

Correlation Analysis of Stem Hardness Traits with Fiber and Yield Related Traits in Core Collections of *Gossypium hirsutum*

Irum Raza

Chinese Academy of Agricultural Sciences Cotton Research Institute

Dao-Wu Hu

Chinese Academy of Agricultural Sciences Cotton Research Institute

Adeel Ahmad

Chinese Academy of Agricultural Sciences Cotton Research Institute

Hongge Li

Chinese Academy of Agricultural Sciences Cotton Research Institute

Shou-Pu He

Chinese Academy of Agricultural Sciences Cotton Research Institute

Mian Faisal Nazir

Chinese Academy of Agricultural Sciences Cotton Research Institute

Xiao-Yang Wang

Chinese Academy of Agricultural Sciences Cotton Research Institute

Yin-Hua Jia

Chinese Academy of Agricultural Sciences Cotton Research Institute

Zhao-e Pan

Chinese Academy of Agricultural Sciences Cotton Research Institute

Peng Zhang

Chinese Academy of Agricultural Sciences Cotton Research Institute

Muhammad Yasir

Chinese Academy of Agricultural Sciences Cotton Research Institute

Muhammad Shahid Iqbal

Chinese Academy of Agricultural Sciences Cotton Research Institute

Xiao-Li Geng

Chinese Academy of Agricultural Sciences Cotton Research Institute

Li-Ru Wang

Chinese Academy of Agricultural Sciences Cotton Research Institute

Bao-Yin Pang

Chinese Academy of Agricultural Sciences Cotton Research Institute

Xiong-Ming Du (✉ dxm630723@163.com)

Research

Keywords: Bending, Compression, Acupuncture, Principle component analysis, Stem hardness

Posted Date: June 18th, 2020

DOI: <https://doi.org/10.21203/rs.3.rs-36287/v1>

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 March 28th, 2021. See the published version at <https://doi.org/10.1186/s42397-021-00082-8>.

Abstract

Background Stem hardness is one of the major influencing factors for plant architecture in upland cotton (*Gossypium hirsutum* L.). Evaluating hardness phenotypic traits is very important for the selection elite lines for resistant to lodging in *Gossypium hirsutum* L. Cotton breeder are interested in using diverse genotypes to enhance fibre quality and high- yield. The research for hardness and its relation with fiber quality and yield were very few. This study was designed to find the relationship of stem hardness traits with fiber quality and yield contributing traits of upland cotton.

Results Experiments were carried out to measure the bending, acupuncture and compression properties of stem from a collection of upland cotton genotypes, comprising 237 accessions. The results showed that the genotypic difference in stem hardness were highly significant among the genotypes, and the stem hardness traits (BL, BU, AL, AU, CL and CU) have a positive association with fiber quality traits and yield related traits. In descriptive statistics result bending (BL, BU) have maximum coefficient of variance and trait fiber length and fiber strength have less coefficient of variance among the genotypes. Principal component analysis (PCA) reduced quantitative characters into nine principal components. The first nine principal components (PC) with Eigen values >1 explained 0.86% of variation among 237 accessions of cotton crop. Both 2017& 2018, PCA results indicated that BL, BU, FL, FE and LI variables contributed their variability in PC1 and BU, AU, CU, FD, LP and FWPB have shown their variability in PC2.

Conclusion We describe here, to the best of our knowledge, the systematic study of the mechanism involved in the regulation of enhancing fiber quality and yield by stem bending strength, acupuncture and compression properties of *Gossypium hirsutum* crop.

1. Introduction

Cotton is one of the most important cash crops and the only major fiber crop in the world. The contribution of cotton to the total fiber used worldwide is about 35 percent (Zhang et al. 2014). Upland cotton (*Gossypium hirsutum* L.) is the largest cultivated species of cotton, occupying more than 90 percent of the world cotton cultivated area, which reflects widespread adaptability and high yield production characteristics (Wendel, 1989; Chen et al. 2007). *Gossypium hirsutum* L. is allotetraploid ($2n = 4x = 52$), and is composed of two ancestral genomes that are designated as At from *Gossypium arboreum* and Dt from *Gossypium raimondii* (Al-Ghazi et al. 2009). Due to long-term natural selection and artificial breeding, a number of cotton germplasm resources for sustainable genetic improvement have been created under varied climatic and cultivating conditions. In the National Gene Bank for Cotton, China, 7,712 *G. hirsutum* accessions are present. All these accessions were collected from cotton producing countries around the world since 1865 when the United States introduced upland cotton (Dai et al. 2016). In order to efficiently use these resources, various efforts have been made to investigate and evaluate cotton diversity (Fang et al. 2017b; Huang et al. 2017; Sun et al. 2017; Wang et al. 2017; Ma et al. 2018).

Yield and quality of the produce are the most important factors for all crops (Fang et al. 2017a). Stem hardness is basic characteristic in the plant architecture of cotton, which is not well studied. Stem hardness may have a relationship with yield and fiber quality. The physical characteristics contributing to the strength of stem are the bending force, puncture force and compression force. The stem bending force is the force at which the trunk bends or breaks under a particular load. The basal portion of the culm internode plays a crucial role in ensuring that the plant remains upright (Peng et al. 2014). The greater carbohydrate accumulation in the base stem could increase the force need to bend the stem (Ishimaru et al. 2008). Stem thickness is a biological indicator for green or dry biomass. The strength (force, stress) and energy requirements are the compressive properties. Therefore, the selection of genotypes with increased stem strength is a useful field indicator (Beeck et al. 2006). Compression properties of stem depend on species, variety, stalk structure, stalk diameter, maturity, moisture and cell structure (Persson 1987). A physical quantitative measurement may enhance selection effectiveness and boost genetic gain, such as penetrometer sorghum measurement (Pedersen & toy 1999). Therefore, greater understanding of these parameters provide a theoretical basis to enhance the physical strength of the stem and basal part of the culm internode, with the aim of obtaining a higher yield and good fiber quality of cotton.

Amorphous fibrils, lignin, and pectin, present in the cell wall, are also known to enhance the strength and hardness of the stem (Mohsenin 1986). Lignin or cellulose generally determines physical strength, as a low content of lignin or cellulose causes a brittle culm (Tanaka et al. 2003). In wheat plant the mechanical strength of the stem provided by cellulose and lignin to the lodging resistance plants (Cai et al. 2019). The selection of elevated stalk strength and resistance to the corn border increases the elements of cell walls in the breeding programme (Li et al. 2016) . Cotton has a high biomass output and a high cellulose and lignin proportion. In mature cotton fibers the secondary cell wall (SCW) includes over 90% cellulose and it differs from all other known species of plant. By contrast, typical SCWs contain 40-50% cellulose in dicotyledonous stem xylem (Huang et al. 2016). The knowledge of stem hardness therefore allows the cell wall to be modified to improve fiber quality and quantity, because plant cell wall has a close association with mechanical and biochemical strength of stem parameters.

