

# Heritability and correlation analysis of morphological and yield traits in genetically modified cotton

**Abdul Rehman**

Cotton Research Institute

**Nida Mustafa**

University of Agriculture Faisalabad Faculty of Agriculture

**Du Xiongming**

Cotton Research Institute

**Muhammad Tehseen Azhar** (✉ [tehseenazhar@gmail.com](mailto:tehseenazhar@gmail.com))

Department of Plant Breeding and Genetics, University of Agriculture, Faisalabad <https://orcid.org/0000-0003-1137-481X>

---

## Research

**Keywords:** Genetic Correlation, Genetic variability, Inheritance, Productivity, Upland cotton

**Posted Date:** July 2nd, 2020

**DOI:** <https://doi.org/10.21203/rs.3.rs-17787/v2>

**License:**   This work is licensed under a Creative Commons Attribution 4.0 International License. [Read Full License](#)

---

**Version of Record:** A version of this preprint was published on August 26th, 2020. See the published version at <https://doi.org/10.1186/s42397-020-00067-z>.

# Abstract

Background Cotton is known for fiber extraction and it is grown in tropical and sub-tropical areas of the world due to having hot weather. Cotton crop has a significant role in GDP of Pakistan. Therefore, the two years research was conducted to estimate heritability and association among various yield contributing parameters of cotton i.e. plant height, number of bolls per plant, number of sympodial branches per plant, seed cotton yield, boll weight, seed index, GOT, fiber length, fiber strength, and fiber fineness. Results Association analysis revealed that seed cotton yield had significant positive correlation with plant height, number of bolls per plant, number of sympodial branches per plant, GOT, staple length and fiber strength. Staple length and fiber strength were negatively linked with each other. Estimates of heritability were high for all of the traits except number of sympodial branches per plant and boll weight. Conclusion The parent IUB-222 was found to be best for plant height, number of bolls per plant, boll weight, GOT, seed cotton yield and seed index. The genotypes namely NIAB-414 and VH-367 were identified as best parents for fiber length, strength and fineness. Among the crosses NIAB-414 × IUB-222 was best for number of bolls per plant, seed index, seed cotton yield and fiber fineness, whereas, cross of NIAB-414 × CIM-632 was good for plant height. The combination of A555 × CIM-632 was best for number of sympodial branches per plant, boll weight, fiber length and strength, and VH-367 × CIM-632 proved best for GOT.

# Introduction

Cotton is also known as white gold due to its white and soft fiber, also called vegetable fibre. The cotton plant was grown like shrub in nature and its fiber is pure cellulose. The cotton fiber is used to spin into yarn which is further used for making socks, curtains and towels etc. Its fiber also consumed in textile industry for cloth making (Stewart and Rossi 2010). Significant amount of oil (16-27%) is extracted from cotton seed and seed cake is used in livestock industry, and oil extracted from cotton seed is used as vegetable oil for making fries, excite etc. because taste of cottonseed oil is similar to coconut oil. In addition, it is an important source of vitamins, fat and antioxidants (Dowd et al., 2010). During 2018-2019 survey, cotton was cultivated on an area of 2,373 thousand hectares with 9.861 million bales production. It shares 0.8% in GDP and 4.5% in value addition (Govt. of Pakistan, 2018-19).

The world population is increasing day by day therefore it is necessary to increase the productivity of crop to meet the requirement of textile industry. Utilization of various breeding tools is one of method to meet the demand of textile industry (Farooq et al., 2014). Understanding the genetic basis of important yield contributing traits is the pre-requisite and information about their relationship must be available to cotton breeder. All of yield related traits are correlated with each other in such a way that increase or decrease in one trait directly effects on other traits. So, estimation of genotypic and phenotypic correlations among these traits are helpful to initiate the breeding programs. The knowledge about association among various plant characters is useful in the selection of appropriate breeding method (Teklewold et al., 2000). Phenotypic correlation shows the visual observation while genotypic correlation estimates the inheritance of characters (Desalegn et al., 2009). It is indicated that number of bolls and number of sympodial branches per plant were positively linked with each other. Weight of a boll had negative relationship with number of bolls per plant. Seed cotton yield and number of bolls were also positively correlated with each other. Heritability values were also high for these traits (Shar et al., 2017). Investigations revealed that association and inheritance for various quantitative and fiber related

parameters of American cotton (Haq et al., 2017). In addition, yield of seed cotton was positively linked with plant height, sympodial branches, monopodial branches and bolls per plant whereas negatively correlated with days to 1<sup>st</sup> flowering. While seed cotton yield had positive correlation with 100-seed weight, number of bolls per plant, plant height and boll weight (Memon et al., 2017; Mukoyi et al., 2018). Lint index, number of bolls per plant, boll weight, sympodial branches/plant and GOT exhibited positive linkage with yield of seed cotton per plant. Heritability was high for number of bolls/plant, monopodial branches/plant, internode distance and sympodial branches/plant (Monisha et al., 2018). High heritability and positive correlation were reported for monopodia/plant, number of bolls/plant, yield of seed cotton and fiber fineness (Khokhar et al., 2017; Komala et al., 2018). Positive correlation and high heritability were observed for plant height, sympodial branches, number of bolls, boll weight, seed cotton yield and fiber fineness. Hence, it is concluded that these traits may be considered as selection criteria for improvement in seed cotton yield (Jarwar et al., 2018; Rathinavel et al., 2017). The presented research was planned to determine correlation among various yield contributing traits due to increased demand of cotton in the country. In addition, heritability of these parameters was also computed which could be used for the selection of suitable traits from certain parents for the development of new germplasm of upland cotton.

