions

This study was conducted during 2018-2019 in the Tashkent region of Uzbekistan (41.389°N and 69.465°E). This region experiences cold winters and long hot and dry summers. The annual photoperiod (light/dark) is 16/8 h. This study involved the genetic potential and genetic aspects of four upland cultivars, i.e., Ishonch, Navbakhor-2, Tashkent-6, and S-6524, and their 12 F1 diallel hybrids under optimal and deficit irrigation conditions. These parental cultivars have an average fiber production (2.0-2.2 tons/hm²) but varied levels of drought tolerance. The parental cultivars and their F1 diallel hybrids were grown in a randomized complete block design with factorial arrangement and four replications under optimal and deficit irrigation conditions. The cotton genotypes were planted with plant and row spacings of 10 and 60 cm in 50 m long furrows. The soil moisture content was 70% under optimal irrigation conditions and 42% under deficit irrigation conditions as calculated with a moisture tester. Full (optimal) and deficit irrigation conditions were separated by a specified distance.

The temperature increases in April during the cotton sowing season and decreases in late September before the harvesting period. Information on

maximum and minimum temperatures, air humidity, and total rainfall during the study period is provided in Table 1. Sunny days were between 180-185 days. Rainfall varied from 0 mm to 45 mm during the dry season for a period of 5-6 months. The crop requires intensive irrigation throughout the vegetative period. Cotton is irrigated in accordance with a 1-2-1 (preflowering-flowering-boll opening) sequence with 900 m³/hm² of water applied before flowering, two applications of 1200 m^3/hm^2 each during flowering, and 900 m³/hm² prior to the boll-opening (Xamidovand phases Matyakubov, 2019). This sequence is an optimal irrigation protocol that is widely used in cotton production in Uzbekistan. Soil moisture also contributes water during seed germination. A modified irrigation protocol was also developed for irrigation conditions. This deficient protocol follows a 0-1-0 sequence that availability limits water during the flowering, preflowering, and boll development stages and reduces the total irrigation requirement to 1200 m³/hm² water (Xamidov and Matyakubov, 2019). For crop protection purposes, the insecticides Bi-58 (BASF, Germany) and Hexachloran were applied for the control sucking (aphids) and chewing of (bollworm) insects, respectively. The seasonal application of fertilizers was performed during tillage and before irrigation per annum with 250:180:115 NPK kg/hm².

Table 1.	Maximum	and	minimum	temperatures,	air	humidity,	and	total	rainfall	during	the
study per	iod.										

	Maximum		Minir	mum	Average	e relative	Total rainfall (mm)		
Months	temperature (°C)		tempera	temperature (°C)		ity (%)			
	2018	2019	2018	2019	2018	2019	2018	2019	
April	+31	+28	+3	+5	34%	+35	4.87	42.38	
May	+36	+36	+9	+10	26%	+38	1.79	11.25	
June	+37	+36	+14	+16	19%	+39	1.00	6.90	
July	+42	+42	+19	+20	15%	+39	0.00	2.43	
August	+39	+40	+15	+17	14%	+38	0.00	0.08	
September	+32	+36	+12	+10	15%	+33	0.16	1.05	
October	+31	+28	+1	+6	29%	+24	2.6	2.78	

Source of Variation	d.f.	Bolls plant ⁻¹		Boll weight (g)		100-seed weight (g)		Seed cotton yield plant ⁻¹ (g)		Lint (%)	
		Irrigated	Drought	Irrigated	Drought	Irrigated	Drought	Irrigated	Drought	Irrigated	Drought
Replications	3	5.36	12.19	5.49	0.002	0.05	0.19	0.93	21.19	7.77	0.80
Genotypes	15	6.97 ^{NS}	10.72*	0.51**	0.60**	1.98**	0.98**	121.52**	207.81**	11.35**	9.47**
Error	45	3.02	2.54	0.003	0.02	0.08	0.19	32.79	18.31	1.26	0.82

Table 2. Analysis of variance for various traits in upland cotton under optimal and deficit irrigation conditions.

**: Significant at $P \le 0.01$, *: Significant at $P \le 0.05$, NS: Nonsignificant

Data recorded and statistical analysis

The valuable economic traits of the parental genotypes and their F1 hybrids were studied under optimal and deficit irrigation conditions. The data on boll weight, bolls per plant, seed cotton yield per plant, and lint percentage were recorded. These parameters were used to monitor stress conditions for comparison with irrigated conditions. The data were subjected to analysis of variance (Steel et al., 1997). Genotypic, phenotypic, and environmental coefficients of variances were estimated by following Burton and Devane (1953). Broad-sense heritability was calculated in accordance with Honson et al. (1956), and GG was estimated as per Johnson et al. (1955). Correlation coefficients were calculated by using the formulae given by Kown and Torrie (1964).

 $V_g = [Genotypes mean squares - Error mean squares]/Number of replications$

 $V_e = Error mean squares$

 $V_p = V_g + V_e,$ Genotypic coefficient of variation (GCV) = $\sqrt{V_g}/GM \times 100$

Phenotypic coefficient of variation (PCV) = $\sqrt{V_p/GM \times 100}$

Environmental coefficient of variation (ECV) = $\sqrt{V_e/GM} \times 100$

where V_g = Genotypic variance V_p = Phenotypic variance V_e = Environmental variance GM = Grand mean of the trait. Broad-sense heritability (h^2) on an entry mean basis was calculated as

$$(h^2) = Vg/Vp.$$

The expected GG (GG) for each trait was calculated as follows:

$$GG = k. h^2 \sqrt{vp}.$$

The GG as a percentage of mean for each trait was calculated as

$$GG = GG/GM \times 100,$$

where k = 1.40 at 20% selection intensity for a trait

 V_P = Phenotypic variance for a trait

 h^2 = Broad sense heritability for a trait

GG = GG (expected response to selection)

RESULTS AND DISCUSSION

The analysis of variance showed that parental cultivars and F1 populations exhibited significant ($P \le 0.01$) differences for boll weight, 100-seed weight, seed cotton yield per plant, and lint% (Table 2). However, for bolls per plant, the genotypes exhibited significant ($P \le 0.05$) and nonsignificant differences under water deficit and optimum irrigation conditions, respectively. These observations indicated considerable genetic variability and scope among the genotypes for further improvement in yield-related traits. The mean performances of the parents and F_1 hybrids for various traits under study are presented in Tables 3 and 4. The data were further analyzed to estimate the heritability (broad sense), GG, GCV%, PCV%, and ECV% (Table 5). The