Principal component analysis (PCA) has been used extensively in the plant sciences for variable reduction and genotype grouping. This is the most prevalent statistical multivariate method used in environmental studies (Tahri et al. 2005; Yongming et al. 2006). PCA commonly used in the analysis of the relationships between observed variables and in the extraction of a small number of autonomous factors (major component) (Tokalioglu S et al. 2006). It commences with the correlation matrix, it describes the dispersion of the original variables and extracts eigenvalues and eigenvectors (Astel et al. 2008). Eigenvector is a list of coefficients that multiply the original correlated variants to obtain new uncorrelated (orthogonal) principal components that are linearly weighted combinations of the original variables. The number of correlated variables can be reduced to a smaller set of orthogonal factors, which allows the interpretation of a specified multidimensional system by showing correlations between the original variables. The analysis of the correlations also reflects a related response of a given character and also provides a good index for predicting the corresponding change in one character to the

extent of the proportional change in the other. PCA was used by Kamara et al. (2003) to identify maize (*Zea mays* L.) traits which accounted for the majority of variance in the data. Granati et al. (2003) have used PCA to investigate the relationship among *Lathyrus* accessions. Žáková and Benková (2006) identified the traits of 106 Slovak barley accessions as the primary sources of variations in genetic diversity. PCA and cluster analysis were used by Cartea et al (2002), Salihu et al. (2006), respectively to group kale populations and genotypes of winter wheat. The current study examined the multivariate data analysis of agronomic and quality characteristics of a global collection of 237 genotypes.

Some studies have been conducted on stem strength behaviors of different plants; however, for stem hardness characteristics of cotton stalk, no data is reported. The present research therefore seeks to establish a relationship between stem hardness and yield characteristics and quality characteristics in *Gossypium hirsutum*. However, yield is a complicated, multicomponent controlled character. Stem hardness components are less sensitive than yield per se to the environmental changes and are therefore comparatively more likely to improve. Once the nature and extent of relations among these component characteristics and yield are understood, effectiveness of choice in the segregated generation will improve. Therefore, the present research was carried out to assess PCA and correlations of significant *Gossypium hirsutum* characteristics.

2. Materials & Methods

2.1. Cotton Accessions

From a set of 7,362 *G. hirsutum* accessions, preserved at the China National Gene Bank, Cotton Research Institute, Chinese Academy of Agriculture Sciences, Anyang, Henan, 237 cotton genotypes were selected. These accessions have various geographical origins including China, the United States, the former Soviet Union, Australia, Brazil, Pakistan, Mexico, Chad, Uganda and Sudan, which are the world's largest cotton-growing areas.

2.2. Planting and phenotyping

Phenotyping of stem hardness-related features was performed during the normal cotton growing season (mid-April to late-October) at Cotton Research Institute, Anyang, Henan, China (Yellow River command area) for two years i.e., 2017 and 2018. Coordinates of the location are E 114.07° and N 35.85 °, longitude and latitude respectively. All accessions (237) were planted in randomized complete block design with three replicates in the experimental field. Each entry plot had a dimension of 7m × 3m and row-to-row and plant-to-plant distance was 30cm and 76cm respectively. Field management practices were conducted according to the local management scheme. The scoring standards for phenotypic traits in both years were identical. Six stem hardness traits and 14 agronomic traits were characterized.

2.3. Sample preparation for stem hardness traits

The stems were cut and separated from the branches after harvesting the cotton genotypes. Stem samples were air dried for two months in the lab. At the time of hardness testing, the air-dried cotton stem had low humidity content. The stem was equally divided into two parts for the preparation of test samples: Upper and lower ((Additional file 2: Figure S1a).

2.4. Stem Hardness Traits

For each replicate, three plant were selected to test the hardness of the stem. These characteristics were Breaking force of the Upper part (BU), breaking point of lower part (BL), Compression force of upper and lower part (CU and CL), and acupuncture force of the upper and lower part (AU and AL). The YYD-1 SS testing system (TOP Instrument Co., Zhejiang, China) was used to measure all hardness characteristics of 15 cm segment from lower, and upper part of stem (Additional file 2: Figure S2b). The tester was set perpendicular to the culm at the middle, under gradual loading, and the breaking force was measured when the culm was pushed to breaking point. The maximum force in Mega Newtons needed in order to break, puncture & compress the center of the two segments of the stem (upper and lower) was recorded.

2.5. Agronomic traits

Days to first flower opening, FD (days) were calculated from the date of sowing to the day when first flowers bloomed on 50 percent of the plants in each plot. Plant height (PH) recorded from the base of plant above ground to the tip of the plant. Ten consecutive plants were selected for plant height in each plot. From each accession, 30 naturally opened bolls were harvested randomly to calculate boll weight (BW) in grams and to gin the fiber. The Seed Index (SI) was calculated after counting and weighing 100 cotton seeds. Fiber samples were separately weighed to calculate the Lint percentage (LP) and Fiber weight per boll (FWPB) in grams. The Lint index (LI) was calculated based on SI and LP data.

$$\text{Lint Index} = \frac{\text{Seed Index} \times \text{Lint Percentage}}{1 - \text{Lint Percentage}}$$

Fiber samples were examined in the Cotton Quality Test Center in Anhui, China for fiber-quality characteristics using a high-volume instrument (HFT9000). Data on the fiber length (FL, mm), fiber strength (cN/tex), micronaire value (Mic, $\mu\text{g/inch}$), elongation percentage (EP, %), Length uniformity (LU, %), Spinning Consistency Index (SCI) recorded. Average of the three replicates in the same year is defined to be phenotypic information per accession.

2.6. Statistical analysis

For the evaluation of phenotypic traits statistics, Minitab 18 and R were used. The primary impacts of the experimental variables and their relationships were analyzed by the analysis of variances (ANOVA). The significance level for ANOVA was set at $p \leq 0.05$. R software (package "corrplot") was used for calculating and plotting correlation. Principal component analysis was performed using Minitab 18.