## Materials And Methods

The experiment was performed at two places, first in glasshouse and then at cotton research area of the Department of Plant Breeding and Genetics, University of Agriculture, (latitude 31°25'N, longitude 73°09'E and altitude 184.4 m from sea level) Faisalabad, Pakistan. Experimental material was collected from Cotton Research Group of the department of Plant Breeding and Genetics, University of Agriculture, Faisalabad, Pakistan. Five genotypes namely, A-555, IUB-222, VH-367, NIAB- 414 and CIM-632 were grown in earthen pots during November 2017 in greenhouse conditions available with the department. The optimal growing conditions *i.e.* temperature (25-35°C) light intensity (25,000-30,000 lux) and humidity (44-49%) was maintained for germination and growth of the plants in glasshouse. At two leaf stage one healthy seedling per pot was kept while others were thinned. These five parents were crossed to make all possible combinations in full diallel at the appearance of buds. Some of the buds from parents were selfed. Later on, cotton seed from 20 crosses along with their parents was picked, ginned and sown at cotton farm during second week of May 2018. The parents along with F<sub>1</sub> population planted in three replications followed by randomized complete block design (RCBD). Row to row and plant to plant distance was 75 cm and 30 cm respectively. All agronomic practices were followed from sowing to harvesting to get good and healthy plant population. Following plant parameters of cotton plant were noted at various time intervals and protocol of each trait is mentioned in the following paragraphs.

**Plant height (cm):** Plant height was measured in cm with the help of measuring rod. The height was measured from the first cotyledonary node to the apical bud at maturity. Average height of seven guarded plants was calculated in each family.

**Number of bolls per plant:** Fully opened bolls were picked and recorded from all the replications of each family. Average number of bolls were calculated for each parent/cross for data analysis.

**Number of sympodial branches per plant:** Sympodial branches are direct fruit bearing branches, these are generally short appearing towards the top of the stem. At maturity, number of sympodial branches of seven guarded plants was counted manually in each replication and then average values were calculated for each parent/cross.

**Seed cotton yield (g):** Seed cotton was picked from mature opened bolls from seven guarded plants. Seed cotton picked during all of pickings was cumulatively weighed in grams by using electronic balance. Average seed cotton yield was calculated for each parent/cross for use in analysis.

**Boll weight (g):** Boll weight was obtained by dividing weight of seed cotton yield from each plant by the number of bolls per plant. Average boll weight was calculated for each genotype for biometrical analysis.

**Seed index (g):** Seed index determined from 100 seed weight from each plant. Cotton seeds were separated from each plant by using single roller ginning machine (McCarthy Roller Gin 1840). A random sample of 100 seeds was obtained from each plant and weighed by using electronic balance to determine the seed index. Mean seed index was calculated for each parent/cross in all of replications.

**Ginning out turn (%):** First seed cotton yield was weighed then ginned with a single roller electrical gin machine (McCarthy Roller Gin 1840). The lint obtained from each sample was weighed separately. Ginning percentage was calculated by using the formula as proposed by by Singh (2004).

$$\text{GOT (\%)} = \frac{\text{Weight of lint}}{\text{Weight of seed cotton}} \times 100$$

**Fiber length (mm), Fiber strength (g/tex) and Fiber fineness (µg/inch):** Fiber traits were measured by high using high volume instrument (Model USTER@ HVI-900 SA) system and means for each parent and cross were calculated.

### Statistical Approaches

The data collected were subjected to analysis of variance following the method of Steel et al. (1997) in order to determine the significant differences in plant characters of upland cotton by using (Minitab Inc., USA). Standard deviation and standard error were calculated by using following formulae.

$$SD = \sqrt{\frac{\Sigma(X - M)^2}{n - 1}}$$

$$SE = SD/\sqrt{n}$$

Genotypic and phenotypic correlation among traits were analyzed by a statistical technique that is known as correlation analysis (Kwon and Torrie, 1964). Whereas, Heritability in broad sense was estimated according to Burton and Devance (1953). Heritability were divided in three classes *i.e.* Low heritability < 0.2, Medium heritability = 0.2-0.5 and High heritability > 0.5.

$$h^2_{(BS)} = (\sigma^2_g / \sigma^2_p)$$

where,

$\sigma^2_g$  = The genotypic variance

$\sigma^2_p$  = The phenotypic variance

$h^2_{(BS)}$  = Heritability broad sense

## Results

The analysis of variance (ANOVA) exhibited significant differences and confirm the presence of variations among genotypes for the traits namely plant height, number of bolls per plant, number of sympodial branches per plant, boll weight, yield of seed cotton, seed index, fiber length, fiber strength and fiber fineness (Table 1). In later stage this data were used for correlation and heritability analysis. The significant results of these traits allowed the researcher to proceed for correlation and heritability analysis.

## Correlation coefficient analysis

Correlation coefficient analysis measures the relationship between various plant characters. The estimation of genotypic and phenotypic correlations among studied traits are helpful for initiating breeding programs. If correlation between two traits is positive and significant then improvement in one trait will exert significant impact on other trait. Hence, selection for one character will also improve the other associated traits. In present study, Plant height exhibited positive and significant linkage with number of bolls/plant, number of sympodial branches per plant, GOT, seed cotton yield, fiber length and fiber fineness whilst non-significant association with boll weight (Table 2). However, plant height showed negative and non-significant relationship with fiber strength. Number of bolls per plant had positive and significant association with plant height, number of sympodial branches per plant, boll weight, seed index, seed cotton yield and fiber strength whereas positive but non-significant relationship with GOT and fiber fineness but found negative and non-significant correlation with fiber length. Number of sympodial branches per plant had positive and significant relationship with all of traits except seed index and staple length. Boll weight had positive and significant correlation with number of bolls per plant, number of sympodial branches per plant, 100 seed weight, staple length and fiber fineness whereas rest had positive but non-significant association except GOT which have positive and significant linkage with plant height, seed cotton yield, number of sympodial branches per plant and fiber length. It had positive but non-significant association with number of bolls per plant. However, GOT had negative and non-significant correlation with boll weight, 100 seed weight, fiber strength and fiber fineness. Seed index had positive and significant linkage with number of bolls per plant, boll weight and fiber length while had positive but non-significant correlation with plant height, number of sympodial branches/plant, yield of seed cotton and fiber fineness. It had negative and non-significant relationship with GOT and fiber strength.