Parental genotypes	Irrigated/	Bolls plant ^{-1}	Boll weight (g)	100-seed	Seed cotton vield plant ⁻¹	Lint (%)
and F1	Drought	·	5 (5)	weight (g)	, (g)	
hybrids	-	M ± SE	M ± SE	$M \pm SE$	M ± SE	M ± SE
Ishonch	Irrigated	16.45 ± 0.63	5.62 ± 0.06	12.20 ± 0.10	60.18 ± 3.00	36.31 ± 0.40
	Drought	11.50 ± 0.89	5.00 ± 0.06	12.09 ± 0.11	42.09 ± 1.69	37.54 ± 0.45
Navbakhor-2	Irrigated	14.80 ± 0.53	5.80 ± 0.06	12.93 ± 0.10	57.18 ± 3.38	36.47 ± 0.56
	Drought	10.30 ± 0.85	5.13 ± 0.06	12.73 ± .013	42.14 ± 1.42	38.80 ± 0.38
Tashkent-6	Irrigated	16.95 ± 0.74	5.75 ± 0.09	12.60 ± 0.16	62.40 ± 2.32	35.77 ± 0.47
	Drought	7.90 ± 057	4.34 ± 0.04	11.44 ± 0.15	27.17 ± 0.80	37.62 ± 0.08
C-6524	Irrigated	17.35 ± 0.85	5.08 ± 0.07	12.20 ± 0.09	63.97 ± 1.73	36.90 ± 0.78
	Drought	8.30 ± 0.65	4.29 ± 0.09	10.99 ± 0.13	27.10 ± 0.93	37.20 ± 0.11
Ishonch ×	Irrigated	15.20 ± 0.69	6.24 ± 0.02	14.23 ± 0.10	69.07 ± 2.28	41.13 ± 0.55
Navbakhor-2	Drought	12.80 ± 0.65	5.34 ± 0.08	12.58 ± 0.13	49.86 ± 3.05	41.17 ± 0.39
Ishonch ×	Irrigated	17.00 ± 1.06	5.73 ± 0.04	14.90 ± 0.08	74.15 ± 2.86	39.68 ± 0.70
Tashkent-6	Drought	8.95 ± 1.05	4.82 ± 0.08	12.18 ± 0.27	46.12 ± 2.11	39.84 ± 0.51
Ishonch \times C-	Irrigated	15.50 ± 0.54	5.51 ± 0.08	13.52 ± 0.08	70.74 ± 3.20	36.87 ± 0.49
6524	Drought	11.25 ± 0.45	4.84 ± 0.04	12.02 ± 0.11	43.52 ± 2.00	40.26 ± 0.048
Navbakhor-2	Irrigated	14.50 ± 1.02	5.70 ± 0.07	13.33 ± 0.15	75.65 ± 3.63	39.27 ± 0.44
× Ishonch	Drought	10.30 ± 0.85	5.34 ± 0.05	$12.05 \pm 0,19$	45.37 ± 3.05	42.72 ± 0.48
Navbakhor-2	Irrigated	15.40 ± 0.51	6.22 ± 0.07	13.20 ± 0.07	65.64 ± 2.39	37.08 ± 0.31
× Tashkent-6	Drought	8.00 ± 0.52	5.43 ± 0.05	11.14 ± 0.12	38.49 ± 1.28	41.53 ± 0.65
Navbakhor-2	Irrigated	16.901.29	6.31 ± 0.07	13.23 ± 0.21	66.71 ± 4.75	39.08 ± 0.14
× C-6524	Drought	8.60 ± 0.67	5.56 ± 0.07	11.68 ± 0.95	41.81 ± 2.58	40.59 ± 0.30
Tashkent-6 ×	Irrigated	19.20 ± 0.73	5.45 ± 0.05	13.04 ± 0.17	77.48 ± 2.43	36.24 ± 0.75
Ishonch	Drought	9.10 ± 0.71	4.54 ± 0.07	11.12 ± 0.21	35.04 ± 1.08	40.13 ± 0.51
Tashkent-6 ×	Irrigated	17.30 ± 0.92	5.38 ± 0.06	13.19 ± 0.10	68.62 ± 2.37	36.05 ± 0.93
Navbakhor-2	Drought	7.90 ± 0.63	4.46 ± 0.08	11.06 ± 0.22	33.26 ± 1.65	38.33 ± 0.51
Tashkent-6 ×	Irrigated	14.65 ± 0.65	5.52 ± 0.05	12.67 ± 0.09	64.75 ± 1.94	39.06 ± 0.68
C-6524	Drought	7.05 ± 0.49	4.61 ± 0.08	11.74 ± 0.22	28.44 ± 1.89	39.33 ± 0.45
C-6524 ×	Irrigated	15.20 ± 0.90	5.42 ± 0.06	13.92 ± 0.22	70.24 ± 2.53	37.88 ± 0.51
Ishonch	Drought	7.60 ± 0.34	4.73 ± 0.06	11.81 ± 0.18	32.61 ± 1.45	38.97 ± 0.70
C-6524 ×	Irrigated	16.20 ± 1.00	5.40 ± 0.07	13.07 ± 0.13	67.17 ± 2.45	37.34 ± 0.40
Navbakhor-2	Drought	8.20 ± 0.44	4.64 ± 0.04	11.63 ± 0.21	32.86 ± 0.93	39.27 ± 0.56
C-6524 ×	Irrigated	14.60 ± 0.69	5.23 ± 0.07	12.91 ± 0.09	63.79 ± 1.82	40.45 ± 0.57
Tashkent-6	Drought	8.10 ± 0.35	4.57 ± 0.10	11.49 ± 0.22	32.42 ± 1.22	40.64 ± 0.75

Table 3. Mean performance of parental genotypes and their F1 hybrids for various traits in upland cotton under optimal and deficit irrigation conditions.

Table 4.	Basic statistics	in parental	cultivars	and their I	F1 hybrids	for various	traits in uplar	۱d
cotton.								