3. Results

3.1. Stem hardness variations among the genotypes

The ANOVA given in shows Tab. 1. that genotypic difference in stem hardness were highly significant for traits like bending (BL and BU), and compression CU ($p > 0.05$). Basic descriptive statistics (mean, standard deviation, minimum, maximum and coefficient of variance) of all the genotypes for morphological, yield and fiber traits were studied (Additional file 1: Table S1) It was observed that maximum coefficient of variance (26.67% & 20.40) was recorded in bending (BU & BL) respectively, which mean and SD was (0.08 & 0.19) and (0.02 & 0.04) followed by acupuncture upper and lower (17.81 & 15.76) compression upper and lower (14.96 & 13.45), seed index (14) and fibre length (14) with a mean and SD of (0.07, 0.08, 0.54, 0.82, 12.30 & 30.39) and (0.01, 0.01, 0.08, 0.11, 1.43 & 1.82) respectively. The traits like fibre elongation (1.47) and fibre strength (3.43) have comparably less coefficient of variance among the genotypes. Similar result was found in 2017 data.

Table .1 ANOVA for 20 traits in 2017 and 2018.

Trait	Sum of square		Mean square		F Value		P value		R ² (adj)
	G	Y	G	Y	G	Y	G	Y	
BL	1.05	0.3	0	0.3	1.49	102.8	**	***	31.54
AL	0.05	0.09	0	0.09	0.88	341.1	ns	***	39.72
CL	2.78	3.39	0.011	3.39	1.19	343.16	ns	***	45.04
BU	0.19	0.03	0	0.03	1.6	60.21	***	***	29.72
AU	0.02	0	0	0	1.13	14.94	ns	***	8.54
CU	3.26	0.08	0.01	0.08	1.24	7.33	*	**	11.71
PH	307	184.8	130.32	184.84	2.92	4.14	***	*	49.09
BW	128.80	294.80	0.55	294.75	1.02	551	ns	***	53.98
GP	489	564	207.22	5646.6	3.3	89.85	***	***	57.16
LP	8163.1	110.1	34.58	110.05	8.54	27.17	***	***	79.24
LI	485.55	208.75	2.057	208.75	7.57	768.48	***	***	83.06
FWPB	43.97	49.23	0.18	49.23	2.09	552.3	***	***	63.09
SI	801.82	328.83	3.39	328.83	9.4	910.14	***	***	85.95
FS	1221.7	68.37	5.177	68.37	0.86	11.36	ns	**	0
FL	1339.1	671.9	5.674	671.89	7.14	845.55	***	***	82.9
UP	691.97	11.96	2.932	11.959	2.57	10.47	***	**	44.49
SCI	105	0	447.86	0.103	3477.25	0.8	***	ns	99.94
FD	101	197	43.1	19754.4	2.04	935.48	***	***	71.38
FE	31.5	0	0.13	0	3.33	0	***	ns	53.76
MI	86.54	0.04	0.36	0.04	4.84	0.59	***	ns	65.67

G, genotype; **Y**, Year; **ns**, non-significant; * and ***, significant at P < 0.05 and P < 0.001, respectively.

BL: bending lower; **AL**: acupuncture lower; **CL**: compression lower; **BU**: bending upper; **AU**: acupuncture upper; **CU**: compression upper;

PH: plant height; **BW**: boll weight; **GP**: growth period; **LP**: lint percentage; **LI**: lint index; **FWPB**: flowering weight per boll;

SI: seed index; **FS**: fibre strength; **FL**: fibre length; **UP**: uniformity percentage; **SCI**: spinning consistancy index;

The results showed (Additional file 1: Table S2) the variation among different varieties. Based on the bending trait values all 237 accessions of cotton were differentiated in to two groups, higher stem hardness (HSH); those varieties that have the higher value of bending trait and lower stem hardness (LSH); the varieties, which have lower values of bending traits. In Table S2. Only mentioned six HSH genotypes and six LSH genotypes.

3.2. Principal component analysis

Principal component analysis was performed to Only the Principal component (PCs) with an eigenvalues higher than 1 maintained according to the Kaiser (1960) criterion. Thus PC1, PC2, PC3, PC4, PC5 and PC6 (Tab. 2) were selected as these represented 0.23%, 0.14%, 0.12%, 0.08%, 0.08% and 0.06% of progeny variation, respectively, and accounted for 0.73% of the overall diversity. PC7, PC8 and PC9 variances represented a cumulative percentage of 0.78%, 0.82% and 0.86% respectively. Table 2 summarizes the PCs and the eigenvectors, which were estimated on the average of twenty variables. All twenty traits contributed to the total variation in PC1, but Fibre length (FL), uniformity percentage (UP), fibre elongation (FE), Lint index (LI), Bending lower (BL) and bending upper (BU) have contributed more in PC1. PC1 is a weighted average of these characters indicating that fibre quality traits have significant importance for this component. On the other hand, other traits are less important to PC1. While In PC2, all variables are contributed but the main contributors to variation were bending upper (BU), acupuncture upper (AU), compression upper (CU), days to flowering (FD), lint percentage (LP) and fiber weight per boll (FEPB) So yield traits have more contribution in PC2. For PC3, bending lower (BL), compression lower (CL), compression Upper (CU) bending upper (BU), Maturity (M) and Spinning consistency index (SCI) are the most important trait, while multiple traits contributed to the other PCs in varying proportions and the same trend as found in 2017 result.

Table. 2 Eigenvectors and Eigen analysis of the Correlation Matrix for the nine principal components traits associated with Stem hardness Performance in *Gossypium hirsutum* accessions