Seed cotton yield had positive and significant association with plant height, number of bolls per plant, number of sympodial branches per plant, GOT, staple length and fiber strength. Seed index showed positive but non-significant correlation with boll weight, seed index and fiber fineness whereas negative and non-significant association with fiber strength. Fiber length presented positive and significant linkage with plant height, boll weight, GOT, seed index, fiber fineness and seed cotton yield while positive but non-significant correlation with number of sympodial branches per plant. Staple length exhibited negative and significant association with fiber strength while negative and non-significant relationship with number of bolls/plant and fiber fineness. Fiber strength had positive and significant association with number of bolls per plant, number of sympodial branches per plant, and seed cotton yield whereas positive and non-significant correlation with boll weight. Whereas, revealed negative and significant linkage with fiber length while had negative and non-significant relationship with plant height, GOT and seed index. Fiber fineness had positive and significant correlation with plant height, number of sympodial branches per plant, boll weight, and staple length whereas positive and non-significant association with number of bolls per plant and seed index. It showed negative and non-significant linkage with GOT, yield of seed cotton and fiber strength.

## Estimates of heritability

Heritability (B.S) were ranged from medium to high for various yield and fiber related traits (Table 3). The traits namely plant height, number of bolls per plant, GOT, seed index, seed cotton yield, fiber length, strength and fineness showed high heritability estimates *i.e.* 74.48, 53.87, 90.65, 53.42, 54.56, 52.95, 59.66 and 70.42 respectively. While the traits including sympodial branches per plant and boll weight exhibited medium

estimates of heritability *i.e.* 48.06 and 46.66. The process of selection could be useful for characters with high heritability values.

### Mean comparison for certain traits

Among parents, the genotype A555 had lowest mean value 99.93 cm, while CIM-632 exposed maximum mean value 124.73 cm for plant height (Figure 1-A). Whereas, hybrid IUB-222 × NIAB-414 exhibited minimum mean value 88.07 cm, whereas NIAB-414 × CIM-632 showed highest mean value 127.2 cm. VH-367 exposed minimum number of bolls per plant with lowest mean value 16.26 whereas IUB-222 showed maximum value 32.13 followed by CIM-632, NIAB-414 and A555 having mean estimates of *i.e.* 26.53, 23.26, 23.13 correspondingly (Figure 1-B). Crosses, VH-367 × CIM-632 indicated highest mean value 32.93 followed by NIAB-414 × IUB222, NIAB-414 × CIM-632 and A555 × IUB-222, having mean values *i.e.* 32.53, 31, 30.4 and 28.2 respectively. VH-367 showed highest mean value 23.13 followed by A555 (22.66), NIAB-414 (20.93) and IUB-222 (20.93) for number of sympodial branches per plant (Figure 1-C). A555 × IUB-222 revealed minimum mean value 18.86, whereas VH-367 × NIAB-414 indicated maximum number of sympodial branches 25.73. Genotype, CIM-632 had minimum mean value for boll weight 1.83 g whereas VH-367 exposed maximum mean value 2.46 g (Figure 1-D). Hybrids, IUB-222 × VH-367 revealed lowest but VH-367 × NIAB-414 showed maximum boll weight. NIAB-414 had minimum mean value for GOT 38.03 % while IUB-222 had maximum GOT 42.32 % (Figure 1-E). NIAB × IUB-222 showed minimum value 35.46 % whilst hybrid VH-367 × CIM-632 revealed maximum GOT 46.33 % followed by CIM-632 × VH-367, CIM-632 × A555 and A555 × NIAB-414 with estimates of 45.62, 44.33 and 43.22%.

The accessions NIAB-414 and A555 exhibited the minimum and maximum mean estimates 5.07 and 5.64g respectively for 100 seed weight (Figure 1-F). Hybrid CIM-632 × VH-367 presented minimum seed index 4.62 g whereas CIM-632 × A555 exhibited maximum mean value 5.70g. While studying the seed cotton yield it was observed that IUB-222 revealed maximum mean value 74.81g (Figure 1-G) whilst IUB-222 × A555 showed minimum seed cotton yield 42.39 g but the hybrid VH-367 × CIM-632 revealed highest mean value 80.17g as compare to A555 × IUB-222 and NIAB-414 × IUB-222 with mean values of 73.95 and 72.84 respectively. CIM-632 indicated lowest value 24.71mm, while NIAB-414 had maximum fiber length 26.86mm (Figure 1-H). Among the hybrids, A555 × VH-376 had minimum fiber length 24.29 mm, while highest mean value 28.33 mm was shown by CIM-632 × NIAB-414. In case of fiber strength, VH-367 indicated maximum mean value 29.83 g/tex followed by A555, NIAB-414 and CIM-632 having mean values of 26.35 , 26.28 and 25.47 g/tex respectively (Figure 1-I). Furthermore, CIM-632 × IUB-222 had minimum fiber strength 23.29 g/tex, while VH-367 × NIAB-414 revealed highest value 27.96 g/tex. IUB-222 exhibited lowest value 3.31µg/inch for fiber fineness, while NIAB-414 had highest mean value 5.06 µg/inch (Figure 1-J). While among the hybrids IUB-222 × NIAB-414 exposed minimum mean value 3.23 µg/inch, but NIAB-414 × IUB-222 had highest mean value 5.44 µg/inch.

