

## PHENOTYPIC AND GENOTYPIC CORRELATION COEFFICIENTS FOR SOME YIELD AND FIBER QUALITY TRAITS OF SEGREGATING POPULATIONS (F<sub>2</sub>, F<sub>3</sub> AND F<sub>4</sub>) IN SOME EGYPTIAN COTTON CROSSES

Srour, M.S.M.\* and E. F. El-Hashash\*\*

\* Cotton Research Institute, Agricultural Research Center, Giza, Egypt.

\*\* Agronomy Dep., Fac., of Agric. Al-Azhar Univ., Cairo, Egypt

### ABSTRACT

The present study was undertaken with a view to determine the extent of relationship between seed cotton yield and other economic traits of F<sub>2</sub>, F<sub>3</sub> and F<sub>4</sub> generations in the two Egyptian cotton crosses i.e., cross I [(Giza 89 x Giza 85) x (Giza 86 x Giza 81)] x [(Giza 83 x Giza 80) x Giza 89] and cross II [(Giza 85 x Giza 86) x (Giza 83 x Giza 80) x Giza 89]. The F<sub>2</sub>, F<sub>3</sub> and F<sub>4</sub> generations in the two crosses were grown at Sakha Experimental farm, Cotton Research Institute, Agricultural Research Center, Egypt during 2008, 2009 and 2010 seasons, respectively. The mean squares obtained from analysis of variance showed highly significant ( $P \leq 0.01$ ) differences for most studied traits of F<sub>2</sub>, F<sub>3</sub> and F<sub>4</sub> generations in the two crosses. For the two crosses, the best mean values for boll weight (3.27 and 3.39 g) and 2.5% span length (33.22 and 32.84 mm) traits were detected in F<sub>4</sub> generation, seed cotton yield/plant (175.88 and 197.09 g) and uniformity index (87.01 and 86.54 %) traits in F<sub>3</sub> generation and fiber fineness (3.75 and 3.63) in F<sub>2</sub> generation. While, lint percentage (40.68 and 41.39 %) and fiber strength (10.32 and 10.44 g/tex) traits exhibited highest mean values of the crosses I and II in F<sub>4</sub> and F<sub>3</sub> generations, respectively. In general the magnitude of genotypic correlation coefficient was higher than those of phenotypic correlations coefficient of most studied traits for F<sub>2</sub>, F<sub>3</sub> and F<sub>4</sub> generations in the two crosses. Seed cotton yield for the two crosses exhibited positive and highly significant associations with lint percentage, 2.5% span length, fiber fineness and fiber strength traits in F<sub>2</sub> generation and with lint percentage and 2.5% span length traits in F<sub>3</sub> and F<sub>4</sub> generations, while, the correlation changed for fiber fineness and fiber strength in F<sub>3</sub> and F<sub>4</sub> generations. 2.5% span length, fiber fineness, fiber strength and uniformity index for the two crosses in F<sub>2</sub>, F<sub>3</sub> and F<sub>4</sub> generations had a highly significant positive correlations with boll weight. Lint percentage showed positive and highly significant associations with fiber fineness for F<sub>3</sub> generation in the cross I and with fiber fineness and strength traits for the three generations in the cross II. These results indicating that, the two crosses could be used in improving yield and fiber quality in Egyptian cotton.

**Keywords:** Phenotypic correlation, genotypic correlation, analysis of variance and F<sub>2</sub>, F<sub>3</sub> and F<sub>4</sub> generations.

### INTRODUCTION

Breeding programs continue to develop new cotton varieties to meet the requirements of both producers and consumers. The increase in yield can be possible if the existing genetic resources and information are properly utilized. Correlation between traits can be useful in developing selection criteria, but correlation can also present a morass of interrelationships (Kloth, 1998). Hybridization of two parents followed by five generations of intercrossing via insects improved yield by 9% and maintained fiber strength.

Only two cycles of inter-mating were used to reduce the correlation between lint yield and fiber strength from  $-0.54$  to  $-0.38$  (Meredith and Bridge, 1971). Improving yield and quality are important objectives for crop breeders. In cotton (*Gossypium spp.*), however, yield increases are often associated with reduced fiber quality (Meredith, 1984). Lancon *et al.*, (1993) observed negative association between fiber fineness and fiber strength. Tyagi (1994) found negative correlation of lint length with lint percentage and lint fineness. Larik *et al.*, (1997) demonstrated negative correlation for micronaire and lint length in cotton. Since there were large interactions between accessions generation for the yield and fiber traits studied. Azhar *et al.*, (2004) displayed negative correlation between staple length and fiber fineness, also between lint length and lint fineness. Gutiérrez *et al.*, (2006) mentioned that, the negative association between yield and fiber quality has hampered breeding efforts for the improvement of multiple traits. These linkages between desirable and undesirable loci can slow down genetic progress through traditional breeding programs, such as selfing and selection. It is desirable to break the linkage blocks in cotton that associate undesirable traits with high yield and quality. Ulloa (2006) reported negative correlation between lint percentage and lint strength. Single plant selection in early generations would effectively improve the seed cotton yield and its various additively controlled components (Ali *et al.*, 2008). Preetha and Raveendren (2008) stated that, the association of boll weight, seed cotton yield, lint percentage, 2.5% span length, fiber fineness, fiber strength and uniformity index traits has shown breakdown of linkage between high yield and poor quality and further the shift noticed in correlation coefficients values as generation advanced from  $F_3$  to  $F_4$  revealing unstabilized nature of the population. Basal, *et al.*, (2009) denoted that, the most important fiber quality parameters, UHM, fiber strength, and UI, were negatively associated with the most basic within-boll lint yield components, L/S, and F/S. Chaudhary *et al.*, (2010) found a wide variation in  $F_2$ ,  $F_3$  and  $F_4$  generations for fiber quality traits, especially fiber strength and also for seed cotton yield and important yield components. Transgressive segregation was observed for fiber strength, length and other traits. The objective of the present study was to estimate phenotypic and genotypic correlation coefficients between some traits for  $F_2$ ,  $F_3$  and  $F_4$  generations in two Egyptian cotton crosses (*Gossypium barbadense* L.).