Traits	Years	PC1	PC2	PC3	PC4	PC5	PC6	PC7	PC8	PC9
BL	2017	0.18	0.42	0.02	-0.04	-0.12	-0.13	0.07	-0.01	0.12
	2018	0.26	0.09	0.38	0.05	0	-0.15	-0.08	0.13	0.01
AL	2017	0.04	0.16	0.17	-0.23	0.47	-0.4	0.04	-0.1	0.48
	2018	0.09	0.1	0.19	-0.1	-0.57	0	-0.04	-0.09	-0.11
CL	2017	0.11	0.28	-0.05	0	-0.4	0	0.13	0.16	0.37
	2018	0.138	0.01	0.29	0	0.5	-0.05	0.01	-0.28	0.03
BU	2017	0.19	0.4	0.05	0	-0.19	-0.13	-0.02	-0.13	-0.21
	2018	0.22	0.18	0.43	0.05	-0.05	-0.11	0.02	0.04	0.02
AU	2017	0.1	0.26	0.14	-0.26	0.43	-0.22	-0.02	0.06	-0.27
	2018	0.06	0.13	0.15	-0.13	-0.53	0.02	-0.06	-0.4	0.15
CU	2017	0.15	0.4	0.04	-0.01	-0.22	-0.01	0.02	0.07	-0.42
	2018	0.17	0.17	0.39	0.01	0.28	-0.08	0	-0.18	-0.06
FS	2017	-0.07	0.02	0.05	-0.05	-0.11	-0.16	-0.86	0.42	0.07
	2018	0.32	0.09	-0.29	-0.09	0.02	-0.12	-0.05	-0.05	0.21
FL	2017	0.29	-0.08	0.34	0.19	0	-0.05	0.02	0.09	0.02
	2018	0.37	0.06	-0.19	-0.06	0.03	-0.04	0.12	-0.08	-0.01
M	2017	0.14	-0.02	-0.51	-0.25	-0.02	-0.07	0.11	0.36	0.13
	2018	0	-0.31	0.19	-0.02	0.03	0.24	-0.54	0.11	0.5
UP	2017	0.31	-0.08	0.18	0.19	0.12	0.03	0.06	0.16	-0.02
	2018	0.34	-0.1	-0.1	-0.12	0	0.04	-0.17	0.06	0.17
FE	2017	0.24	-0.07	-0.01	0.15	0.05	0.07	0.27	0.54	0.19
	2018	0.35	-0.07	-0.16	-0.12	0.01	0.06	-0.03	0.01	0.31
SCI	2017	0.19	-0.07	0.47	0.3	-0.05	0.09	-0.07	-0.02	0.12
	2018	0.35	0.13	-0.27	-0.12	0.01	-0.17	0.19	-0.02	-0.03
FD	2017	0.23	0.17	-0.23	0.06	0.34	0.41	-0.18	-0.07	0
	2018	0.08	0.31	0.01	0.37	0	0.433	0.19	0.15	0.2
PH	2017	0.21	0.08	-0.16	0.12	-0.15	0.02	-0.23	-0.48	0.4

	2018	0.16	-0.17	0.18	-0.14	-0.11	-0.16	0.14	0.75	-0.11
BW	2017	0.29	-0.21	0.04	-0.39	-0.17	0.07	-0.02	-0.14	0
	2018	-0.04	-0.08	-0.07	0.54	-0.1	-0.52	-0.03	0	0.22
GP	2017	0.26	0.14	-0.11	-0.02	0.27	0.5	-0.16	0.01	-0.02
	2018	0.09	0.36	-0.06	0.37	-0.05	0.36	0.04	0.16	0.05
LP	2017	0.27	-0.19	-0.25	0.24	0.06	-0.36	-0.05	-0.06	-0.18
	2018	0.22	-0.42	0.06	0.08	-0.05	0.3	0.17	-0.13	-0.23
LI	2017	0.32	-0.26	-0.03	-0.14	-0.04	-0.18	-0.05	-0.03	-0.15
	2018	0.29	-0.23	-0.03	0.2	-0.02	0.23	-0.27	-0.07	-0.5
FWPB	2017	0.34	-0.24	-0.12	-0.1	-0.09	-0.15	-0.03	-0.15	-0.06
	2018	0.13	-0.37	-0.01	0.48	-0.11	-0.18	0.1	-0.12	0.04
SI	2017	0.06	-0.11	0.33	-0.59	-0.15	0.27	0.01	0.02	0.02
	2018	0.06	0.32	-0.16	0.13	0.03	-0.16	-0.65	0.1	-0.32
Eigenvalue	2017	5.42	3.07	1.78	1.55	1.36	1.23	0.99	0.87	0.75
	2018	4.61	2.9	2.55	1.69	1.66	1.33	0.98	0.79	0.71
Proportion	2017	0.27	0.15	0.08	0.07	0.06	0.06	0.05	0.04	0.03
	2018	0.23	0.14	0.12	0.08	0.08	0.06	0.04	0.04	0.03
cumulative	2017	0.27	0.42	0.51	0.59	0.66	0.722	0.772	0.81	0.855
	2018	0.23	0.37	0.504	0.58	0.67	0.73	0.78	0.82	0.86
BL: bending lower; AL: acupuncture lower; CL: compression lower; BU: bending upper; AU: acupuncture upper; CU: compression upper;										
PH: plant height; BW: boll weight; GP: growth period; LP: lint percentage; LI: lint index; FWPB: flowering weight per boll;										
SI: seed index; FS: fibre strength; FL: fibre length; UP: uniformity percentage; SCI: spinning consistancy index;										
FD: days to flowering; FE: fibre elongation; MI: Micornire index										

3.3. Stem hardness correlation with fiber quality traits

The result of the 2018 correlation of stem hardness indicated that bending lower (BL) has a positive association with fiber length, micronaire value, uniformity percentage, fibre elongation, spinning

consistency index and days to flowering (Fig. 1). Bending upper (BU) was a positively correlation with FL, UP, SCI, and FD. Compression lower (CL) were a positive correlation with FL, M, SCI and FD, while compression upper (CU) was a positively correlation with LU, SCI and FD. Same result were found in 2017 correlation of stem hardness with fibre related traits (Additional file 2: Figure S2).

3.4. Stem hardness correlation with yield related traits and morphological traits

In Fig. 2, the 2018 correlation result showed that Bending lower (BL) was highly positively associated with PH, GP, LP and FWPB, while bending upper (BU) was in positive association with PH, GP and LI. Acupuncture lower (AL) showed positive correlation with PH. Compression (CL and CU) have a positive correlation with PH, GP, BW and LI. We also found the same correlation trend of all traits in 2017 data (Additional file 2: Figure S3).

4. Discussion

In the last decade, there has been great progress in developing new cotton genotypes for better fiber quality and higher yield. The stem associated characteristics such as bending, acupuncture and compression may be used to determine yield and quality of the fiber. One reason for influencing crop quality and yield is plant height (Tang JH 2007). The fiber quality parameters on which textile processing and the quality of the item rely, are fiber strength, length therefore premium pricing are charged for these quality features (Hussain K 2010).