## Discussion

Plant height was positively linked with sympodial branches per plant, number of bolls per plant, GOT, seed cotton yield, staple length and fiber fineness (Table 2). Azhar and Ajmal (1999), Rao and Gopinath (2013) and Shahzad et al. (2015) also had similar findings. Tulasi et al. (2012) also observed positive association with GOT, fiber length and fineness. Heritability (B.S) for plant height was 74.48% (Table 3). Kapoor and Kaushik (2003), Ahmad et al. (2011) and Baloch et al. (2015) also found high heritability 94%, 81% and 96.4% correspondingly for plant height. High heritability estimates indicated that selection for plant height can be

effective. Bolls per plant had positive association with plant height, boll weight, sympodial branches per plant, seed index, seed cotton yield and fiber strength. Ahmad and Azhar (2000), Djaboutou et al. (2005), Gul et al. (2014), Magadum et al. (2012), Alkuddsi et al. (2013) and Farooq et al. (2014), also found same results. Heritability value for bolls per plant was 53.87% (Table 3). Desalegn et al. (2009), Ahmad et al. (2011), Baloch et al. (2015) and Rathinavel et al. (2017) estimated 59%, 88%, 93% and 60.21% high broad sense heritability respectively for bolls per plant. High estimates of heritability revealed that successful and effective selection can be helpful in the improvement of this trait.

Sympodial branches per plant had positive relationship with plant height, number of bolls per plant, boll weight, seed cotton yield, GOT, fiber strength and fiber fineness (Table 2). Pujer et al. (2014), Joshi et al. (2006), Anandan (2009) indicated that sympodial branches/plant positively correlated with seed cotton yield, plant height, GOT and boll weight. Whereas, Killi et al. (2005) found that sympodial branches per plant were positively linked with fiber strength. Rauf et al. (2004) also observed that sympodial branches per plant had positive relationship with number of bolls per plant and fiber fineness. Moderate heritability for this trait was observed *i.e.* 48.06% (Table 3). Ahmed et al. (2006), Mustafa et al. (2007), Neelima and Reddy (2008) and Kulkarni et al. (2011) also observed medium heritability 50.72%, 59%, 61.30% and 43% respectively for sympodial branches per plant. Boll weight was positively linked with bolls per plant, sympodial branches per plant, 100 seed weight, staple length and fiber fineness (Table 2). Jatt et al. (2007) revealed that boll weight had positive association with yield of seed cotton. Abdullah et al. (2016) and Shaheen and Yaseen (2014) observed that boll weight was positively correlated with fiber length, fiber fineness and sympodial branches per plant. Do Thi et al. (2008) and Kale et al. (2007) reported that boll weight positively linked with seed index and number of bolls per plant. Whilst heritability value was moderate 46.66% for this trait (Table 3). Huangjun and Myers (2011), Naveed et al. (2004) and Ahmed et al. (2006) estimated 57%, 22% and 50.0% medium heritability respectively for boll weight.

GOT had positive relationship with plant height, seed cotton yield, sympodial branches per plant and fiber length (Table 2). Monicashree and Balu (2018), Pujer et al., (2014) and Chattha et al. (2013) observed that GOT had positive linkage with plant height and sympodial branches per plant and yield of seed cotton. Shahzad et al. (2015) observed that GOT had positive association with staple length. Heritability for GOT was 90.65% (Table 3). Devidas et al. (2017), Shahzad et al. (2015), Kumar and Katageri (2017) and Jarwar et al. (2018) found high heritability values 72.5%, 80.73%, 90.0% and 85.46% for GOT. Seed index had positive linkage with bolls per plant, boll weight and fiber length (Table 2). Patil (2010), Komala et al. (2018), Memon et al. (2017), Isong et al. (2017), Ashokkumar and Ravikesavan (2010), Shabbir et al. (2016) and Méndez et al. (2012) depicted similar findings. Heritability (B.S) for this trait was 53.42% (Table 3). Dhivya et al. (2014), Kaleri et al. (2016), Kumar and Katageri (2017) and Rajamani et al. (2015) estimated 60.01%, 72.24%, 51.63% and 66.72% heritability correspondingly for seed index. Significant progress is possible through selection for this character.

Seed cotton yield had positive association with plant height, number of bolls per plant, sympodial branches per plant, GOT, fiber length and fiber strength (Table 2). Majeedano et al. (2014), Joshi et al. (2006), Gite et al. (2006) and Latif et al. (2015) indicated that seed cotton yield was positively linked with plant height, sympodial branches per plant and number of bolls/plant.. Monisha et al. (2018) determined positive correlation among GOT, fiber strength and seed cotton yield. Heritability value for seed cotton yield was 54.56% (Table 3). Desalegn et al. (2009), Reddy and Reddy (2007), Hussain et al. (2010), Ullah et al. (2015) and Ahmad et al. (2011) estimated 61%, 80%, 50%, 98% and 76% heritability respectively for this trait. This trait could be improved