## **MATERIALS AND METHODS**

### **Plant materials, growing conditions and character measurements:**

Two crosses, the cross I [(Giza 89 x Giza 85) x (Giza 86 x Giza 81)] x [(Giza 83 x Giza 80) x Giza 89] and cross II [(Giza 85 x Giza 86) x (Giza 83 x Giza 80) x Giza 89] were developed for this study.  $F_{1s}$  derived from the two intra-specific crosses were self-pollinated to developed  $F_2$  populations. One hundred single plants from  $F_2$  generation were phenotype for fiber quality traits as well as seed cotton yield and some important yield components. These single plants were self-pollinated to advance to further filial generations up to  $F_4$  generation which constituted the material for the study.  $F_2$ ,  $F_3$  and  $F_4$  generations were raised along with the parents at Sakha

experimental farm, Cotton Research Institute, Agricultural Research Center, Egypt during 2008, 2009 and 2010 seasons, respectively. The experimental design used in the three seasons was a randomized complete blocks design (RCBD) with three replications. The parents were grown in two row plots; the  $F_2$ 's were raised in 10 rows,  $F_3$ 's and  $F_4$ 's in 5 row plots. Each row was 4 m long and 0.6 m wide. Hills were spaced at 0.4 m and thinned at one plant/hill. Selected plants in each single plant progeny were observed on their biometrical and fiber quality traits and recorded. All the recommended cultural practices of cotton production in the area were done. Measurements were taken on seven traits including boll weight (B.W. g), seed cotton yield/plant (S.C.Y. /P g), lint percentage (L. %), fiber length (mm) at 2.5 % span length (2.5 % S.L.), fiber fineness (F.F.), fiber strength (F.S. g/tex) and uniformity index (U.I. %). The fiber properties were measured using HVI according to (ASTMD – 4650 – 86) at the Lab. of Cotton Research Institute, Agricultural Research Center.

**Statistical analysis:**

The averages for each character were subjected to analysis of variance and covariance following the method of Sing and Chaudhary (1985). In all generations, the phenotypic and genotypic correlation coefficients were calculated to determine the degree of association among the different characters in the two Egyptian cotton crosses. The estimates of phenotypic and genotypic correlation coefficients were worked out by using the formulae suggested by Edhaie *et al.* (1993). Both genotypic and phenotypic correlation coefficients were compared against table r-values given in Fisher and Yates (1953) table at (n-2) degrees of freedom at the probability levels of 0.05 and 0.01 to test their significance.

## **RESULTS AND DISCUSSION**

The estimates of correlation among traits are useful for planning a breeding programme to synthesize a genotype with desirable traits. The objective of the present study was to find correlation among the seed cotton yield trait and traits related to fiber in cotton.

**I- Analysis of variance:**

Results mean squares for  $P_1$ ,  $P_2$ ,  $F_2$ ,  $F_3$  and  $F_4$  generations in the two crosses are Table 1 which demonstrated that, the two parents in the two crosses were insignificant traits except seed cotton yield/plant for  $P_1$  in the cross II and 2.5% span length for  $P_2$  in the two crosses which were exhibited highly significant differences ( $P \leq 0.01$ ). On the other hand, boll weight, seed cotton yield/plant and lint percentage traits showed highly significant differences ( $P \leq 0.01$ ) of  $F_2$ ,  $F_3$  and  $F_4$  generations in the two crosses. As for 2.5% S.L. of  $F_2$  generation in the cross I and  $F_3$  generation in the cross II were highly significant and significant, respectively. However, mean squares for fiber fineness were highly significant for the three segregating generations in the two crosses except  $F_2$  generation in the cross II which was insignificant. In the same time, the mean squares of  $F_2$  and  $F_4$  generations in the cross I and  $F_3$  generation in the cross II were displayed highly significant differences for fiber strength. Finally, the uniformity index demonstrated

highly significant differences for F<sub>2</sub> and F<sub>3</sub> generations in the cross II and the cross I, respectively. Meredith (1990) reported that F<sub>2</sub> hybrids had significantly longer and finer lint than the parents; however, the improvements were too small to be of practical value. He suggested that F<sub>2</sub> hybrids have the genetic potential for increasing cotton yields and fiber quality. McCarty *et al.*, (2003) stated that, the analysis of variance for seed cotton yield, lint percentage, 2.5% span length, fiber fineness, fiber strength and uniformity index were significantly different in F<sub>2</sub> and F<sub>3</sub> generations. Srouf *et al.*, (2010) mentioned that, the analysis of variance for all traits studied manifested highly significant differences of F<sub>3</sub> generations in the two crosses, while, the F<sub>2</sub> generation exhibited highly significant for seed cotton yield/plant and uniformity index in the first cross, and boll weight, seed cotton yield/plant and uniformity index in the second cross.

**Table 1: Mean squares for seven traits of P<sub>1</sub>, P<sub>2</sub>, F<sub>2</sub>, F<sub>3</sub> and F<sub>4</sub> populations in the two Egyptian cotton crosses.**

Traits	Generations	P <sub>1</sub>	P <sub>2</sub>	F <sub>2</sub>	F <sub>3</sub>	F <sub>4</sub>
	Crosses					
B.W. (g)	Cross I	0.14	0.06	6.56**	0.17**	0.26**
	Cross II	0.05	0.06	0.26**	0.35**	0.31**
S.C.Y./P (g)	Cross I	1400.86	361.55	6660.48**	14213.55**	1095.01**
	Cross II	1687.45**	361.55	7861.72**	20386.44**	3696.99**
L. %	Cross I	0.57	3.84	8.15**	6.89**	4.68**
	Cross II	1.79	3.84	15.63**	4.76**	5.33**
2.5% S.L. (mm)	Cross I	0.25	0.45**	0.41**	22.04	0.59
	Cross II	0.49	0.45**	7.61	1.17*	0.29
F.F.	Cross I	0.05	0.13	0.38**	0.18**	0.06**
	Cross II	0.15	0.13	0.18	0.30**	0.56**
F.S. (g/tex)	Cross I	0.04	0.05	0.92**	0.35	0.29**
	Cross II	0.24	0.05	0.52	0.70**	0.05
U.I. (%)	Cross I	0.11	0.11	0.35	1.58**	0.55
	Cross II	0.96	0.11	1.49**	0.88	0.53

\* and \*\*: The shifts were significant and highly significant, respectively.