Our breeding program goal for *G. hirsutum* was to identify high-yield genotypes, some agronomic features that are easily evaluated and linked with these characteristics could be used as markers (Biyun Chen et al. 2014). In this study, we observed that the bending, compression and acupuncture related to stem hardness have a positive and substantial correlation with the fiber length, spinning consistency and flowering times. May (2002) speculated that enhanced fiber strength could demand more energy; therefore, higher strength genotypes produce fewer than lower strength lines. Pettigrew (2001 & 2008) reported that an increase in light and temperature also increased the strength, the difference was however not enough to cause a yield penalty. Our findings showed a positive association of length uniformity, micronaire values with stem hardness characteristics. Fiber fineness was positively associated with fiber length and fiber strength by Killi et al. (2005). There was a negative association of fiber fineness with a fiber strength and fiber uniformity ratio. The fiber strength showed positive correlation with fiber uniformity. Mature cotton fibers are approximately 95% cellulose with other polysaccharides such as arabinose, galactose and xylose (Meinert and Delmer 1977) and pectin (Meinert and Delmer, 1977; Wang et al. 2010). These are important for determining fiber strength by joining cellulose fibrils. A direct correlation between cellulose molecular weight and fiber strength was reported by Timpa and Ramey (1994). Though the metabolic cost of these polysaccharides is higher (Amthor 2010), a higher metabolic cost, unless transport of complex polysaccharides was an issue, seems unlikely to be a yield drain for such small fractions of the fiber. These fundamental reasons for negative association yield and fiber strength should be investigated further. Fiber diameter reduced from bottom to top, possibly because cell

wall thickness decreased. Similarly, this can be explained by the fact that development in cell walls depends upon the accumulation of metabolism products (cellulose, hemicellulose, lignin, waxes etc.) that rises with maturity (C. Ververis et al. 2003). The major requirement for increasing rice grain yield is to enhance the physical strength of the culm in order to enhance the breaking-type lodging resistance (Hirano et al. 2014). It has therefore been concluded that stem-related characteristics like bending stress have determined the morphology and the quality of the culm, such as cellulose, lignin, pectin inside the cell wall, which have a direct relationship to high yield and crop quality.

The Lint Index is the main feature and contributes significantly to the lint percentage increase. The promising cultivars showed a maximum lint percentage due to the close association of the lint index (LI) with the lint percentage. Our current findings have demonstrated an important correlation of the LI & growth period (GP) with the bending (BL and BU) and Compression (CL and CU). Bending lower (BL) were positively associated for lint yield features such as LI and GOT%. The GOT% has a positive and substantial correlation with the lint index, according to (Hussain K 2010). Positive observations were found by Scholl and Miller (1976), while Tyagi (1994) stated that the GOT% was negatively associated with the lint index. Positive associations between lint index and lint percentage suggest an increased cottonseed yield. This result indicates that the lint index and the lint percentage are significant elements for enhancing cotton yield and should be considered during breeding program.

Plant height (PH) is one of the major morphological features, which plays a key part and closely associated with plant bolls (if there is no lodging) with the ultimate positive impact on cotton yield. Because of the lodging risk and appropriate for mechanical picking, cotton breeders are mostly interested in short-stature plants, but the plant height is strongly associated with bolls per plant and seed yields (Khan 2003). The present correlation results showed that Bending, Acupuncture & Compression have a positive correlation with PH. The stability and adaptability of *G. hirsutum* cultivars were studied by Meena et al. (2007) and reported varied values for yield components and plant height. The varieties of upland cotton were also evaluated by Suinaga et al. (2006) and it was found that the plant height was associated positively with the seed cotton yield and bolls per plant. The positive correlation between plant height and cotton yield seed was observed by Khan (2003), Soomro et al. (2005) and Taohúa and Haiping (2006), and their research showed that plant height contributed 70% of the total variability in seed cotton yield. Therefore, it is concluded that in cotton crop, height of plant is desirable if no lodging occurred.

Obtaining high seed yield per unit is one of the most important challenges in *G. hirsutum* breeding. Several agronomic traits are important for improving yield traits. Boll weight is the second major yield component and have a greater contribution in enhancement of seed cotton yield. The similar proportion and variation for boll weight with regard to the cotton seed yield of was observed by Khan (2003) and Copur (2006). For yields and other economic characters, Taohua and Haipeng (2006) and Meena et al. (2007) assessed various *G. hirsutum* varieties, significant variations were observed for boll weight and the effect on cotton seed yield was positive. The correlation result showed that Bending lower (BL) and compression upper (CU) have a positive correlation with BW, GP and FEPB. Batool et al. (2010) and

Makhdoom et al. (2010) also stated that boll weight had been found to be positive for yield and had a higher contribution to the yield improvement for upland cotton. Results also revealed that the boll weight following the bolls per plant had positive effect on seed cotton yield. Therefore, it is concluded that boll weight is an important yield component and should be kept in mind while breeding for seed cotton yield.

5. Conclusions

It may be concluded from the present study that cotton fiber quality and yield can be improved by selecting types having high strength of stem. Stem hardness related traits bending, acupuncture and compression show positive association with fiber and yield related traits. Also enhancing the stem strength has proven to be an effective approach to decrease stem lodging risk. Because stem lodging is a persistent problem to decrease yield. Thus during future breeding programs these parameters also kept in mind during selection, as they were the major attributes of the cotton quality and yield. Recurrent selection could be followed to accumulate genes for the said traits in any population. In addition, the phenotypic data for stem hardness may be used in our subsequent genome-wide association studies for *G. hirsutum*.

Declarations

Acknowledgments

Not applicable

Authors' contributions

I. R, X.D designed the study. I.R, D.Wu H, H. L, S.P.H, M. F. N, M.Y, S.I, X.Y.W, Y.H.J, Z.P, Z.P, X.L.G, W.L.R and B.Y.P performed the experiment and collected data. I.R and A.A analyzed the data. I.R wrote the manuscript. X.D, A.A, D.Wu H review the manuscript. Supervision: X.D. All authors read and approved the final version of manuscript.

Funding

This work was supported by funding from the National Key Technology R&D Program, the Ministry of Science and Technology (2016YFD0100306-2016YFD0100203), the National Natural Science Foundation of China (grants 31671746).

Availability of data and materials

No other data related to this study is available at this time.

Ethics approval and consent to participate

Not applicable.