Traits	Irrigated/ Drought	Grand mean	Minimum	Maximum	Variance	CV (%)
Bolls plant ⁻¹	Irrigated	16.09	11.60	20.60	4.07	0.12
	Drought	9.12	6.60	14.20	4.95	0.24
Boll weight (g)	Irrigated	5.65	5.03	6.32	0.12	0.06
	Drought	4.88	4.23	5.69	0.16	0.08
100-seed weight (g)	Irrigated	13.20	11.94	15.12	0.53	0.05
	Drought	11.75	10.58	13.27	0.38	0.05
Seed cotton yield plant ^{-1} (g)	Irrigated	67.36	50.61	85.41	52.42	0.11
	Drought	37.39	25.66	55.46	63.57	0.21
Lint (%)	Irrigated	37.89	34.25	41.98	3.98	0.05
	Drought	39.62	37.03	44.45	2.88	0.04

Traits	Irrigated /Drought	GV	GCV (%)	PV	PCV (%)	EV	ECV (%)	h²	GG	GG (%)
Bolls plant ⁻¹	Irrigated	0.98	6.17	4.01	12.45	3.02	10.81	0.24	1.01	6.31
	Drought	2.05	15.68	4.58	23.50	2.54	17.50	0.44	1.97	21.60
Boll weight	Irrigated	0.12	6.30	0.13	6.39	0.004	1.06	0.97	0.72	12.82
	Drought	0.14	7.82	0.16	8.34	0.02	2.90	0.88	0.52	10.74
100-seed	Irrigated	0.47	5.23	0.55	5.63	0.08	2.08	0.86	1.32	10.04
weight	Drought	0.19	3.78	0.39	5.34	0.19	3.77	0.50	0.64	5.52
Seed cotton	Irrigated	22.18	6.99	54.97	11.00	32.79	8.50	0.40	6.17	9.16
yield plant ⁻¹	Drought	47.37	18.41	65.69	21.67	18.32	11.44	0.72	12.06	32.25
Lint (%)	Irrigated	2.52	4.19	3.79	5.13	1.26	2.97	0.66	2.67	7.05
	Drought	2.16	3.71	2.98	4.36	0.82	2.29	0.72	2.51	6.51

Table 5. Genotypic, phenotypic, and environmental variances and coefficient of variation, broad sense heritability, and GG under optimal and deficit irrigation conditions.

GV: Genotypic variance; GCV%: Genotypic coefficient of variance; PV: Phenotypic variance; PCV%: Phenotypic coefficient of variance; EV: Environmental variance; ECV%: Environmental coefficient of variance; h²: Heritability (broad sense); Genetic gain: GG

Table 6. Correlation coefficients among yield-related traits under optimal irrigation (below diagonal) and deficit irrigation (above diagonal) conditions.

Traits	Bolls plant ⁻¹	Boll weight	100-seed weight	Seed cotton yield plant ⁻¹	Lint%
Bolls plant ⁻¹	_	0.467***	0.395**	0.421***	0.033 ^{NS}
Boll weight	-0.045 ^{NS}	-	0.327**	0.396**	0.033 ^{NS}
100-seed weight	-0.099 ^{NS}	0.372**	-	0.001 ^{NS}	0.546***
Seed cotton yield plant ⁻¹	0.763***	0.554***	0.718***	_	0.350**
Lint (%)	-0.377**	0.425***	0242 ^{NS}	0.15 ^{NS}	

**: Significant at $P \le 0.01$, *: Significant at $P \le 0.05$, NS: Nonsignificant

correlation coefficients among various yield related traits are presented in Table 6.

Mean performance

Under optimal irrigation, bolls per plant fell in the ranges of 14.80-17.35 and 14.50–19.20 in parental cultivars and F₁ hybrids, respectively (Tables 3 and 4). Among parental genotypes, the cultivars C-6524 (17.35), Tashkent-6 (16.95), and Ishonch (16.45) produced the maximum bolls per plant, whereas Navbakhor-2 (14.80) presented the lowest bolls per plant. Among the F_1 hybrids, Tashkent-6 \times Ishonch (19.20) displayed the maximum bolls per plant, followed by Tashkent-6 \times Navbakhor-2 (17.30), whereas the lowest bolls per plant were recorded for the F1 hybrid Navbakhor-2 \times Ishonch (14.50) under optimal irrigation conditions.

Under deficit irrigation conditions, the bolls per plant fell in the ranges of

7.90-11.50 and 7.05-12.80 in parental cultivars and F₁ hybrids, respectively (Tables 3 and 4). Among the parental genotypes, Ishonch (11.50)and Navbakhor-2 (10.30)provided the maximum number of bolls per plant, followed by cultivar C-6524 (8.30). However, the lowest number of bolls per plant was recorded for the cultivar Tashkent-6 (7.90). Among F₁ hybrids, Ishonch × Navbakhor-2 (12.80) displayed the maximum bolls per plant, followed by two other F1 hybrids, i.e., Ishonch \times C-6524 (11.25) and Navbakhor-2 × Ishonch (10.30). However, the lowest bolls per plant were recorded for the F1 hybrid Tashkent-6 \times C-6524 (7.05) under water deficit conditions.

Under optimal irrigation, boll weight ranged from 5.08 g to 5.80 g and from 5.23 g to 6.31 g in parental cultivars and F_1 hybrids, respectively (Tables 3 and 4). The heaviest boll weight in parents was recorded for cultivar Navbakhor-2

(5.80 g), followed by Tashkent-6 (5.75 g)then by Ishonch (5.62 g), whereas the lowest boll weight was observed for cultivar C-6524 (5.08 g). Among F1 cross combinations, Navbakhor-2 × C-6524 (6.31 g) and Ishonch × Navbakhor-2 (6.24 g), followed by Navbakhor-2 \times Tashkent-6 (6.22 g), showed the maximum boll weight. The lowest boll weight was recorded for F1 hybrids, i.e., C-6524 \times Tashkent-6 (5.23 g), Tashkent- $6 \times \text{Navbakhor-2}$ (5.38 g), and C-6524 \times Navbakhor-2 (5.40 g), under optimal irrigation conditions.

Under deficit irrigation, in parental cultivars and F₁ hybrids, boll weight varied from 4.29 g to 5.13 g and to 4.46 g to 5.56 g, respectively (Tables 3 and 4). Among parental genotypes, the bigger bolls were recorded for Navbakhor-2 (5.13 q), followed by Ishonch (5.00 g). The lowest boll weight was observed for Tashkent-6 (4.34 g) and C-6524 (4.29 g). Among F_1 hybrids, the highest boll weight was recorded in hybrid Navbakhor-2 × C-6524 (5.56 g), followed by Navbakhor-2 \times Tashkent-6 (5.43) and Ishonch X Navbakhor-2 and its reciprocal (5.34 g). F_1 hybrids i.e., Tashkent-6 × C-6524 (4.61 g) and Tashkent-6 × Ishonch (4.54 g)g), followed by Tashkent-6 \times Navbakhor-2 (4.46 g) showed the minimum boll weight under water deficit conditions.