through selection process. Fiber length was positively linked with plant height, boll weight, GOT, seed index, fiber fineness and seed cotton yield. Fiber length had negative correlation with fiber strength (Table 2). Ali and Awan (2009) and Echekwu (2001) indicated that fiber length was negatively associated with fiber strength. Bechere et al. (2014) indicated that fiber length had positive linkage with GOT. Killi et al. (2005) determined positive association among fiber length, plant height and seed cotton yield. Abbas et al. (2013) observed that staple length was positively associated with fiber fineness. Shabbir et al. (2016) observed that fiber length had positive association with seed index. Khan and Azhar (2000) found that fiber length had positive relationship with boll weight. Heritability in broad sense for fiber length was 52.95% (Table 3). Killi et al. (2005), Abbas et al. (2013), Khan and Azhar (2000) and Ahmed et al. (2006) found 94%, 52%, 96% and 56% heritability estimates respectively for fiber length. It is concluded from results that selection can be useful for fiber length. Fiber strength had positive association with bolls per plant, sympodial branches per plant, and seed cotton yield whereas had negative linkage with fiber length (Table 2). Ahmad and Azhar (2000), Thiyagu et al. (2010) and Farooq et al. (2014) found that fiber strength was positively correlated with yield of seed cotton and number of bolls per plant. Ali and Awan (2009) revealed that fiber strength was negatively linked with fiber length. For this trait heritability value was 59.66% (Table 3). Desalegn et al. (2009), Killi et al. (2005), Shahzad et al. (2015), Rasheed et al. (2009) and Khokhar et al. (2017) determined 33%, 73%, 62%, 70% and 68% heritability for this character. Fiber fineness was positively correlated with plant height, sympodial branches per plant, boll weight, and staple length (Table 2). Ali and Awan (2009), Zeng and Meredith (2009), Tang and Xiao (2014) and Yaqoob et al. (2016) found positive linkage between fiber fineness and staple length. Abbas et al. (2013) and Altaher and Singh (2003) revealed that fiber fineness had positive linkage with plant height, sympodial branches per plant. Abdullah et al. (2016) reported that fiber fineness was positively correlated with boll weight. Heritability value for fiber fineness was 70.42% (Table 3). Hendawi et al. (1999) and Lu et al. (2002) estimated 67% and 73% heritability respectively for fiber fineness.

## Conclusion

The correlation results from this study would be helpful to breed cotton cultivars with good yield and quality characters. Broad sense heritability was high for all of parameters which provides the strong evidence that selection in early generations can improve the performance of these traits.

## Declarations

### Affiliations of authors

<sup>1</sup>Zhengzhou Research Base, State Key Laboratory of Cotton Biology, Zhengzhou University, <sup>2</sup>Zhengzhou 450000, China. <sup>3</sup>Institute of Cotton Research, Chinese Academy of Agricultural Sciences, Anyang 455000, China.

<sup>3</sup>Department of Plant breeding and genetics, University of Agriculture Faisalabad, Pakistan.

### Ethics approval and consent to participate

Not applicable.

## Consent for publication

Not applicable.

## Availability of data and material

Not applicable.

## Conflict of interest

Authors declare that they have no conflict of interest for the publication of the manuscript.

## Funding

This work was supported by the department of Plant Breeding and Genetics, Faculty of Agriculture, University of Agriculture, Faisalabad, Pakistan.

## Authors' contributions

NM and AR conducted experiment and wrote the initial draft of the manuscript. MTA played role in designing and statistics of the experiment, in addition MTA supervised NM for her master studies. DX proofread the manuscript before submission to JCR.

## References

- Abbas H, Mahmood A, Ali Q, Khan M, Nazeer W, Aslam T, Zahid W: Genetic variability, heritability, genetic advance and correlation studies in cotton (*Gossypium hirsutum* L.). *Int Res J Microbiol.* 2013, 4(6):156-161.
- Abdullah M, Numan M, Shafique MS, Shakoor A, Rehman S, Ahmad M: Genetic variability and interrelationship of various agronomic traits using correlation and path analysis in cotton (*Gossypium hirsutum* L.). *Acad J Agri Res.* 2016, 4(6):315-318.
- Ahmad M, Azhar FM: Genetic correlation and path coefficient analysis of oil and protein contents and other quantitative characters in F2 generation of *Gossypium hirsutum* L. *Pak J Biol Sci.* 2000, 3(6):1049-1051.
- Ahmad M, Khan NU, Muhammad F, Khan SA, Munir I, Bibi Z, Shaheen S: Genetic potential and heritability studies for some polygenic traits in cotton (*Gossypium hirsutum* L.). *Pak J Bot* 2011, 43(3):1713-1718.
- Ahmed HM, Kandhro MM, Laghari S, Abro S: Heritability and genetic advance as selection indicators for improvement in cotton (*Gossypium hirsutum* L.). *J Biol Sci* 2006, 6(1):96-99.
- Ali MA, Awan SI: Inheritance pattern of seed and lint traits in cotton (*Gossypium hirsutum*). *Int J Agric Biol* 2009, 11(1):44-48.
- Alkuddsi Y, Rao MG, Patil S, Joshi M, Gowda T: Correlation and path coefficient analysis between seed cotton yield and its attributing characters in intra hirsutum cotton hybrids. *Mol Plant Breed.* 2013, 4.

- Altaher A, Singh R: Yield component analysis in upland cotton (*Gossypium hirsutum* L.). J Indian Soc Cotton Improv 2003, 28(3):151-157.
- Anandan A: Studies on choice of characters for breeding for seed cotton yield and fibre quality traits in cotton. Crop Improv. 2009, 36(1):35-37.
- Ashokkumar K, Ravikesavan R: Genetic studies of correlation and path coefficient analysis for seed oil, yield and fibre quality traits in cotton (*G. hirsutum* L.). Aust J Basic App Sci. 2010, 4(11):5496-5499.
- Azhar F, Ajmal S: Diallel analysis of oil content in seed of *Gossypium hirsutum* L.[Pakistan]. J Genet Breed. (Italy) 1999.
- Baloch M, Baloch A, Baloch M, Mallano I, Baloch A, Baloch N, Abro S: Association and heritability analysis for yield and fiber traits in promising genotypes of cotton (*Gossypium hirsutum* L.). Sindh Uni Res J. (Science Series) 2015, 47(2).
- Bechere E, Zeng L, Boykin D: Correlation and path-coefficient analyses of lint yield and other traits in upland cotton (*Gossypium hirsutum* L.). Crop Improv. 2014, 28(6):852-870.
- Burton GW, Devane dE: Estimating heritability in tall fescue (*Festuca arundinacea*) from replicated clonal material 1. Agron J. 1953, 45(10):478-481.
- Chattha WS, Farooq J, Ahmad A, Kang SA, Naveed-UI-Haq M: Correlation analysis of quality and yield contributing traits in upland cotton (*Gossypium hirsutum* L.). Int J Modern Agri 2013, 2:95-101.
- Desalegn Z, Ratanadilok N, Kaveeta R: Correlation and heritability for yield and fiber quality parameters of Ethiopian cotton (*Gossypium hirsutum* L.) estimated from 15 (diallel) crosses. Kasetart J(Nat Sci) 2009, 43:1-11.
- Devidas AA, Narayan SA, Prakash PN: Study of genetic variability, heritability and genetic advance in some genotypes of egyptian cotton (*Gossypium barbadense* L.). J Global Biosci. 2017, 6(4):4954-4957.
- Dhivya R, Amalabalu P, Pushpa R, Kavithamani D: Variability, heritability and genetic advance in upland cotton (*Gossypium hirsutum* L.). Afr J Plant Sci. 2014, 8(1):1-5.
- Djaboutou C, Alabi S, Echewku C, Orakwue F: Variability and interrelationship of some agronomic and fibre quality traits in multi-adversity cotton (*Gossypium hirsutum* L.). Agric Trop Subtrop. 2005, 38(3-4):7-12.
- Do Thi HA, Ravikesavan R, Iyanar K: Genetic advance and heritability as a selection index for improvement of yield and quality in cotton. J Cotton Res Develop. 2008, 22(1):14-18.
- Dowd MK, Boykin DL, Meredith Jr WR, Campbell BT, Bourland FM, Gannaway JR, Glass KM, Zhang J: Fatty acid profiles of cottonseed genotypes from the national cotton variety trials. J Cotton Sci. 2010, 14: 64-73.
- Echekwu C: Correlations and correlated responses in upland cotton (*Gossypium hirsutum* L.). Tropicultura 2001, 19(4):210-212.