Consent for publication

Not applicable

Competing interests

The authors declare that they have no competing interests

References

1. Al-Ghazi, Bourot Y, Arioli S, Dennis T, Llewellyn ES. Transcript Profiling During Fiber Development Identifies Pathways in Secondary Metabolism and Cell Wall Structure That May Contribute to Cotton Fiber Quality. *Plant Cell Physiol.* 2009; 50: 1364-1381.
2. Amthor J.S . From sunlight to phytomass: on the potential efficiency of converting solar radiation to phyto-energy. *New Phytol.* 2010; 188: 939-959.
3. Astel A, Astel K, Biziuk M. PCA and multidimensional visualization techniques united to aid in the bioindication of elements from transplanted Sphagnum palustre moss exposed in the Gdansk City area. *Environ Sci Pollut Res Int.* 2008; 15 :41-50.
4. Beeck CP, Wroth J, Cowling WA, Genetic variation in stem strength in field pea (*Pisum sativum L.*) and its association with compressed stem thickness. *Aust J Agr Res.* 2006; 57: 193-199.
5. Biyun C, Kun X, Jun L, et al. Evaluation of yield and agronomic traits and their genetic variation in 488 global collections of *Brassica napus L.* *Genet Resour Crop Evol.* 2014; 61: 979–999.
6. Batool S, Khan NU, Makhdoom K, et al. Heritability and genetic potential of upland cotton genotypes for morpho-yield traits. *Pak. J. Bot.* 2010; 42(2): 1057-1064.
7. Cai T, Peng D, Wang R, et al. Can intercropping or mixed cropping of two genotypes enhance wheat-lodging resistance? *Field Crop Res.* 2019; 239: 10-18.
8. Cartea M.E, Picoagea A, Soengas P, Ordás A. Morphological characterization of kale populations from northwestern Spain. *Euphytica*, 2002; 129: 25-32.
9. Chen ZJ, Scheffler BE, Dennis E, et al. Toward sequencing cotton (*Gossypium*) genomes. *Plant Physiol.* 2007; 145:1303–10. <https://doi.org/10.1104/pp.107.107672>.
10. Ververis C, Georghiou K, Christodoulakis N, et al. Fiber dimensions, lignin and cellulose content of various plant materials and their suitability for paper production, *Industrial Crops and Products.* 2003; 19: 245–254.
11. Copur O. Determination of yield and yield components of some cotton cultivars in semi-arid conditions. *Pak. J. Biol. Sci.* 2006; 9(14): 2572-2578.
12. Dai PH, Sun JL, Jia YH, et al. Construction of core collection of upland cotton based on phenotypic data. *J Plant Genet Resour.* 2016; 17.
13. Fang C, Ma Y, Wu S, Liu Z, et al. Genome-wide association studies dissect the genetic networks underlying agronomical traits in soybean. *Genome Biol.* 2017; 18: 161.

14. Fang L, Wang Q, Hu Y, Jia Y, et al. Genomic analyses in cotton identify signatures of selection and loci associated with fiber quality and yield traits. *Nat Genet.* 2017; 49: 1089-1098.
15. Granati E, Bisignano V, Chiaretti D, Crino P, Polignano BG, et al. Characterization of Italian and exotic *Lathyrus* germplasm for quality traits. *Genet. Res. Crop Evol.* 2013; 50: 273-280.
16. Hirano K, Okuno A, Hobo T, et al. Utilization of stiff culm trait of rice smos1 mutant for increased lodging resistance. *PLoS One.* 2014; 9: e96009.
17. Huang C, Nie XH, Shen C, et al. Population structure and genetic basis of the agronomic traits of upland cotton in China revealed by a genome-wide association study using high-density SNPs. *Plant Biotechnology Journal.* 2017; 15: 1374-1386.
18. Huang JF, Chen F, Wu SY, Li J, Xu WL. Cotton GhMYB7 is predominantly expressed in developing fibers and regulates secondary cell wall biosynthesis in transgenic Arabidopsis. *Sci China Life Sci.* 2016; 59: 194-205.
19. Hussain K, Khan IA, Sadaqat HA, Amjad M, Genotypic and phenotypic correlation analysis of yield and fiber quality determining traits in upland cotton (*Gossypium hirsutum*). *Int. J. Agric. Biol.* 2010; 12: 348–352.
20. Ishimaru K, Togawa E, Ookawa T, et al. New target for rice lodging resistance and its effect in a typhoon. *Plant.* 2008; 227: 601-609.
21. Kamara AY, Kling JG, Menkir A, Ibikunle O. Agronomic performance of maize (*Zea maysL.*) breeding lines derived from low nitrogen maize population. *J. Agric. Sci.* 2003; 141: 221-230.
22. Killi F, Efe L, Mustafayev S. Genetic and environmental variability in yield, yield components and lint quality traits of cotton. *Int. J. Agric. Biol.* 2005; 7: 1007-1010.
23. Khan NU. Genetic analysis, combining ability and heterotic studies for yield, its components, fiber and oil quality traits in upland cotton (*G. hirsutum L.*). Ph.D. Dissert. Sindh Agric. Univ. Tandojam, Pakistan; 2003.
24. Li K, Wang H, Hu X. et al. Genome-Wide Association Study Reveals the Genetic Basis of Stalk Cell Wall Components in Maize. *PLoS One.* 2016; 11: e0158906.
25. Ma Z, He S, Wang X. et al. Resequencing a core collection of upland cotton identifies genomic variation and loci influencing fiber quality and yield. *Nat Genet.* 2018; 50, 803-813.
26. Meinert MC, Delmer DP. Changes in biochemical composition of the cell wall of the cotton fiber during development. *Plant Physiol.* 1977; 59: 1088-1097.
27. Meena R.A, Monga D, and Kumar R. Undescriptive cotton cultivars of north zone: an evaluation. *J. Cotton Res.* 2007; 21(1): 21-23.
28. Makhdoom K, Khan NU, Batool S. et al. Genetic aptitude and correlation studies in *G. hirsutum L.* *Pak. J. Bot.* 2010 ; 42(3): 2011-2017.
29. Mohsenin NN. Physical Properties of Plant and Animal Materials, Seconded. Gordon and Breach, Science Publishers Inc., New York. 1986; 58-76.