Under optimal irrigation, 100-seed weight ranged from 12.20 g to 12.93 g and 12.67 g to 14.90 g in parental cultivars and F₁ hybrids, respectively (Tables 3 and 4). The maximum 100-seed displayed weight was by cultivar Navbakhor-2 (12.93 g), followed by Tashkent-6 (12.60 g). The parental genotypes C-6524 and Ishonch showed the minimum and same 100-seed weight. The F1 hybrids i.e., Ishonch × Tashkent-6 (14.90 g) and Ishonch × Navbakhor-2 (14.23 g), followed by C-6524 \times Ishonch (13.92 g), exhibited the highest 100-seed weight. However, F1 hybrid Tashkent-6 \times C-6524 revealed the lowest 100-weight (12.67 g) with optimum irrigation condition.

Under deficit irrigation condition, the 100-seed weight of the parental

cultivars and F₁ hybrids ranged from 10.99 g to 12.73 g and 11.06 g to 12.58 g, respectively (Tables 3 and 4). Among parental cultivars, the highest 100-seed weight was displayed by Navbakhor-2 (12.73 g), followed by Ishonch (12.09 g). The cultivars Tashkent-6 (11.44 g) and C-6524 (10.99 g) showed the least 100-seed weight. Among F1 hybrids, the highest 100-seed weight was revealed by two hybrids i.e., Ishonch × Navbakhor-2 (12.58 g) and Ishonch × Tashkent-6 (12.18 g). The minimum 100-seed weight was recorded in F1 hybrids i.e., Tashkent- $6 \times \text{Navbakhor-2}$ (11.06 g) and its reciprocal (11.14 g), and Tashkent-6 \times Ishonch (11.12 g) with deficit irrigation condition.

Under optimal irrigation, the seed cotton yield per plant ranged from 57.18 g to 63.97 g and 63.79 g to 77.48 g in cultivars F_1 hybrids, parental and respectively (Tables 3 and 4). The parental cultivar C-6524 (63.97 q) displayed maximum seed cotton yield per plant, followed by two other cultivars, i.e., Tashkent-6 (62.40 g) and Ishonch (60.18 g). The parental genotype Navbakhor-2 (57.18 g) was recorded with the lowest seed cotton yield per plant. Among F_1 hybrids, Tashkent-6 \times Ishonch (77.48 g) and its reciprocal (74.15 and q) Navbakhor-2 \times Ishonch (75.65 g) showed the maximum seed cotton yield per plant. The lowest seed cotton yield per plant was obtained in the F1 hybrid C-6524 \times Tashkent-6 (63.79 g) and its reciprocal Tashkent-6 \times C-6524 (64.75 g).

Under deficit irrigation conditions, the seed cotton yield per plant of the parental genotypes and F1 hybrids ranged from 27.10 g to 42.14 g and 28.44 g to 77.48 g, respectively (Tables 3 and 4). Under water deficit conditions, the parental genotype Navbakhor-2 (42.14 g), followed by the cultivar Ishonch (42.09 g), displayed the maximum yield per plant. The cultivar Tashkent-6 (27.17 q), followed by C-6524 (27.10 g), exhibited the lowest seed cotton yield per plant. The F1 hybrids Ishonch × Navbakhor-2 (49.86 g), Ishonch \times Tashkent-6 (46.12), and Navbakhor-2 \times Ishonch (45.37 g) showed the maximum seed cotton yield per plant. The lowest seed cotton yield per plant was recorded in the F1 populations Tashkent-6 \times C-6524 (28.44 g), C-6524 \times C-Tashkent-6 (32.42 g), and Tashkent-6 \times Navbakhor-2 (33.26 g) under deficit irrigation.

Under optimal irrigation, the lint% of the parental genotypes and F1 hybrids ranged from 35.77% to 36.99% and 36.05% to 41.13%, respectively, (Tables 3 and 4). In parental cultivars, the highest lint% was exhibited by the parental genotype C-6524 (36.90%), followed by Navbakhor-2 (36.47%). The lowest lint% was recorded for the parental cultivar Tashkent-6 (35.77%). Similarly, among F1 hybrids, the cross combinations Ishonch × Navbakhor-2 (41.13%) and C- $6524 \times C$ -Tashkent-6 (40.45%) displayed the maximum lint%. The lowest lint% was recorded for the F1 hybrids Tashkent-6 \times Navbakhor-2 (36.05%) and Tashkent-6 \times Ishonch (36.24%) under optimal irrigation.

Under water deficit conditions, the lint% of the parental genotypes and F1 hybrids varied from 37.20% to 38.80% and from 38.33% to 42.72%, respectively (Tables 3 and 4). The mean performance of the parents for lint% revealed that the parental genotype Navbakhor-2 (38.80%), followed by Tashkent-6 (37.62%), displayed the maximum lint%. The lowest lint% was recorded for the parental cultivar C-6524 (37.20%).Similarly, among F1 hybrids, the cross combinations Navbakhor-2 × Ishonch (42.72%) and Navbakhor-2 × Tashkent-6 (41.53%) displayed the maximum lint%. The lowest lint% was recorded for the F1 hybrids Tashkent-6 Navbakhor-2 × (38.33%) and C-6524 Ishonch х (38.97%).

Overall, the mean performance of the genetic material for various yieldrelated traits revealed a considerable amount of genetic variability. Under the optimal irrigation condition, no one genotype was superior for all traits. However, under deficit irrigation condition, the parental cultivars Ishonch and Navbakhor-2 and their hybrids were found to be superior for all traits. Previous studies on the assessment of cotton germplasm under various environmental conditions also reported a significant magnitude of genetic variability for seed cotton yield and its components (Islam *et al.*, 2013; Srinivas *et al.*, 2014; Dahiphale *et al.*, 2015; Raza *et al.*, 2016; Nizamani *et al.*, 2017; Marguba *et al.*, 2020).