## II- Mean performance:

Data on mean in respect of fiber quality traits, seed cotton yield and important yield components in P<sub>1</sub>, P<sub>2</sub>, F<sub>2</sub>, F<sub>3</sub> and F<sub>4</sub> generations in the two crosses are given in Table 2. According to these obtained data, the F<sub>4</sub> generations had the highest means for boll weight and 2.5% span length traits in the two crosses (3.27 g – 3.39 g and 33.22 mm – 32.84 mm, respectively), and for lint percentage and fiber strength traits (40.68% - 10.32 g/tex, respectively) in the cross I. While, the maximum seed cotton yield/plant and uniformity index traits for the two crosses (175.88 g – 197.09 g and 87.01 % – 86.54 %, respectively) and the highest mean lint percentage (41.39 %) and fiber strength (10.44 g/tex) for the cross II were counted in F<sub>3</sub> generations. On the other hand, the best mean fiber fineness values of 3.75 – 3.63 were observed for F<sub>2</sub> generation in the two crosses, respectively. Data presented in Table 1 showed that, the cross II was superior for boll weight, seed cotton yield/plant, lint percentage and fiber fineness traits in five

populations, but, the cross I was superior for 2.5% S.L. and uniformity index traits in P<sub>1</sub>, P<sub>2</sub>, F<sub>2</sub>, F<sub>3</sub> and F<sub>4</sub> populations. Preetha and Raveendren (2008) reported that, the mean performance of F<sub>3</sub> and F<sub>4</sub> generations for boll weight, seed cotton yield, lint percentage, 2.5 percent span length, bundle strength, uniformity ratio and fiber fineness traits were found to be intermediate between parental values.

**III- Genotypic and phenotypic correlation coefficients:**

Selection for specific character is known to result in correlated response in certain other characters (Falconer, 1981). Generation, satisfactions progress in breeding programme depends much on the genetic variability of different traits under selection. Seed yield is the most important target character in most of the crop species and it is the most complex one. Improvement in the seed yield can be achieved by indirect selection through other easily observable characters. But this needs a good understanding of the association of different traits with seed yield and their possible association among themselves.

**Table 2: Means of seven traits in P<sub>1</sub>, P<sub>2</sub>, F<sub>2</sub>, F<sub>3</sub> and F<sub>4</sub> generations of the two Egyptian cotton crosses.**

Traits	Generations	P <sub>1</sub>	P <sub>2</sub>	F <sub>2</sub>	F <sub>3</sub>	F <sub>4</sub>
	Crosses					
B.W. (g)	Cross I	2.97	3.17	2.65	2.87	3.27
	Cross II	2.97	3.17	2.76	3.13	3.39
S.C.Y./P (g)	Cross I	100.85	114.07	107.03	175.88	102.45
	Cross II	77.89	114.07	126.92	197.09	130.76
L. %	Cross I	37.24	39.79	35.59	39.05	40.68
	Cross II	35.44	39.79	39.84	41.39	30.9
2.5% S.L. (mm)	Cross I	32.95	31.66	32.65	32.16	33.22
	Cross II	32.25	31.66	30.9	32.73	32.84
F.F.	Cross I	4.40	4.65	3.75	4.24	4.44
	Cross II	4.03	4.65	3.63	4.23	4.52
F.S. (g/tex)	Cross I	10.13	10.23	10.16	10.11	10.32
	Cross II	9.90	10.43	9.89	10.44	10.18
U.I. (%)	Cross I	86.37	85.81	84.59	87.01	86.83
	Cross II	85.95	85.81	83.55	86.54	86.13

The Genotypic and phenotypic correlation coefficients of seed cotton yield with other quantitative characters for the two crosses in different populations are presented in Tables 3, 4, 5, 6, 7 and 8. The genotypic correlation coefficients ( $r_g$ ) were higher as compared to phenotypic correlation coefficients ( $r_p$ ) in most of the traits indicating the effects of environment suppressed the phenotypic relationship between these traits. In few cases, moreover, phenotypic correlation coefficients were same with or higher than their genotypic correlation coefficients suggesting that both environmental and genotypic correlations in these cases acted in the same direction and finally maximize their expression at phenotypic level. Hussain *et al.*, (2009) denoted that phenotypic correlation was usually different in magnitude or even in direction as compared with the correlation of component effects.

**A- Cross I:**

A general observation of data (Tables 3, 4 and 5) revealed that in segregating generation's seed cotton yield/plant showed a significant or highly significant associations with lint percentage ( $r_p = 0.30$ ), 2.5% S.L. ( $r_g = -0.55$  and  $r_p = -0.51$ ), fiber fineness ( $r_g = -0.39$  and  $r_p = -0.30$ ) and uniformity index ( $r_g = -0.99$ ) in  $F_2$  generation, with lint percentage ( $r_g = 0.27$ ), fiber strength ( $r_g = 1.00$  and  $r_p = 0.65$ ) and uniformity index ( $r_g = 0.51$  and  $r_p = 0.43$ ) in  $F_3$  generation and with lint percentage ( $r_g = 0.44$ ), 2.5% S.L. ( $r_g = 0.56$  and  $r_p = 0.27$ ), fiber fineness ( $r_g = -0.56$  and  $r_p = -0.33$ ), fiber strength ( $r_g = -0.97$  and  $r_p = -0.58$ ) and uniformity index ( $r_g = -0.51$  and  $r_p = -0.35$ ) in  $F_4$  generation.