30. Peng DL, Chen XG, Yin YP. et al. Lodging resistance of winter wheat (*Triticum aestivum* L.): Lignin accumulation and its related enzymes activities due to the application of paclobutrazol or gibberellin acid. *Field Crop Res.* 2014;157: 1-7.
31. Persson S. Mechanics of cutting plant material. American Society of Agricultural Engineers, St. Joseph, Mich., USA. 1987;
32. Peddersen JF, Toy JJ. Measurement of sorghum stalk strength using the missourie – modified electronic rent penetrometer, Mydica. 1999; 44: 155-158.
33. Pettigrew WT. Environmental effects on cotton fiber carbohydrate concentration and quality. *Crop Sci.* 2001; 41: 1108–1113.
34. Pettigrew WT. The effect of higher temperatures on cotton lint yield production and fiber quality. *Crop Sci.* 2008; 48: 278–285.
35. Salihu S, Grausgruber H, Ruckenbauer P. Agronomic and quality performance of international winter wheat genotypes grown in Kosovo. *Cereal Res. Commun.* 2006; 34: 957-964.
36. Sun Z, Wang X, Liu Z. et al. Genome-wide association study discovered genetic variation and candidate genes of fibre quality traits in *Gossypium hirsutum* L. *Plant Biotechnol J.* 2017; 15: 982-996.
37. Scholl RL, Miller PA. Genetic associations between yield and fiber strength in upland cotton. *Crop Sci.* 1976; 16: 780–783.
38. Suinaga F A, Bastos CS, Rangel LEP. Phenotypic adaptability and stability of cotton cultivars in Mato Grosso State, Brazil. *Pesquisa Agropecuaria Trop.* 2006; 36(3): 145-150.
39. Soomro AR, Kakar RG, Ali H, and S.A. Abid. Comparison of yield and its components in some commercial cotton varieties. *Indus J. Plant Sci.* 2005; 4(4): 545-552
40. Tahri M, Benyaich F, Bounakhla M. et al. Multivariate analysis of heavy metal contents in soils, sediments and water in the region of Meknes (central Morocco). *Environ Monit Assess.* 2005; 102: 405-417.
41. Tanaka K, Murata K, Yamazaki M. et al. Three distinct rice cellulose synthase catalytic subunit genes required for cellulose synthesis in the secondary wall. *Plant Physiol.* 2003; 133: 73-83.
42. Tokalioglu S, Kartal S. Multivariate analysis of the data and speciation of heavy metals in street dust samples from the organized industrial district in Kayseri (Turkey), *Atmos. Environ.* 2006; 40 : 2797–2805.
43. Tang JH, Teng WT, Yan JB. et al. Genetic dissection of plant height by molecular markers using a population of recombinant inbred lines in maize. *Euphytica.* 2007; 155: 117–124.
doi: [10.1007/s10681-006-9312-3](https://doi.org/10.1007/s10681-006-9312-3)
44. Timpa JD, Ramey HH. Relationship between cotton fiber strength and cellulose molecular-weight distribution - HVI calibration standards. *Text. Res. J.* 1994; 64: 557–562.
45. Tyagi AP. Correlation coefficient and selection indices in upland cotton (*Gossypium hirsutum* L.). *Indian J. Agric. Res.* 1994; 28: 189–196.

46. Taohua Z. and Z. Haipeng. Comparative study on yield and main agri-characters of five hybrids coloured cotton varieties. *J. Anhui Agric. Univ.* 2006; 33(4): 533-536
47. Wang H, Guo Y, Lv F. et al. The essential role of GhPEL gene, encoding a pectate lyase, in cell wall loosening by depolymerization of the de-esterified pectin during fiber elongation in cotton. *Plant Mol Biol.* 2010; 72: 397-406.
48. Wang MJ, Tu LL, Lin M. et al. Asymmetric subgenome selection and cis-regulatory divergence during cotton domestication. *Nat Genet.* 2017; 49.
49. Wendel JF. New World tetraploid cottons contain Old World cytoplasm. *Proc Natl Acad Sci U S A.* 1989; 86: 4132-4136.
50. Yongming H, Peixuan D, Junji C, Posmentier ES. Multivariate analysis of heavy metal contamination in urban dusts of Xi'an, Central China. *Science of the Total Environment.* 2006; 355: 176-186.
51. Žáková M, Benková M. Characterization of spring barley accessions based on multivariate analysis. *Commun. Biom. Crop Sci.* 2006; 1: 124-134.
52. Zhang JF, Fang H, Zhou HP. et al. Genetics, Breeding, and Marker-Assisted Selection for Verticillium Wilt Resistance in Cotton. *Crop Sci.* 2014; 54: 1289-1303.