GCV, PCV, and ECV

The ranges of genotypic, phenotypic and environmental variances for various traits in parental genotypes and their F1 hybrids are presented in Table 5. Under optimal irrigation condition, the GCV and PCV values varied from 4.19% to 6.99% and from 5.13% to 12.45%, respectively. Under deficit irrigation, the estimates of GCV and PCV ranged from 3.71% to 18.41% and 4.36% to 23.50% respectively. Under the optimal irrigation regime, the highest GCV and PCV effects were found for seed cotton yield per plant (6.99%) and 11.00%, respectively), followed by bolls per plant (6.17% and 12.45%, respectively) and boll weight (6.30% and 6.39%, respectively). The lowest values of GCV and PCV were observed for 100-seed weight (5.23% and 5.63%, respectively) and lint% (4.19%) and 5.13%, respectively).

Under deficit irrigation conditions, the highest GCV and PCV effects were found for seed cotton yield per plant (18.41% and 21.67%, respectively), followed by bolls per plant (15.68% and 23.50%, respectively) and boll weight (7.82% and 8.34%, respectively) (Table 5). The lowest GCV and PCV effects were noted for lint% (3.71% and 4.36%, respectively) and 100-seed weight (3.78% and 5.34%, respectively). Saleem et al. (2016) reported the highest genetic variability in segregating populations for economically important traits and declared it as a prerequisite for the success in breeding programs. Adequate information regarding genotypic variances make selection in breeding populations effective

given that plant traits are mostly influenced by environmental effects (Mahaingam *et al.*, 2013).

The 100-seed weight and lint% showed low GCV and PCV values, suggesting limited room for the further improvement in these traits. For traits with low estimates of GCV and PCV, the breeders should search for the source of highest genetic variability for further improvement. Cotton breeders must exploit Uzbek germplasm from diverse sources to identify the genetic variability in breeding populations. Moderate PCV and GVC for yield traits suggest that these characters can be improved by using vigorous selection methods (Girase and Mehatne, 2002; Harshal, 2010). Kowsalya and Raveendran (1996) provided similar suggestions in their studies on upland cotton. However, some of the past studies showed higher values of GCV and PCV for seed cotton yield and yield related traits (Ahuja and Tuteja, 2000; Girase and Mehetre, 2002; Ganesan and Raveendran, 2007). In upland cotton genotypes, boll weight showed moderate estimates of PCV and GCV and indicated the possibility of improvement in yield and yield component traits by further selection in segregating populations (Rao and Reddy, 2001; Prasad et al., 2005).

Under optimal and deficit irrigation conditions, the ECV ranged from 2.97% to 10.81% and from 2.29% to 17.50%, respectively (Table 5). The highest ECV values were recorded for bolls per plant (10.81% and 17.50%), followed by seed per plant (8.50% and cotton yield 11.44%) under deficit and optimal conditions, irrigation respectively, suggesting that these traits were considerably influenced by environmental factors. The selection of genotypes in early generations with moderate to high GCV and PCV is also recommended for improvement in yield and yield component traits (Ahsan et al., 2015; Shao et al., 2016; Adsare et al., 2017). Thus, genetic variability with high heritability estimates play an important role in the inheritance and improvement of yield-related traits in upland cotton (Ahmed et al., 2011).

coefficient of The phenotypic variation has the same contribution as the coefficient of genotypic variation. In this study, the GCV and PCV for all the traits showed close resemblance, indicating that these traits were less influenced by the environment. The highest GCV and PCV were observed for boll weight, 100-seed weight, and seed cotton yield per plant under water deficit conditions, and effective selection is recommended to isolate highly promising cotton lines. Past studies also reported similar type of observations for yield-related traits in upland cotton populations (Dheva and Potdukhe, 2002; Preetha and Raveendran, 2007; Amir et al., 2012; Abbas et al., 2013).

Heritability (broad sense)

The present investigation revealed the highest broad sense heritability estimates for boll weight, 100-seed weight, seed cotton yield, and lint% (Table 5). low heritability (bs) However, was reported for bolls per plant. The high heritability estimates highlighted the genetic variance importance of and depicted that variation among the populations for majority of the traits, except bolls per plant, is under the influence of genetic factors. Past studies reported high heritability and greeter genetic variability estimates for seed cotton yield and yield component traits in upland cotton genotypes (Khan et al., 2010; Batool et al., 2010; Ahmed et al., 2011; Aziz et al., 2014; Raza at al., 2016; Nizamani et al., 2017). Additional genetic influence with lower environmental factors predicted that most of the traits are controlled by additive gene action. The present results were also in accordance with the results of Abbas et al. (2013) and Dhivya et al. (2014) for upland cotton genotypes and were also in agreement with earlier findings of Ahuja and Tuteja (2000), Girase and Mehetre (2002), Prasad et al. (2005), and Pujer et al. (2014), who reported similar values of broad-sense heritability for seed cotton yield and yield-contributing traits in in upland cotton. For yield-related traits, the greater genetic variance might be effective in selection and improvement because of additive gene action (Rokadia and Vaid, 2003; Eswari *et al.*, 2017).

High heritability is a determinant of genotype flexibility in the selection process. In this investigation, the high heritability coupled with high GG observed for boll weight and seed cotton yield per plant reflected the preponderance of additive gene action in the inheritance of these traits. Johnson et al. (1955) and Swarup and Chaugale (1962) indicated that high heritability is not always an indication of high GG. If the transmission of heredity from generation to generation is mainly due to nonadditive gene effects. then the expected GG will be low; however, if some additive gene effects exist, then the expected GG will be high (Panse, 1957). Ahuja and Tuteja (2000), Girase and Mehetre (2002), Kumari and Chamundeshwari (2005), Kale et. al. (2006), and Preetha and Raveendran (2007)also reported the highest heritability and GG values for seed cotton yield and yield-related traits in upland genotypes.

In this study, under water deficit conditions and considering genetic variability, heritability and GG, selection would be effective for bolls per plant and boll weight besides yield per plant for developing high-yielding cotton cultivars. Hence, the pediaree breeding method will be the rewarding method for improving the traits under investigation. Characters with high heritability and GG can be used as tools in the selection process, and such traits are controlled by additive gene effects that are less affected by the environment (Panes and Suxhatme, 1995). Under optimal and deficit irrigation conditions, the GG, as the percentage of means for various traits, ranged from 6.31% to 12.82% and 5.52% to 32.25%. A higher GG as the percentage of mean was recorded for seed cotton yield per plant (9.16% and 32.25%), followed by boll weight (12.82% and 10.74%), bolls per plant (6.31% and 21.6%), 100-seed weight (10.04% and 5.52%) and lint%

(7.05% and 6.51%) under optimum and deficit irrigation conditions, respectively.