Boll weight had highly significant positive correlation with 2.5% S.L. ( $r_g = 0.41$ ), fiber strength ( $r_g = 0.40$  and  $r_p = 0.29$ ) and uniformity index ( $r_g = 1.00$ ) in  $F_2$  generation, with 2.5% S.L. ( $r_g = 1.12$  and  $r_p = 0.37$ ), fiber strength ( $r_g = 0.80$  and  $r_p = 0.46$ ) and uniformity index ( $r_g = 0.56$  and  $r_p = 0.43$ ) in  $F_3$  generation and with fiber fineness ( $r_g = 0.44$  and  $r_p = 0.39$ ) in  $F_4$  generation. However, it was significant or highly significant negatively correlated with lint percentage ( $r_g = -0.33$ ) in  $F_3$  generation and with lint percentage ( $r_g = -0.55$  and  $r_p = -0.45$ ), fiber strength ( $r_g = -0.44$  and  $r_p = -0.39$ ) and uniformity index ( $r_g = -0.51$  and  $r_p = -0.34$ ) in  $F_4$  generation.

**Table 3: Estimates of genotypic ( $r_g$ ) and phenotypic ( $r_p$ ) correlation coefficients among seven traits for  $F_2$  generation in cross I.**

Traits	Traits	B.W.	L. %	2.5%S.L.	F.F.	F.S.	U.I
	Parameters						
S.C.Y.	$r_g$	-0.09	0.10	-0.55**	-0.39**	0.04	-0.99**
	$r_p$	-0.03	0.30**	-0.51**	-0.30**	0.10	0.06
B.W.	$r_g$		0.08	0.41**	0.08	0.40**	1.00**
	$r_p$		0.07	0.17	0.06	0.29**	0.10
L. %	$r_g$			-0.46**	-0.18	-0.05	-1.00**
	$r_p$			-0.13	-0.14	-0.02	-0.15
2.5%S.L.	$r_g$				-1.18**	0.61**	1.50**
	$r_p$				-0.13	0.24**	0.53**
F.F.	$r_g$					0.09	0.99**
	$r_p$					0.10	0.45**
F.S.	$r_g$						1.11**
	$r_p$						0.29**

\* and \*\*: The shifts were significant and highly significant, respectively.

**Table 4: Estimates of genotypic ( $r_g$ ) and phenotypic ( $r_p$ ) correlation coefficients among seven traits for  $F_3$  generation in cross I.**

Traits	Traits						
	Parameters	B.W.	L. %	2.5%S.L.	F.F.	F.S.	U.I
S.C.Y.	$r_g$	0.26	0.27*	0.14	0.23	1.00**	0.51**
	$r_p$	0.24	0.26	0.06	0.20	0.65**	0.43**
B.W.	$r_g$		-0.33*	1.12**	0.11	0.80**	0.56**
	$r_p$		-0.25	0.37**	0.17	0.46**	0.43**
L. %	$r_g$			0.10	0.44**	0.01	0.15
	$r_p$			0.04	0.37**	0.09	0.06
2.5%S.L.	$r_g$				-0.27	1.50**	1.33**
	$r_p$				-0.10	0.63**	0.54**
F.F.	$r_g$					0.21	0.23
	$r_p$					0.08	0.14
F.S.	$r_g$						1.19**
	$r_p$						0.67**

\* and \*\*: The shifts were significant and highly significant, respectively.

**Table 5: Estimates of genotypic ( $r_g$ ) and phenotypic ( $r_p$ ) correlation coefficients among seven traits for  $F_4$  generation in cross I.**

Traits	Traits						
	Parameters	B.W.	L. %	2.5%S.L.	F.F.	F.S.	U.I
S.C.Y.	$r_g$	0.22	0.44**	0.56**	-0.56**	-0.97**	-0.51**
	$r_p$	0.09	0.26	0.27*	-0.33*	-0.58**	-0.35**
B.W.	$r_g$		-0.55**	-0.15	0.44**	-0.44**	-0.51**
	$r_p$		-0.45**	-0.12	0.39**	-0.39**	-0.34*
L. %	$r_g$			-0.65**	-0.82**	-0.21	0.07
	$r_p$			-0.38**	-0.59**	-0.07	0.01
2.5%S.L.	$r_g$				-0.03	-0.37**	1.34**
	$r_p$				-0.18	-0.26	0.55**
F.F.	$r_g$					0.39**	0.38**
	$r_p$					0.35**	0.29**
F.S.	$r_g$						0.39**
	$r_p$						0.19

\* and \*\*: The shifts were significant and highly significant, respectively.

The genotypic and phenotypic correlation coefficients indicated that lint percentage had positive or negative and highly significant associations with 2.5% S.L. ( $r_g = -0.46$ ) and uniformity index ( $r_g = -1.00$ ) in  $F_2$  generation, with fiber fineness ( $r_g = 0.44$  and  $r_p = 0.37$ ) in  $F_3$  generation and with 2.5% S.L. ( $r_g = -0.65$  and  $r_p = -0.38$ ) and fiber fineness ( $r_g = -0.82$  and  $r_p = -0.59$ ) in  $F_4$  generation.