- Farooq J, Anwar M, Riaz M, Farooq A, Mahmood A, Shahid M, Rafiq M, Ilahi F: Correlation and path coefficient analysis of earliness, fiber quality and yield contributing traits in cotton (*Gossypium hirsutum* L.). J Anim Plant Sci. 2014, 24(3).
- Gite V, Misal M, Kalpande H: Correlation and path analysis in cotton (*Gossypium hirsutum* L.). J Cotton Res Develop. 2006, 20(1):51-54.
- Government of Pakistan. 2018-19. Pakistan Economic Survey. Ministry of Finance, Economic Advisor's Wing, Islamabad.
- Gul S, Khan N, Batool S, Baloch M, Munir M, Sajid M, Khakwani A, Ghaloo S, Soomro Z, Kazmi S: Genotype by environment interaction and association of morpho-yield variables in upland cotton. J Anim Plant Sci 2014, 24(1):262-271.
- Haq A, Khan N, Raza H, Gul S, Akbar S, Khan S, Muhammad S, Ali M, Khan H, Khan S: Genetic attributes of F3 populations and their parental lines in upland cotton. J Anim Plant Sci. 2017, 27(2).
- Hendawi F, Radi M, Abdel-Hamid A, Ismail R: Inheritance of fiber traits in some cotton crosses. Egypt J Agron. 1999, 21: 15-36.
- Huangjun L, Myers GO: Combining abilities and inheritance of yield components in influential upland cotton varieties. Aust J Crop Sci. 2011, 5(4):384.
- Hussain S, Nawab N, Ali MA, Hussain A, Nawaz MA, Malik T: Evolution of performance, genetic divergence and character association of some polygenic traits in upland cotton. J Agric Soc Sci 2010, 6(4):79-82.
- Isong A, Balu PA, Ramakrishnan P: Association and principal component analysis of yield and its components in cultivated cotton. Electr. J Plant Breed. 2017, 8(3):857-864.
- Jarwar AH, Wang X, Wang L, Jarwar ZH, Ma Q, Fan S: Genetic Advancement, Variability and Heritability in Upland Cotton (*Gossypium hirsutum* L.). J Envir Agric Sci. 2018: 6: 24-31.
- Jatt T, Abro H, Larik A, Soomro Z: Performance of different cotton varieties under the climatic conditions of Jamshoro. Pak J Bot 2007, 39(7):2427-2430.
- Joshi H, Chovatia P, Mehta D: Genetic variability, character association and component analysis in upland cotton. Indian J Agric Res. 2006, 40(4):302-305.
- Kale U, Kalpande H, Annapurve S, Gite V: Yield components analysis in American cotton (*Gossypium hirsutum* L.). Madras Agric J 2007, 94(7-12):156-161.
- Kaleri AA, Baloch AW, Baloch M, Wahocho NA, Abro TF, Jogi Q, Soomro AA, Marri A, Bhutto LA: Heritability and correlation analysis in Bt and non-Bt cotton (*Gossypium hirsutum* L.) genotypes. Pure and App Bio. 2016, 5(4):1.
- Kapoor C, Kaushik S: Variability, heritability and genetic advance studies in cotton (*Gossypium hirsutum* L.). J Cotton Res Dev 2003, 17:242.