High heritability and GG were observed for seed cotton yield per plant and boll weight and were considered as highly reliable during selection. High heritability and moderate GG were reported for lint% in upland cotton genotypes (Muhammad et al., 2004). In present studies, some of the genotypes were identified as potential donors for improvement in different traits. Therefore, direct selection can be more effective for improvement in majority of the traits. Johnson et al. (1955) suggested that heritability estimates in conjunction with high GG were usually helpful in predicting its resultant effects for selecting promising genotypes.

Correlation coefficient

Under optimal irrigation condition, bolls per plant revealed a significant ($P \le 0.01$) positive association with seed cotton yield but negative associations with all other traits (Table 6). Boll weight exhibited significant ($P \le 0.01$) positive correlations with 100-seed weight, seed cotton yield, and lint%. Hundred-seed weight exhibited significant (P ≤ 0.01) а positive relationship with seed cotton yield but a nonsignificant positive relationship with lint%. Seed cotton yield showed a nonsignificant positive association with lint%.

Under water deficit conditions, bolls per plant revealed a significant ($P \leq$ 0.01) positive association with all the traits except lint% (Table 6). Boll weight exhibited a significant ($P \le 0.01$) positive correlation with 100-seed weight and seed cotton yield but a nonsignificant positive relationship with lint%. Hundred-seed weight exhibited a significant ($P \leq 0.01$) positive association with lint% but a nonsignificant positive relationship with seed cotton yield. Seed cotton yield showed a significant ($P \leq 0.01$) positive association with lint%. Overall, seed cotton yield revealed highly significant positive associations with all the traits, except for 100-seed weight under water

deficit condition, under optimal and deficit irrigation conditions.

Past studies revealed the positive correlation of seed cotton yield with bolls per plant and boll weight (Alkuddsi *et al.*, 2013; Jaloliddin *et al.*, 2020; Zeeshan *et al.*, 2020). Previous studies also reported the positive correlation of seed cotton yield with lint% and boll weight (Amanov *et al.*, 2020). Tohir *et al.* (2018) mentioned the positive correlation of seed cotton yield with bolls per plant, boll weight, and 100-seed weight under water deficit conditions.

CONCLUSIONS

The cotton cultivars Ishonch and Navbakhor-2 and their F1 hybrids were more stable and performed better than genotypes under both other water regimes. These parental genotypes and their hybrids can be used as source materials for developing genotypes with tolerance to drought conditions. Broadsense heritability was the highest for boll weight, 100-seed weight, seed cotton yield, and lint% but was low for bolls per plant. Genotypic and phenotypic variances for various traits were greater under water deficit conditions than under the optimal irrigation regime. Overall, under optimal and deficit irrigation conditions, seed cotton yield revealed a highly significant positive association with the majority of the traits.

ACKNOWLEDGEMENTS

The authors are grateful to the Zangi-Ata Experimental Field Station for providing space and resources to carry out this work.

REFERENCES

Abbas HG, Mahmood A, Ali Q, Khan, MA, Nazeer W, Aslam T, Zahid W (2013). Genetic variability heritability genetic advance and correlation studies in cotton (*G. hirsutum* L.). *Int. Res. J. Microbiol.* 4(6): 156-161.

- Adsare AN, Abhay NS (2017). Study on genetic variability for the quantitative traits in some genotypes of upland cotton (*G. hirsutum* L.). *Biosci. Discov.* 8(3): 365-368.
- Ahmad M, Khan NU, Mohammad F, Khan SA, Munir I, Bibi Z, Shaheen S (2011). Genetic potential and heritability studies for some polygenic traits in cotton (*G. hirsutum* L.). *Pak. J. Bot.* 43(3): 1713-1718.
- Ahsan MZ, Majidano MS, Bhutto H, Soomro AW, Panhwar FH, Channa AR, Sial KB (2015). Genetic Variability, coefficient of variance, heritability and genetic advance of some *G. hirsutum* L. accessions. *J. Agric. Sci.* 7(2): 147-151.
- Ahuja SL, Tuteja OP (2000). Variability and association analysis for chemical components imparting resistance to *G. hirsutum* cotton. *J. Cotton Res. Dev.* 14(1): 19-22.
- Ali MA, Khan IA (2007). Assessment of genetic variation and inheritance mode of some metric traits in cotton (*G. hirsutum* L.). *J. Agric. Soc. Sci.* 3: 112-116.
- Alkuddsi Y, Rao MR, Patil SS, Joshi M, Gowda TH (2013). Correlation and path coefficient analysis between seed cotton yield and its attributing characters in intra *G. hirsutum* cotton hybrids. *Mol. Plant Breed*. 4: 214-219.
- Amanov B, Abdiev F, Shavkiev J, Mamedova F, Muminov K (2020). Valuable economic indicators among hybrids of peruvian cotton genotypes. *Plant Cell Biotechnol. Mol. Biol.* 21(67-68): 35-46.
- Amir S, Farooq J, Bibi A, Khan SH, Saleem MF (2012). Genetic studies of earliness in *G. hirsutum* L. Int. J. Agro. Vet. Med. Sci. 6(3): 189-207.
- Aziz U, Afzal J, Iqbal M, Naeem M (2014). Selection response, heritability and genetic variability studies in upland cotton. *J. Appl. Environ. Biol. Sci.* 4(8S): 400-412.
- Babar M, Saranga Y, Iqbal Z, Arif M, Zafar Y, Ρ Lubbers Ε, Chee (2009). Identification of QTLs and impact of selection from various environments (dry vs. irrigated) on the genetic relationships among the selected cotton lines from F6 population using a phylogenetic approach. Afr. 1 Biotechnol. 8: 4802-4810.
- Baloch MJ (2004). Genetic variability and heritability estimates of some polygenic

traits in upland cotton. *Pak. J. Sci. Ind. Res.* 42(6): 451-454.