Results indicate that fiber strength and uniformity index was correlated in  $F_2$  [ $(r_g = 0.61$  and  $r_p = 0.24)$  and ( $r_g = 1.50$  and  $r_p = 0.53$ ), respectively] and in  $F_3$  [ $(r_g = 1.50$  and  $r_p = 0.63)$  and ( $r_g = 1.33$  and  $r_p = 0.54$ ), respectively]. Meanwhile, uniformity index ( $r_g = 1.34$  and  $r_p = 0.55$ ) in  $F_4$  generation had positive and highly significant correlations at genotypic and phenotypic level with 2.5% span length. However 2.5% span length

demonstrated negative highly significant associations with fiber fineness ( $r_g = -1.18$ ) in  $F_2$  generation and fiber strength ( $r_g = -0.37$ ) in  $F_4$  generation.

The positive and highly significant associations were recorded between fiber fineness and uniformity index in  $F_2$  generation ( $r_g = 0.99$  and  $r_p = 0.45$ ) and  $F_4$  generation ( $r_g = 0.38$  and  $r_p = 0.29$ ), fiber fineness and fiber strength ( $r_g = 0.39$  and  $r_p = 0.35$ ) in  $F_4$  generation, and fiber strength and uniformity index in  $F_2$  generation ( $r_g = 1.11$  and  $r_p = 0.29$ ),  $F_3$  generation ( $r_g = 1.19$  and  $r_p = 0.67$ ) and  $F_4$  generation ( $r_g = 0.39$ ).

**B- Cross II:**

The phenotypic ( $r_p$ ) and genotypic ( $r_g$ ) correlation coefficients (Tables 6, 7 and 8), indicated that the seed cotton yield/plant had positive and highly significant with lint percentage ( $r_p = 0.29$ ), 2.5% span length ( $r_p = 0.27$ ), fiber fineness ( $r_g = 0.52$  and  $r_p = 0.27$ ) and fiber strength ( $r_g = 0.47$  and  $r_p = 0.35$ ) in  $F_2$  generation and with 2.5% span length ( $r_g = 0.66$ ) in  $F_4$  generation. However, seed cotton yield/plant was highly significant and negatively correlated with uniformity index ( $r_g = -1.17$ ) in  $F_2$  generation, with lint percentage ( $r_g = -0.52$  and  $r_p = -0.49$ ), 2.5% span length ( $r_g = -0.51$  and  $r_p = -0.38$ ), fiber fineness ( $r_g = -0.73$  and  $r_p = -0.65$ ) and fiber strength ( $r_g = -0.55$  and  $r_p = -0.48$ ) in  $F_3$  generation and with lint percentage ( $r_g = -0.53$  and  $r_p = -0.52$ ) and uniformity index ( $r_g = -0.45$ ) in  $F_4$  generation. While, negative and significant associations were found between seed cotton yield/plant and boll weight ( $r_g = -0.29$  and  $r_p = -0.29$ ) in  $F_3$  generation and with uniformity index ( $r_p = -0.33$ ) in  $F_4$  generation.

Boll weight revealed positive and significant or highly significant correlations with fiber fineness ( $r_g = 0.27$ ) and uniformity index ( $r_p = 0.22$ ) in  $F_2$  generation, with 2.5% span length ( $r_g = 0.66$  and  $r_p = 0.48$ ), fiber strength ( $r_g = 0.38$  and  $r_p = 0.31$ ) and uniformity index ( $r_g = 0.71$  and  $r_p = 0.48$ ) in  $F_3$  generation and with fiber strength ( $r_g = 0.33$ ) in  $F_4$  generation. However, boll weight was found to be negatively and highly significantly correlated with fiber fineness ( $r_g = -0.49$  and  $r_p = -0.48$ ) and uniformity index ( $r_g = -0.93$  and  $r_p = -0.66$ ) in  $F_4$  generation.

**Table 6: Estimates of genotypic ( $r_g$ ) and phenotypic ( $r_p$ ) correlation coefficients among seven traits for  $F_2$  generation in cross II.**

Traits	Traits						
	Parameters	B.W.	L. %	2.5%S.L.	F.F.	F.S.	U.I
S.C.Y.	$r_g$	0.06	-0.00	-0.02	0.52**	0.47**	-1.17**
	$r_p$	0.11	0.29**	0.27**	0.27**	0.35**	-0.06
B.W.	$r_g$		0.14	0.12	0.27**	0.12	-0.14
	$r_p$		0.16	0.11	0.07	0.10	0.22*
L. %	$r_g$			0.10	0.60**	0.31**	-0.39**
	$r_p$			0.04	0.19*	0.21*	-0.09
2.5%S.L.	$r_g$				0.02	0.31**	-0.64**
	$r_p$				0.03	0.08	-0.00
F.F.	$r_g$					0.01	-0.15
	$r_p$					-0.05	-0.01
F.S.	$r_g$						-1.17**
	$r_p$						0.03

\* and \*\*: The shifts were significant and highly significant, respectively.



**Table 7: Estimates of genotypic ( $r_g$ ) and phenotypic ( $r_p$ ) correlation coefficients among seven traits for  $F_3$  generation in cross II.**

Traits	Traits						
	Parameters	B.W.	L. %	2.5%S.L.	F.F.	F.S.	U.I
S.C.Y.	$r_g$	-0.29*	-0.52**	-0.51**	-0.73**	-0.55**	0.11
	$r_p$	-0.29*	-0.49**	-0.38**	-0.65**	-0.48**	0.07
B.W.	$r_g$		0.25	0.66**	-0.00	0.38**	0.71**
	$r_p$		0.24	0.48**	0.00	0.31*	0.48**
L. %	$r_g$			0.22	0.49**	0.37**	-0.01
	$r_p$			0.19	0.41**	0.29*	0.04
2.5%S.L.	$r_g$				0.39**	0.46**	1.02**
	$r_p$				0.34*	0.39**	0.70**
F.F.	$r_g$					0.13	-0.41**
	$r_p$					0.16	-0.16
F.S.	$r_g$						0.51**
	$r_p$						0.39**

\* and \*\*: The shifts were significant and highly significant, respectively.