- Khan AI, Azhar FM: Estimates of heritabilities and pattern of association among different characters of *Gossypium hirsutum* L. Pak J Agri sa Vol 2000, 37:1-2.
- Khokhar ES, Shakeel A, Maqbool MA, Anwar MW, Tanveer Z, Irfan MF: Genetic Study of Cotton (*Gossypium hirsutum* L.) Genotypes for Different Agronomic, Yield and Quality Traits. Pak J Agric Res. 2017, 30(4).
- Killi F, Efe L, Mustafayev S: Genetic and environmental variability in yield, yield components and lint quality traits of cotton. Int J Agri Biol 2005, 7(6):1007-1010.
- Komala M, Ganesan NM, Kumar M: Genetic variability, Heritability and Correlation Analysis in F2 Populations of Ratoon Upland Cotton Hybrids. Int J Agric Envi Biotech. 2018, 11(6):815-827.
- Kulkarni A, Nanda H, Patil S: Study of genetic parameters on yield, yield contributing and fibre quality characters in upland cotton (*Gossypium hirsutum* L.). J Cotton Res Dev. 2011, 25(1):22-24.
- Kumar NM, Katageri I: Genetic variability and heritability study in F2 population of *Gossypium barbadense* L. cotton for yield and its components. Int J Curr Microbiol App Sci. 2017, 6(6):975-983.
- Kwon S, Torrie J: Heritability and interrelationship among traits of two soybean populations. Crop sci. 1964, 4(2):196-198.
- Latif A, Bilal M, Hussain SB, Ahmad F: Estimation of genetic divergence, association, direct and indirect effects of yield with other attributes in cotton (*Gossypium hirsutum* L.) using biplot correlation and path coefficient analysis. Trop. Plant Res. 2015, 2(2):120-126.
- Lu Y, Zhen Z, Zhen G, Ju P: Genetic stability of fibre quality in upland cotton. Cotton Sci. 2002, 14:67-70.
- Magadum S, Banerjee U, Ravikesavan R, Thiyagu K, Boopathi NM, Rajarathinam S: Association analysis of yield and fibre quality characters in interspecific population of cotton (*Gossypium* spp.). J Crop Sci Biotech. 2012, 15(3):239-243.
- Majeedano MS, Ahsaan M, Somroo A, Channa A: Heritability and correlation estimates for some yield traits of *Gossypium hirsutum*. Am Res Thoughts 2014, 1:781-790.
- Memon S, Gandahi AWBN, Yasir TA, Sarki SM, Wasaya A, Mallano IA, Ali M, Baloch AM, Khetran AS: Evaluation of genetic divergence, character associations and path analysis in upland cotton genotypes. Pure App Bio. 2017, 6(4):1516-1521.
- Méndez-Natera JR, Rondón A, Hernández J, Merazo-Pinto JF: Genetic studies in upland cotton. III. Genetic parameters, correlation and path analysis. Sabrao J Breed Genet. 2012, 44(1):112-128.
- Minitab IN. Minitab 16 statistical software. URL:[Computer software]. State College, PA: Minitab, Inc.(www.minitab.com). 2010.
- Monicashree C, Balu PA: Association and Path Analysis Studies of Yield and Fibre Quality Traits in Intraspecific Hybrids of Upland Cotton (*Gossypium hirsutum* L.). Research Journal of Agricultural Sciences 2018, 9(5):1101-1106.

- Monisha K, Premalatha N, Sakthivel N, Kumar M: Genetic variability and correlation studies in upland cotton (*Gossypium hirsutum* L). Elect J Plant Breed. 2018, 9(3):1053-1059.
- Mukoyi F, Gasura E, Makunde G: Implications of correlations and genotype by environment interactions among cotton traits. African Crop Science Journal 2018, 26(2):219-235.
- Mustafa A, Elsheikh Y, Babiker E: Genetic variability and character association and selection criteria in Cotton (*Gossypium hirsutum* L.). Sudan J Agric Res. 2007. 8: 43-50.
- Naveed M, Azhar F, Ali A: Estimates of heritabilities and correlations among seed cotton yield and its components in *Gossypium hirsutum* L. Int J Agri Biol 2004, 6(4):712-714.
- Neelima S, Reddy VC: Genetic parameters of yield and fibre quality traits in American cotton (*Gossypium hirsutum* L.). Indian J Agric Res. 2008, 42(1):67-70.
- Patil H: Variability and correlation analysis by using various quantitative traits in released Bt cotton hybrids. J Cotton Res Dev. 2010, 24(2):141-144.
- Pujer SK, Siwach S, Sangwan R, Sangwan O, Deshmukh J: Correlation and path coefficient analysis for yield and fibre quality traits in upland cotton (*Gossypium hirsutum* L). J Cotton Res Dev. 2014, 28:214-216.
- Rajamani S, Sumalatha P, Gopinath M: Studies on genetic parameters of seed cotton yield and fibre traits in upland cotton (*Gossypium hirsutum* L.). J Cotton Res Dev. 2015, 29(1):36-38.
- Rao P, Gopinath M: Association analysis of yield and fibre quality characters in upland cotton (*Gossypium hirsutum* L.). Aust J Basic App Sci. 2013, 7(8):787-790.
- Rasheed A, Malik W, Khan A, Murtaza N, Qayyum A, Noor E: Genetic evaluation of fiber yield and yield components in fifteen cotton (*Gossypium hirsutum*) genotypes. Int J Agric Biol 2009, 11:581-585.
- Rathinavel K, Kavitha H, Priyadharshini C: Assessment of genetic variability and correlation analysis of seed and seed cotton yield attributing traits of tetraploid cotton genotypes (*G. hirsutum* L.). Electr J Plant Breed. 2017, 8(4):1275-1283.
- Rauf S, Khan TM, Sadaqat HA, Khan AI: Correlation and path coefficient analysis of yield components in cotton (*Gossypium hirsutum* L.). Int J Agric Biol 2004, 6(4):686-688.
- Reddy YR, Reddy C: Genetic variability for yield components and fibre characters in cotton. Plant Arch. 2007, 7(2):759-761.
- Shabbir RH, Bashir QA, Shakeel A, Khan MM, Farooq J, Fiaz S, Noor M: Genetic divergence assessment in upland cotton (*Gossypium hirsutum* L.) using various statistical tools. J Global Inno Agric Soc Sci. 2016, 4(2):62-69.
- Shaheen M, Yaseen M: Path analysis based on genetic association of yield components in upland cotton. Life Sci Int J. 2014 Vol: 8 : 2988-2994

Shahzad MT, Ijaz F, Khan O, Saleem B, Hassan U: Correlation, Path Analysis & Heritability Among Some Yield and Fibre Related Traits of *Gossypium hirsutum* L. Cotton Genom Genet. 2015, 6.