- Batool S, Khan NU, Makhdoom K, Bibi Z, Hassan G, Marwat KB, Farhatullah F, Raziuddin M, Khan IA (2010). Heritability and genetic potential of upland cotton genotypes for morphoyield traits. *Pak. J. Bot.* 42(6): 1057-1064.
- Bozorov T, Shavkiev J, Usmanov R, Nabiev S, Khamdullaev SH (2016). Metabolome profiling during water deficiency in medium-fiber cotton varieties of *G. hirsutum* L. species. *Problems of Modern Sci. Edu.* 33(75): 10-12.
- Burton GW (1952). Quantitative inheritance in grasses. *Proceed. sixth Int. Cong.* pp. 277-283.
- Burton GW, Devane EM (1953). Estimating heritability in fall fescue (Festecdcirunclindcede) from replicated clonal material. *Agron. J.* 45: 478-481.
- Chandio MA, Kalwar MS, Baloch GM (2003). Gene action for some quantitative characters in upland cotton (*G. hirsutum* L.). *Pak. J. Sci. Indus. Res.* 46(13): 295-299.
- Dahiphale KD, Deshmukh JD, Jadhav AB, Bagade AB (2015). Genetic variability and correlation for yield and its attributing traits in cotton (*G. hirsutum* L). *Int. J. Trop. Agric.* 33(1): 15-22.
- Dhamayanathi KPM, Manickam S, Rathinaval K (2010). Genetic variability studies in *G. barbadense* L. genotypes for seed cotton yield and its components. *Electr. J. Plant. Breed*. 1(4): 961-965.
- Dheva NG, Potdukhe NR (2002). Studies on variability and correlations in upland cotton for yield and its components. *J. Indian Soc. Cotton Imp.* 148-152.
- Dhivya R, Amalabalu P, Pushpa R, Kavithamani D (2014). Variability, heritability and genetic advance in upland cotton (*G. hirsutum* L.). *Afr. J. Plant Sci.* 8(1): 1-5.
- Eswari KB, Sudheer Kumar S, Gopinath, Rao MVB (2017). Genetic variability heritability and genetic advance studies in cotton. *Int. J. Dev. Res.* 7(1): 10902-10904.
- Fayetteville AR. Ahmad RT, Malik TA, Khan IA, Jaskani MJ (2009). Genetic analysis of some morpho-physiological traits related to drought strss in cotton (*G. hirsutum* L.). *Int. J. Agric. Biol.* 1: 235-240.
- Ganesan KN, Raveendranm TS (2007). Enhancing the breeding value of

genotype through genetic selection in cotton (*G. hirsutum* L.). *Bull. Inst. Trop. Agr., Kyushu Uni*. 30:1-10.

- Girase VS, Mehetre S S (2002). Variability, heritability and genetic advance studies in cotton (*G. hirsutum* L.). *J. Cotton Res, Dev.* 16(1): 81-82.
- Hanson CH, Robinson HF, Comstock RE (1965). Biometrical studies of yield in segregating populations of Korean Lespedeze. *Agron. J.* 48:314-318.
- Harshal EP (2010). Variability and correlation analysis by using various quantitative traits in released Bt cotton hybrids. J. Cotton Res. Dev. 24(2):141-144.
- Hinze LL, Dever JK, Percy RG (2012). Molecular variation among and within improved cultivars in the U.S. cotton germplasm collection. *Crop Sci*. 52: 222-230.
- Islam MK, Akhteruzzaman M, Sharmin D (2013). Multivariate and genetic component analysis of new cotton (*G. hirsutum* L.) genotypes. *Bangladesh J. Prog. Sci. Technol.* 11(2): 185-190.
- Johnson HW, Robinson HE, Comstock RE (1955). Estimate of genetic and environmental variability in soybean. *Agron. J.* 47: 314-318.
- Kale UV, Kalpande HV, Gunjkar AS, Gite VK (2006). Variability, heritability and genetic advance studies in cotton. *J. Maharashtra Agric. Uni.* 31(3): 389-390.
- Khan NU, Marwat KB, Hassan G, Farhatullah, Batool S, Makhdoom K, Ahmad W, Khan HU (2010). Genetic variation and heritability for cottonseed, fiber and oil traits in *G. hirsutum* L. *Pak. J. Bot.* 42(1): 615-625.
- Khokhar ES, Shakeel A, Maqbool MA, Anwar MW,Tanveer Z, Irfan MF (2017). Genetic study of cotton (*G. hirsutum* L.) genotypes for different agronomic, yield and quality traits. *Pak. J. Agric. Res.* 30(4): 363-372.
- Kowsalya R, Raveendran TS (1996). Genetic variability and D2 analysis in upland cotton. *Crop Res.* 12(1): 36-42.
- Krieg DR (2000): Cotton Water Relations.
 Special Report In: Oosterhuis, D. M. (ed.) Proc. Cotton Research Meeting and Summaries of Cotton Research.
 Arkansas Agric. Exp. Station, USA.
- Kumari SR, Chamundeshwari N (2005). Studies on genetic variability, heritability and genetic advance in cotton (*G. hirsutum* L.). *Res. on Crops*. 6(1): 98-99.

- Kwon SH, Torrie JH (1964). Heritability and interrelationship among traits of two soybean population. *Crop Sci.* 4: 194–202.
- Magadum S, Banerjee U, Ravikesavan R, Gangapur D, Boopathi NM (2012). Variability and heritability analysis of yield and quality traits in interspecific population of cotton (*Gossypium Spp.*). *Bioinfolet* 9(4A): 484-485.
- Mahalingam A, Saraswathi R, Ramalingam J, Jayaraj T (2013). Genetics of floral traits in cytoplasmic male sterile (CMS) and restorer lines of hybrid rice (*Oryza sativa* L.). *Pak. J. Bot.* 45(6): 1897-1904.
- Marguba MR, Abdulaxat AA, Mansur MK, Ilkham DK, Hilola XM (2020). Indicators of abiotic and biotic stresses of local and foreign cotton (*G. hirsutum* L.). *Plant Cell Biotechnol. Mol. Biol.* 21(43&44): 8-15.
- Mishra PK, Ram RB, Kumar N (2015). Genetic variability, heritability, and genetic advance in strawberry (*Fragaria* × *ananassa Duch.*). *Turk J Agric. For.* 39: 451-458
- NabievCM, Usmanov RM, Khamdullaev SA, Shavkiev JS (2020). Study of physiological indicators of the water balance of plants and morphological signs of leaf of fine-fiber varieties in different irrigation regimes. *J. Biol. Uzbekistan* 1: 51-58.
- Nizamani F, Baloch MJ, Baloch AW, Buriro M, Nizamani G.H, Nizamani MR, Baloch IA (2017). Genetic distance, heritability and correlation analysis for yield and fiber quality traits in upland cotton genotypes. *Pak. J. Biotechnol.* 14(1): 29-36.
- Panes VG, Sukhatme PV (1995). Statistical methods for agricultural workers. 3rd ICAR, New Delhi.
- Panse VG (1957). Genetics of quantitative characters in relation to plant breeding. *Indian J. Genet. Plant Breed*. 17(3): 318-328.
- Patil NP, Salve AN, Adsare AD (2017). Combining ability studies over environments for sucking pest and yield in upland cotton. *J. Global Biosci.* 6(4): 4918-4934.
- Patil YB, Madalageri BB, Biradar BD, Hoshmani RM (1996). Variability studies in okra (Abelmoschus esculentus L. Moench). Karnataka J. Agric. Sci. 9: 289–293