**Table 8: Estimates of genotypic ( $r_g$ ) and phenotypic ( $r_p$ ) correlation coefficients among seven traits for  $F_4$  generation in cross II.**

Traits	Traits						
	Parameters	B.W.	L. %	2.5%S.L.	F.F.	F.S.	U.I
S.C.Y.	$r_g$	-0.05	-0.53**	0.66**	-0.07	0.13	-0.45**
	$r_p$	-0.05	-0.52**	0.26	-0.07	0.08	-0.33*
B.W.	$r_g$		0.21	-0.13	-0.49**	0.33*	-0.93**
	$r_p$		0.20	-0.06	-0.48**	0.23	-0.66**
L. %	$r_g$			-1.33**	0.39**	-0.62**	-0.04
	$r_p$			0.54**	0.37**	-0.38**	-0.05
2.5%S.L.	$r_g$				-0.77**	1.59**	-1.47**
	$r_p$				-0.28*	0.31*	-0.28*
F.F.	$r_g$					-0.83**	0.78**
	$r_p$					-0.49**	0.53**
F.S.	$r_g$						0.41**
	$r_p$						0.05

\* and \*\*: The shifts were significant and highly significant, respectively.

Lint percentage showed positively significant or highly significant associations with fiber fineness [( $r_g = 0.60$  and  $r_p = 0.19$ ) and ( $r_g = 0.49$  and  $r_p = 0.41$ )] and fiber strength [( $r_g = 0.21$  and  $r_p = 0.31$ ) and ( $r_g = 0.37$  and  $r_p = 0.29$ )] in  $F_2$  and  $F_3$  generations, respectively, and with 2.5% span length ( $r_p = 0.54$ ) and fiber fineness ( $r_g = 0.39$  and  $r_p = 0.37$ ) in  $F_4$  generation. On the other hand, lint percentage observed negative and highly significant associations with uniformity index ( $r_g = -0.39$ ) in  $F_2$  generation and with 2.5% span length ( $r_g = -1.33$ ) and fiber strength ( $r_g = -0.62$  and  $r_p = -0.38$ ) in  $F_4$  generation.

2.5% span length exhibited significant or highly significant positive associations with fiber strength in  $F_2$  ( $r_g = 0.31$ ) and  $F_4$  ( $r_g = 1.59$  and  $r_p = 0.31$ ) generations and fiber fineness ( $r_g = 0.39$  and  $r_p = 0.34$ ), fiber strength

( $r_g = 0.46$  and  $r_p = 0.39$ ) and uniformity index ( $r_g = 1.02$  and  $r_p = 0.70$ ) in  $F_3$  generation. While, 2.5% span length displayed significant or highly significant negative association with uniformity index ( $r_g = -0.64$ ) in  $F_2$  generation and with fiber fineness ( $r_g = -0.77$  and  $r_p = -0.28$ ) and uniformity index ( $r_g = -1.47$  and  $r_p = -0.28$ ) in  $F_4$  generation.

Highly significant associations were observed between fiber fineness and uniformity index ( $r_g = -0.41$ ) in  $F_3$  generation, fiber fineness and fiber strength ( $r_g = -0.83$  and  $r_p = -0.49$ ) in  $F_4$  generation, fiber strength and uniformity index ( $r_g = -1.17$ ) in  $F_2$  generation, fiber strength and uniformity index ( $r_g = 0.51$  and  $r_p = 0.39$ ) in  $F_3$  generation, fiber fineness and uniformity index ( $r_g = 0.78$  and  $r_p = 0.53$ ) in  $F_4$  generation and fiber strength and uniformity index ( $r_g = 0.41$ ) in  $F_4$  generation.

In other studies, Lancon *et al.*, (1993) observed negative association between fiber fineness and fiber strength. Tyagi (1994) observed positive association between lint percentage with seed cotton yield and fiber fineness traits in their studies. While, negative associations were found between fiber length and seed cotton yield, lint percentage and fiber fineness traits. Shah (1995) revealed the existence of negative associations among boll weight and lint percentage traits. Azhar and Hussain (1998) reported positive association between seed cotton yield and boll weight. Kloth (1998) mentioned that correlations between fiber fineness and fiber strength were found at the phenotypic level, but were non-existent at the genotypic level. Both elongation and strength were correlated genotypically, but not phenotypically. Most traits were highly correlated in  $F_2$  and  $F_3$  generations; however, seed cotton yield and lint yield were not correlated in  $F_2$  and  $F_3$  (McCarty *et al.*, 2003). Azhar *et al.*, (2004) detected negative correlation between staple length and fiber fineness. Naveed *et al.*, (2004) stated that association among boll weight and lint percentage was negative and similarly boll weight and lint percentage were found to have negative association with seed cotton yield. Rauf *et al.*, (2004) revealed that boll weight had non-significant correlation at genotypic level but negative and significant correlation at phenotypic level with seed cotton yield. Positive association of fiber fineness and uniformity index, and negative association of uniformity index and fiber strength were in accordance with the results of Zhang and Xiao (2005). The study of phenotypic correlation indicated significant positive association of seed cotton yield per plant with number of bolls per plant, boll weight and lint yield per plant in  $F_2$  generation (Basamma, 2007). Preetha and Raveendren (2008) noticed significant positive association of boll weight with seed cotton yield, and fiber length has shown significant negative association with fiber fineness, fiber strength and uniformity index traits for  $F_3$  and  $F_4$  generations. They added that, positive associations was significant boll weight and fiber length in  $F_3$  generation but not in  $F_4$  generation, and the significant negative association of uniformity index and fiber strength was observed in  $F_4$  generation. Hussain *et al.*, (2009) reported that a significant positive association with boll weight and seed cotton yield was found. Significant genotypic variation for lint yield and fiber quality including fiber bundle strength, span lengths, short fiber and fineness in the populations were found (Zeng *et al.*, 2007 and Zeng and Meredith, 2009).