Shar T, Baloch M, Arain M, Jatoi W, Lochi R: phenotypic associations, regression coefficients and heritability estimates for quantitative and fiber quality traits in upland cotton genotypes. Pak J Agric Eng VetSci. 2017, 33(2):142-152.

Singh, P. (2004). Cotton breeding. Kalyani Publishers Ludhiana New Delhi Noida (U.P) Hyderabad Chennai Kolkata Cuttack India. pp:295.

Steel, R.G.D., J.H. Torrie and D.A. Dickey (1997). Principles and procedures of statistics: A biometrical approach. (3<sup>rd</sup> ed.) McGraw Hill, New York.

Stewart L, Rossi J: Using cotton byproducts in beef cattle diets. *Bulletin* 1311. 2010.

Tang F, Xiao W: Genetic association of within-boll yield components and boll morphological traits with fibre properties in upland cotton (*Gossypium hirsutum* L.). Plant Breed. 2014, 133(4):521-529.

Teklewold A, Jayaramaiah H, Jagadeesh BN. Correlations and path analysis of physio-morphological characters of sunflower (*Helianthus annuus* L.) as related to breeding method. Helia. 2000;23(32):105-14.

Thiyagu K, Nadarajan N, Rajarathinam S, Sudhakar D, Rajendran K: Association and path analysis for seed cotton yield improvement in interspecific crosses of cotton (*Gossypium* spp). Electr J Plant Breed. 2010, 1(4):1001-1005.

Tulasi J, Lal MA, Murthy J, Rani YA: Correlation and path analysis in american cotton. Electr J Plant Breed. 2012, 3(4):1005-1008.

Ullah K, Usman Z, Khan N, Ullah R, Saleem FY, Khattak SI, Ali M: Genetic diversity for yield and related traits in upland cotton genotypes. Pak J Agric Res. 2015, 28(2): 118-125.

Yaqoob M, Fiaz S, Ijaz B: Correlation analysis for yield and fiber quality traits in upland cotton. Commun Plant Sc. 2016, 6(3/4):55-60.

Zeng L, Meredith WR: Associations among lint yield, yield components, and fiber properties in an introgressed population of cotton. Crop Sci. 2009, 49(5):1647-1654.

## Tables

**Table 1. Analysis of variance of various quantitative traits for F<sub>1</sub> population and parents in upland cotton formulated by using MINITAB 16.**

SOV	DF	PH	NB	SB	BW	GOT	SI	SCY	FL	FS	FF
Rep.	2	2.21 <sup>ns</sup>	2.88 <sup>ns</sup>	4.39 <sup>*</sup>	4.77 <sup>*</sup>	5.88 <sup>ns</sup>	0.29 <sup>ns</sup>	4.63 <sup>ns</sup>	4.65 <sup>*</sup>	5.68 <sup>*</sup>	0.35 <sup>ns</sup>
Gen.	24	3.92 <sup>**</sup>	2.17 <sup>*</sup>	1.93 <sup>*</sup>	1.86 <sup>*</sup>	10.70 <sup>**</sup>	2.15 <sup>*</sup>	16.21 <sup>**</sup>	2.13 <sup>*</sup>	2.48 <sup>**</sup>	3.37 <sup>**</sup>
Err.	48	-	-	-	-	-	-	-	-	-	-

\*\* = Highly significant  $p \leq 0.01$ , \* = Significant  $p \leq 0.05$

Where, PH (plant height), BP (bolls per plant), SB (number of sympodial branches per plant), BW (boll weight), GOT (ginning out turn), SI (seed index), SCY (seed cotton yield), FL (fiber length), FS (fiber strength) and FF (fiber fineness).

**Table 2. Correlation among various traits of Upland cotton grown in filed conditions by using correlation analysis**

Traits	PH	BP	SB	BW	GOT	SI	SCY	FL	FS
<b>BP</b>	0.16*								
<b>SB</b>	0.63**	0.17*							
<b>BW</b>	0.05	0.21**	0.07*						
<b>GOT</b>	0.34**	0.02	0.22*	-0.04					
<b>SI</b>	0.08	0.25*	0.09	0.51**	-0.02				
<b>SCY</b>	0.29**	0.54**	0.39**	0.18	0.28**	0.02			
<b>FL</b>	0.20*	-0.04	0.06	0.20*	0.19*	0.17*	0.03*		
<b>FS</b>	-0.16	0.19*	0.27*	0.12	-0.06	-0.05	0.33**	-0.02*	
<b>FF</b>	0.08*	0.04	0.17*	0.27*	-0.14	0.26	-0.06	0.11*	-0.05

Where, PH (plant height), BP (bolls per plant), SB (number of sympodial branches per plant), BW (boll weight), GOT (ginning out turn), SI (seed index), SCY (seed cotton yield), FL (fiber length), FS (fiber strength) and FF (fiber fineness).

**Table 3: Range of heritability of various traits of cotton grown in field conditions according ti the formula fiven by Burton and Devance (1953)**

<b>Traits</b>	<b>Heritability (%)</b>	<b>Status</b>
Plant height	74.48	High
Sympodial branches/plant	48.06	Medium
Number of bolls/plant	53.87	High
Boll weight	46.66	Medium
Ginning out turn	90.65	High
Seed index	53.42	High
Seed cotton yield	54.56	High
Fiber length	52.95	High
Fiber strength	59.66	High
Fiber fineness	70.42	High

High > 0.5, Medium 0.2-0.5, Low < 0.2

## Figures



**Figure 1**

Mean comparison for plant height of parents and crosses of upland cotton where, A for PH (plant height), B-BP (bolls per plant), C- SB (number of sympodial branches per plant), D- BW (boll weight), E- GOT (ginning out turn), F- SI (seed index), G- SCY (seed cotton yield), H- FL (fiber length), I- FS (fiber strength) and J- FF (fiber fineness).