- Pettigrew WT (2004). Physiological consequences of moisture deficit stress in cotton. *Crop Sci*. 44: 1265-1272.
- Poehlman JM, Sleper DA (1995). Breeding field crops. 4th Ed. Panima Publishing Corporation. New Dehli, India.
- Prasad US, Reddy VC, Narisi Reddy A (2005). Studies on genetic variability in American cotton (*G. hirsutum* L.). *Karnataka J.Agric. Sci.* 18 (4): 1095-1098
- Preetha S, Raveendran TS (2007). Genetic variability and association studies in three different morphological groups of cotton (*G. hirsutum* L.). *Asian J. Plant Sci.* 6(1): 122-128.
- Pujer S, Siwach SS, Deshmukh J, Sangwan RS, Sangwan O (2014). Genetic variability, correlation and path analysis in upland cotton (*G. hirsutum* L.). *Electr. J. Plant Breed*. 5(2): 284-289.
- Radin, JW, Maune, JR, Reaves LL, French OF (1992). Yield enhancement in cotton by frequent irrigations during fruiting. *Agron J.* 84(4): 551-557.
- Ramanjinappa V, Arunkumar KH, Hugar A, Shashibhaskar MS (2011). Genetic variability in okra (*Abelmoschus esculentus L. Moench*). *Plant Arch*. 11: 435–437.
- Rao GN, Reddy MSS (2001). Studies on heritability and variability for yield and its components in*G. hirsutum* cottons. *J. Cotton Res. Dev.* 15(1): 84-86.
- Raza H, Khan NU, Khan SA, Gul S, Latif A, Hussain I, Khan J, Raza S, Baloch M (2016). Genetic variability and correlation studies in F₄ populations of upland cotton. J. Anim. Plant Sci. 26(4): 1048-1055.
- Rokadia P, Vaid B (2003). Variability parameters in American cotton (*G. hirsutum* L.). *Ann. Arid Zone* 42(1): 105-106.
- Rosenow D, Quisenberry J, Wendt C, Clark L (1983). Drought tolerant sorghum and cotton germplasm. *Agric. Water Manag.* 7:207-222
- Saleem S, Kashif M, Hussain M, Khan A, Saleem FA (2016). Genetic behavior of morpho-physiological traits and their role for breeding drought tolerant wheat. *Pak. J. Bot.* 48(3): 925-932.
- Saranga Y, Rudich J, Marani A (1991). The relations between leaf water potential of cotton plants and environmental and plant factors. *Field Crops Res.* 28: 39-46.

- Shao D, Wang T, Zhang H, Zhu J, Tang F (2016). Variation, heritability and association of yield, fiber and morphological traits in a near long staple upland cotton population. *Pak. J. Bot.* 48(5): 1945-1949.
- Shavkiev J, Hamdullaev SH, Nabiev S, Usmanov R, Bozorov T, Erjigitov D (2019a). Water sensitivity and tolerance indices upon productivity in upland cotton and other economic valuable traits. *Bull. Gulistan State Uni*. 2: 64-68.
- Shavkiev J, Nabiev S, Azimov A, Khamdullaev S, Amanov B, Matniyazova H, Nurmetov K (2020). Correlation coefficients between physiology, biochemistry, common economic traits and yield of cotton cultivars under full and deficit irrigated conditions. *J. Crit. Rev.* 7(4):131-136.
- Shavkiev J, Nabiev S, KhamdullaevSh, Usmanov R, Chorshanbiev N (2019b). Physiologic-biochemical and yield traits parameters of cotton varieties under different water irrigated regimes. *Bull. Agrarian Sci. Uzbekistan* 78(4(2): 157-162.
- Srinivas B, Bhadru D, Rao MVB, Gopinath M (2014). Genetic studies in yield and fiber quality traits in American cotton (*G. hirsutum* L.). 2014. *Agric. Sci. Digest* 34(4): 285-288.
- Steel RG., Torrie JH, Dickey DA (1997). Principles and procedures of statistics: A biometrical approach, 3rd edition. McGraw Hill Book Co. Inc. New York.

- Swarup V, Chaugale BS (1962). Studies on genetic variability in sorghum. Phenotypic variation and heritable component in some quantitative characters contributing towards yield. Indian J. Genet. Plant Breed. 22: 31-36.
- Tohir AB, Rustam MU, Yang H, Shukhrat A H, Sardorbek M, Jaloliddin S, Saidgani N, Zhang D, Alisher A A (2018). Effect of water deficiency on relationships between metabolism, physiology, biomass, and yield of upland cotton (*G. hirsutum* L.). *J. Arid Land* 10(3): 441– 456.
- Ullah I, Ashraf M, Zafar Y (2008). Genotypic variation for drought tolerance in cotton (*G. hirsutum* L): Leaf gas exchange and productivity. *Flora* -*Morph. Distribution, Funct. Ecol. Plants* 203: 105-115.
- Waqar AS, Malik MF, Akram Z (2016). Genetic variability among quality traits in cotton varieties. *Merit Res. J. Agric. Sci. Soil Sci.* 4(6): 086-089.
- Xamidov MX, Matyakubov BS (2019). Cotton irrigation regime and economical irrigation technologies. Monography Tashkent.
- Zeeshan MK, Wajid AJ, Jay KS, Muhammad IB, Adil AG, Kirshan K, M, Muhammad S, Mitho C (2020). Studies on correlation and heritability estimates in upland cotton (*G. hirsutum* L.) genotypes under the agro- climatic conditions of Tandojam, Sindh, Pakistan. *Pure Appl. Biol.* 9(4): 2272-2278.