In general, the significant positive or negative inter-relationships of the two crosses in  $F_2$ ,  $F_3$  and  $F_4$  generations showed that, there were good associations among traits, reflecting the effectiveness of indirect selection by breeders while making selection of desirable plants among segregating populations. The association between seed cotton yield and lint percentage was either positively significant or just positive of the cross I in  $F_2$ ,  $F_3$  and  $F_4$  generations. In any case their correlation was regarded as favorable one as it indicates the possibility of having simultaneous improvements in both the traits and consequently yield per plant. A strong associations between uniformity index with boll weight, 2.5% span length and fiber strength of the cross I in the three generations, suggested that boll weight, 2.5% span length and fiber strength can be improved indirectly by selecting plants with higher uniformity index per plant. Boll weight was also important factor for improving seed cotton yield and lint percentage and 2.5% span length. Association between seed cotton yield and fiber fineness was not strong as evident from correlation coefficient values except the  $F_3$  generation in the cross II.

### **CONCLUSION**

The genotypic and phenotypic correlation coefficients among different pairs of plant traits for the two crosses in  $F_2$ ,  $F_3$  and  $F_4$  generations indicated that seed cotton yield can be improved by increasing lint percentage and boll weight traits. The results revealed a possibility of selecting plants with desirable attributes of fiber fineness and seed cotton yield in the next segregating generations. Positive genetic associations between staple length and fiber strength of the two crosses indicated that selection for increased value of one trait will result an increase in value of other. The negative correlation between yield and quality traits were changed to positive in cross combination like indicating the possibility of improving yield and quality traits simultaneously.

### **REFERENCES**

- Ali, M.A.; Khan, I.A.; Awan , S.I.; Ali, S. and Niaz, S. (2008). Genetics of fiber quality traits in cotton (*Gossypium hirsutum* L.). *Aust. J. Crop Sci.*, 2: 10-17.
- A.S.T.M, D. 1448-59, 1967, A.S.T.M, D.1445-67 and (A.S.T.M. Designation D- 1447-67).
- Azhar, F.M.; Hussain, S.S. and Khan, I.A. (1998). Association of seed cotton yield with other quantitative plant characters of *Gossypium hirsutum* L. *Pakistan. J. Bio.Sci.*, 2: 700-701.
- Azhar, F.M.; Naveed, M. and Ali, A. (2004). Correlation analysis of seed cotton yield with lint characteristics in *Gossypium hirsutum* L. *International J. Agri. Biology*, 6:656-658.
- Basal, H.; Unay, A.; Canavar, O. and Yavas, I. (2009). Combining ability for fiber quality parameters and within-boll yield components in intraspecific and interspecific cotton populations. *Spain. J. Agric. Res.*, 7(2), 364-374.

- Basamma, K. (2007). Genetic variability in selected F<sub>2</sub> populations of Desi cotton. M.Sc. Thesis, Fac. of Agric., Dharwad Univ., Dharwad.
- Chaudhary, B.; Singh, J. and Chopra, S.K. (2010). Development of recombinant inbred lines for fibre strength and other important traits in cotton (*Gossypium hirsutum*, L.). *Indian J. of Agric. Sci.*, 80 (5): 357–539.
- Edhaie, B.; Barnhast, D. and Waines, J.G. (1993). Genetic analysis of transpiration efficiency, carbon isotopes discrimination and growth characters in bread wheat. In: *Staple isotopes and plant carbon water relations* (J.R. Enleringer, A.E. Hall and G.D. Farguner Ed.) Academic press, London pp: 419-434.
- Falconer, D. S. (1981). *Introduction of Quantitative Genetics*, Longman Inc. Ltd., New York, p. 340.
- Fisher, R.A and Yates, F. (1953). *Statistical Tables for Biological, Agricultural, and Medical Research*. Olive and Boyd, Edinburgh, p.94
- Gutiérrez, O.A.; Daryl, T.B.; Clay, B.C.; Johnie, N.J.; Jack, C.; Jixiang, W. and Clarence, E.W. (2006). Development of Random-mated Populations Using Bulked Pollen Methodology. *Cotton as a Model. The J. of Cotton Science* 10:175–179.
- Hussain, M.; Azhar, F.M. and A.A. Khan (2009). Genetics of inheritance and correlations of some morphological and yield contributing traits in upland cotton. *Pak. J. Bot.*, 41(6): 2975-2986.
- Kloth, R.H. (1998). Analysis of commonality for traits of cotton fiber. *The J. of Cotton Sci.*, 2:17-22.
- Lancon, J.E.; Goze, B. Hau; Bachelier, M. and Chanselme, J.L. (1993). Multisite trial of diallel with four elite parents. III. Correlation between variables. *Cotton Fib. Trop.* 48:11-14.
- Larik, A.S.; Ansari, S.R. and Kumbhar, M.B. (1997). Heritability analysis of yield and quality components in *Gossypium hirsutum* L. *Pak. J. Bot.* 29: 97-101.
- Meredith, W.R., Jr. (1984). Quantitative genetics. p. 132-147. In R. J. Kohel and C.F. Lewis (ed.) *Cotton. Agron. Monogr.* 24. ASA, CSSA, and SSSA, Madison, WI.
- Meredith, W.R., Jr. (1990). Yield and fiber-quality potential for second-generation cotton hybrids. *Crop Sci.* 30:1045-1048.
- Meredith, W.R., Jr., and Bridge, R.R. (1971). Breakup of linkage blocks in cotton, *Gossypium hirsutum* L. *Crop Sci.*, 11: 695-698.
- McCarty, J.C., Jr.; Jenkins, J.N. and Wu, J. (2003). Use of primitive accessions of cotton as sources of genes for improving yield components and fiber properties. *Mississippi Agricultural & Forestry Experiment Station*, March 2003: 1 – 20.
- Naveed, M.; Azhar, F.M. and Ali, A. (2004). Estimates of Heritabilities and Correlations Among Seed Cotton Yield and its Components in *Gossypium hirsutum* L. *Int. J. Agri. Biol.*, 6(4), 2004
- Preetha, S. and Raveendren, T.S. (2008). Genetic appraisal of yield and fiber traits in cotton using interspecific F<sub>2</sub>, F<sub>3</sub>, and F<sub>4</sub> population. *Inter. J. of Integrative Biology*, 3(2): 136 – 142.