Figures

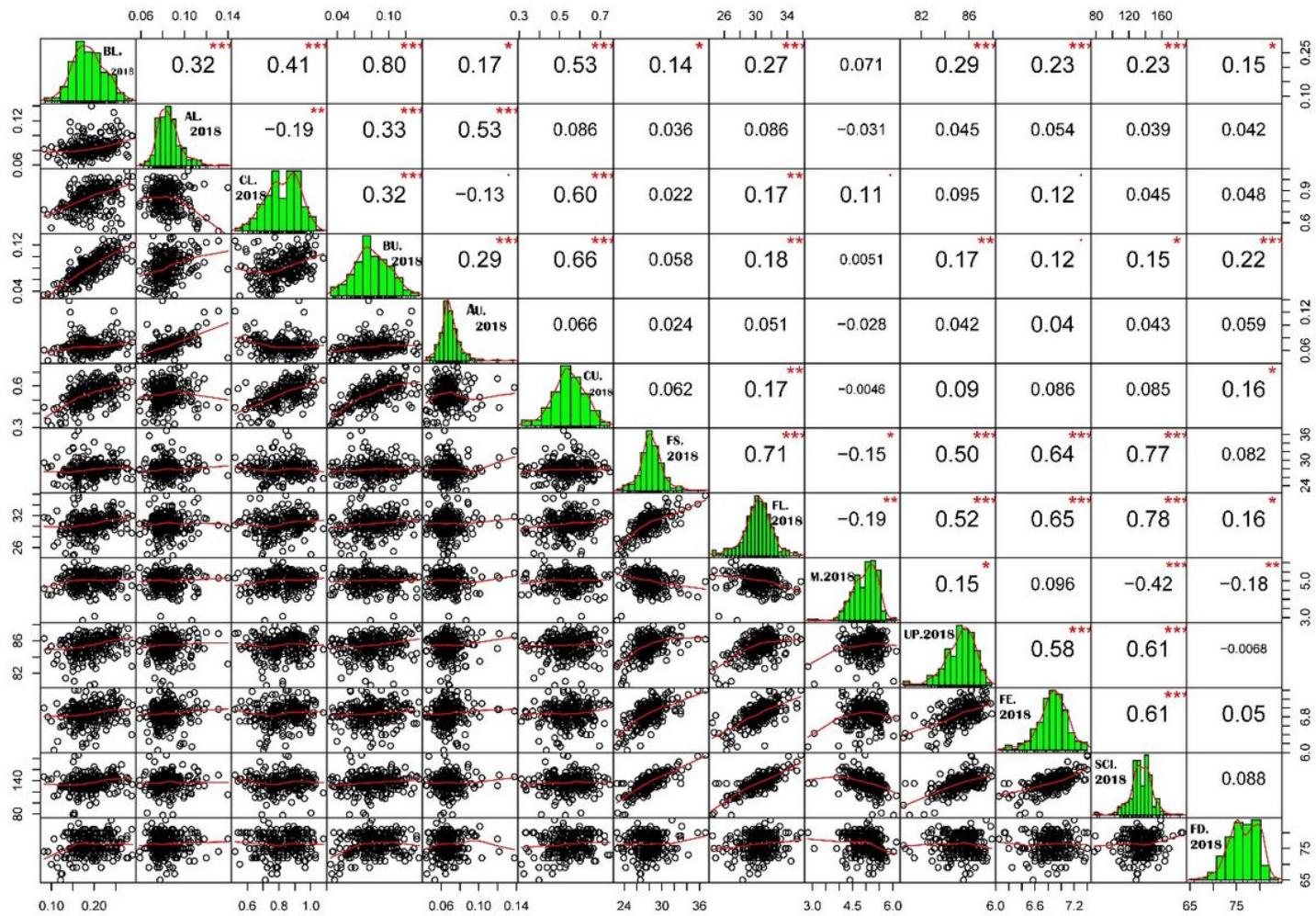


Figure 1

Frequency distribution of phenotypic variation of six stem hardness related traits BL: bending lower; AL: acupuncture lower; CL: compression lower; BU: bending upper; AU: acupuncture upper; CU: compression upper; and seven fiber-related traits and correlation coefficients among the traits in 237 accessions. FS: fiber strength; FL: fiber length; MI: micronaire Index; E: elongation ratio; UP: Uniformity percentage; FE: fibre elongation; SCI: spinning consistency index; FD: days to flowering. *** Indicates extremely significant difference at $P = 0.05$ probability levels.

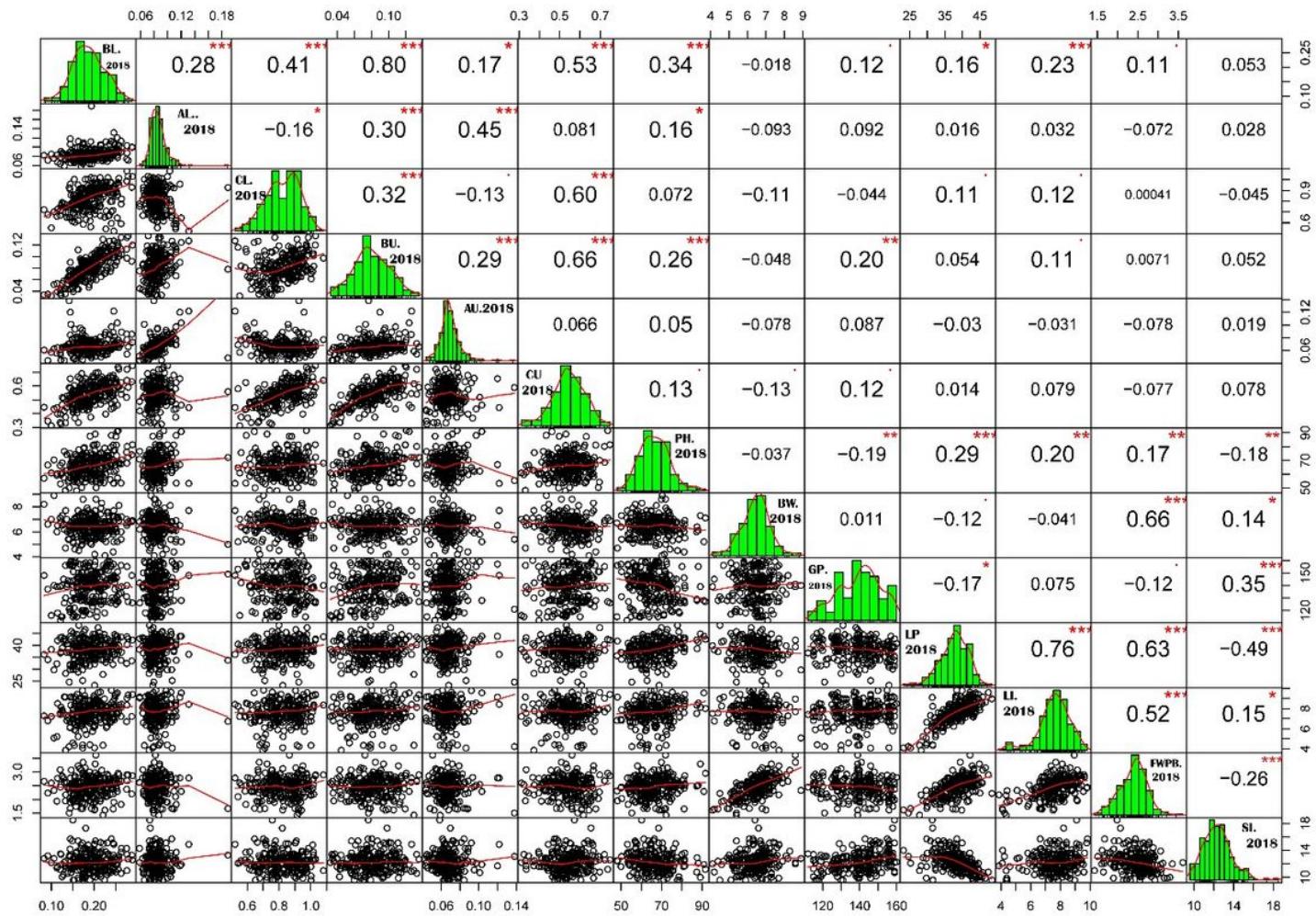


Figure 2

Frequency distribution of phenotypic variation of six-stem hardness related traits BL: bending lower; AL: acupuncture lower; CL: compression lower; BU: bending upper; AU: acupuncture upper; CU: compression upper; and seven yield-related traits and correlation coefficients among the traits in 237 accessions. PH: plant height; BW: boll weight; GP: growth period; LP: lint percentage; LI: lint index; SI: seed index index; FWPR: flower weight per boll. *** Indicates extremely significant difference at $P = 0.05$ probability levels.

Supplementary Files

This is a list of supplementary files associated with this preprint. Click to download.

- [supplementarytablev2.xlsx](#)
- [supplementaryfigure.docx](#)