- Rauf, S.; Khan, T. M.; Sadaqat, H. A. and Khan, A. I. (2004). Correlation and path coefficient analysis of cotton yield. *Int. J. Agri. Biol.*, 6(4): 686 – 688.
- Shah, S.A.H., (1995). Path coefficient and correlation studies in Upland cotton (*G. hirsutum* L.). M.Sc. (Hons.) Thesis. Department of Plant Breeding and Genetics, University of Agriculture Faisalabad
- Singh, R.K and Chawdhury, B.D. (1985). Biometrical methods in quantitative genetic analysis, Kalyani Publications, New Delhi.
- Srour, M.S.M.; Hager, M.A.; Zaazaa, E.I. and El-Hashash, E. F. (2010). Genetic analysis for some yield and fiber quality traits using  $F_2$  and  $F_3$  populations in cotton. *J. Plant Prod., Mansoura Univ.*, 1 (12): 1593 – 1604.
- Tyagi, A.P. (1994). Correlation coefficients and selection indices in upland cotton (*Gossypium hirsutum* L.). *Ind. J. Agric. Res.* 28(3): 189-196.
- Ulloa, M. (2006). Heritability and correlation of agronomic and lint traits in an okra leaf Upland cotton population. *Crop Sci.* 46: 1508-1514.
- Zeng, L., and Meredith, Jr. W.R. (2009). Associations among lint yield, yield components, and fiber properties in an introgressed population of cotton. *Crop Sci.* 49:1647-1654.
- Zeng, L.; Meredith, Jr. W.R.; Boykin, D.L. and Taliercio, E. (2007). Evaluation of an exotic germplasm population derived from multiple crosses among *Gossypium* tetraploid species. *J. Cotton Sci.* 11:118-127.
- Zhang, Z.S. and Xiao, Y.H. (2005). Construction of a genetic linkage map and QTL analysis of fiber related traits in Upland cotton. *Euphytica*, 144: 91 – 99.

معاملات الارتباط الظاهري والوراثي لبعض صفات المحصول وجودة الألياف  
للأجيال الانعزالية (الثاني، الثالث و الرابع) في بعض هجن القطن المصري  
محمود سرور محمود سرور\* و عصام فتحي الحشاش\*\*  
\* معهد بحوث القطن – مركز البحوث الزراعية – الجيزة – مصر.  
\*\* قسم المحاصيل – كلية الزراعة بالقاهرة – جامعة الأزهر – مصر.

أقيمت هذه التجربة لدراسة العلاقة بين المحصول وبعض الصفات الاقتصادية الأخرى  
للأجيال الثاني والثالث والرابع في هجينين من القطن المصري، الهجين الأول [(جيزة 89 x جيزة  
85)x(جيزة 86 x جيزة 81)]x(جيزة 83 x جيزة 80 x جيزة 89)] و الهجين الثاني [(جيزة  
85 x جيزة 86)x(جيزة 83 x جيزة 80 x جيزة 89)]. تم زراعة الأجيال الثاني والثالث  
والرابع لكلاً من الهجينين بمزرعة التجارب بسخا – معهد بحوث القطن – مركز البحوث الزراعية  
– مصر خلال مواسم 2008 – 2009 – 2010 على التوالي. وأوضحت النتائج وجود اختلافات  
عالية المعنوية لمعظم الصفات محل الدراسة للثلاثة أجيال في الهجينين. أن أفضل قيم للمتوسطات  
وجدت لمتوسط وزن اللوزة (3.27 و 3.39 جم) و طول التيلة عند 2.5% (33.22 و 32.84 مم)  
في الجيل الرابع و محصول القطن الزهر/نبات (175.88 و 197.09 جم) و معدل الانتظام  
(87.01 و 86.54%) في الجيل الثالث و لنعومة التيلة (3.75 و 3.63) في الجيل الثاني، بينما  
أظهرت صفتي معدل الحليج (40.68 و 41.39%) و متانة التيلة (10.32 و 10.44 جم/تكس) قيم  
عالية للمتوسط للهجينين الأول والثاني في الجيلين الرابع والثالث على التوالي. وكانت قيم معامل  
الارتباط الوراثي أعلى من قيم معامل الارتباط الظاهري لمعظم الصفات المدروسة للأجيال الثاني  
والثالث والرابع في كلا الهجينين. أظهرت صفة المحصول لكل من الهجينين ارتباطاً عالي المعنوية  
وموجب بصفات معدل الحليج، طول التيلة عند 2.5%، نعومة التيلة، متانة التيلة في الجيل الثاني  
معدل الحليج وطول التيلة عند 2.5% في الجيلين الثالث والرابع، بينما تغير الارتباط لنعومة و متانة  
التيلة في الجيلين الثالث والرابع. كما كان هناك ارتباطاً معنوياً موجباً بين صفات طول التيلة عند  
2.5% و نعومة و متانة التيلة و معدل الانتظام للهجينين في الثلاثة أجيال بمتوسط وزن اللوزة. و بين  
معدل الحليج و نعومة التيلة للجيل الثالث في الهجين الأول و نعومة و متانة التيلة للثلاثة أجيال في  
الهجين الثاني. وأظهرت نتائج الدراسة امكانية استخدام هذين الهجينين في برامج تربية وتحسين  
صفات المحصول والجودة في القطن.

قام بتحكيم البحث

كلية الزراعة – جامعة المنصورة  
مركز البحوث الزراعية

أ.د / عبد الرحيم ليله  
أ.د / حسين يحيى عوض