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DOMINANCE AND HETEROTIC EFFECTS FOR BOLL NUMBER AND BOLL WEIGHT IN UPLAND COTTON HYBRIDS

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SUMMARY

The succeeding study sought to determine the mean performance of parental cultivars and heterotic effects in F_1 hybrids of upland cotton (*Gossypium hirsutum* L.). Six parental genotypes (Guliston, AN-Boyovut-2, Buxoro-102, Yuksalish, Shodlik-11, and Kelajak) underwent crossing to develop 30 complete diallel F_1 hybrids and evaluation in comparison with the standard cultivar (Guliston). The experiment layout in a randomized complete block design (RCBD) had three replications. Analysis of variance revealed significant differences among the parental genotypes and F_1 hybrids for most traits, except the staple length. The F_1 hybrids Guliston \times AN-Boyovut-2 and AN-Boyovut-2 \times Buxoro-102 produced the most bolls per plant, while the F_1 hybrids Guliston \times Yuksalish, Guliston \times AN-Boyovut-2, and Shodlik-11 \times AN-Boyovut-2 showed the maximum boll weight. Parental genotype performance individually did not reliably predict the hybrid performance, indicating the complex genetic interactions in F_1 hybrids. The heterosis analysis showed the hybrids Guliston \times AN-Boyovut-2 and AN-Boyovut-2 \times Buxoro-102 exhibited relative heterosis exceeding 15%–20% and heterobeltiosis above 10%–15% for bolls per plant and boll weight, respectively. The considerable heterotic effects in the F_1 hybrids for bolls per plant and boll weight suggest that utilizing heterosis breeding could be effective.

Keywords: Upland cotton (*G. hirsutum* L.), hybrids, heterosis, heterobeltiosis, dominance, bolls per plant, boll weight, yield-related traits

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Key findings: The upland cotton (G. hirsutum L.) hybrid Gulistan \times AN-Boyovut-2 showed the best performance for boll weight and bolls per plant in the F_1 generation and proved to be the promising genotype for further improvement through breeding programs.

INTRODUCTION

Cotton (Gossypium L.) is the most significant textile crop globally, renowned for its high-quality natural fiber production. During 2017-2018, global cotton production and utilization estimates were at 25.1 million tons, with forecasts predicting an increase to 26.1 million tons by 2026. Cotton supports the textile industry but also plays a vital role in the edible oil industry by providing high-quality cottonseed oil. Globally, approximately 350 million people participate in cotton-related activities, includina production, ainnina, transportation, storage, and textile manufacturing that utilizes cotton fibers. The global cotton market, valued at USD 20 billion annually, thrives due to the unique contribution of the Gossypium genus, which produces single-celled, seed-borne, and longlint fibers (Shavkiev et al., 2023; Khamdullaev et al., 2025).

The leading cotton-producing countries are China, India, and the United States, followed by Pakistan, Brazil, Australia, Uzbekistan, Turkey, Turkmenistan, Burkina Faso, Mali, Greece, and Myanmar. The upland cotton (*G. hirsutum* L.) is a vital cash crop in tropical and subtropical regions. It is crucial in the global textile industry, contributing to an economic impact of USD 600 billion annually (Aslam *et al.*, 2020; Amanov *et al.*, 2020; Samanov *et al.*, 2024; Shavkiev *et al.*, 2025).

In Uzbekistan, cotton is a key cash and industrial crop, being a primary source of fiber, food, and feed. Cotton fibers significantly boost the country's economy. Uzbekistan ranks fourth worldwide for cotton cultivation area and production. However, in Uzbekistan, the yield per unit area remains relatively low compared with other cotton-growing nations (Sanaev et al., 2021; Khamdullaev et al., 2021; Shavkiev et al., 2023). Efforts to develop high-yielding cotton cultivars with improved fiber quality are ongoing to address these constraints and enhance the seed cotton

yield (Nabiyev et al., 2020; Shavkiev et al., 2020, 2022; Chorshanbiev et al., 2023; 2025).

Plant breeding plays a pivotal role in enhancing genetic variation, improving the potential of genotypes, and developing stress resistance to increase crop productivity. Conventional breeding involves hybridization, such as the crossing of different genotypes to develop new high-yielding cultivars with desirable traits for passing on to future generations (Shavkiev *et al.*, 2019a, 2019b; Rajabov *et al.*, 2024).

Breeding research explores possibilities to boost the yield, including the utilization of heterosis (Khan et al., 2017; Narimonov et al., 2023). Heterosis has long been a primary objective in cotton breeding programs. Numerous studies have documented the yield advantages of hybrids over midbetter-parent, and parent, commercial cultivars. However, the major constraints in utilizing heterosis in cotton include the unavailable efficient crossing system, and therefore, producing F₁ hybrid seeds through hand emasculation and pollination is laborintensive (Azimov et al., 2024a).

In cotton, heterosis can enhance seed cotton yield by 10%-20% while also improving fiber quality. A recent review reported that F_1 hybrids exhibited an average useful heterosis of 21.4%, whereas F_2 hybrids averaged 10.7% (Meredith, 1998). Both F_1 and F_2 hybrids demonstrated significantly higher yields than high-yielding parental genotypes and commercial cultivars (Wu *et al.*, 2004).

Heterosis refers to the superior performance of F_1 hybrids relative to their parental cultivars. Though positive heterosis is typically desirable, negative heterosis can be advantageous for specific traits, such as plant height, days to first flowering, micronaire, and gossypol content. For effective hybrid development, a heterosis level of 50% over the promising cultivar and 20% over the best hybrid is considered optimal (Batool and Khan, 2012).

Heterosis is a complex multigenic trait, resulting from extensive gene interaction and physiological processes. F1 hybrids exhibit heterozygosity, which counteracts inbreeding depression. Past studies have shown the causes of heterosis include the cumulative effects of multiple physiological and phenotypic traits (Azimov et al., 2024b). Additionally, Rosas et al. (2010) emphasized the role of gene expression among the parental genotypes and its impact on the phenotype, and the F₁ hybrids mostly exhibit enhanced growth and performance. The presented study aimed to evaluate the performance of F₁ hybrids in comparison with their parental genotypes and the heterosis in F₁ hybrids for bolls per plant and boll weight in upland cotton.

MATERIALS AND METHODS

Genetic material

The breeding material consisted of six parental genotypes of upland cotton ($Gossypium\ hirsutum\ L.$) and their 30 F₁ hybrids, developed using a 6 × 6 complete diallel-crossing system, along with a standard cultivar (Guliston) used as the control. The parental cultivars (Guliston, AN-Boyovut-2, Bukhara-102, Yuksalish, Shodlik-11, and Kelajak) representing a diverse genetic base sustained evaluation for key traits, including the number of bolls per plant and boll weight.

Experimental design and field procedure

The field experiments conducted began during the crop seasons of 2022–2023 and 2023–2024 in Sirdaryo, Uzbekistan (40°25′N, 68°40′E). The study comprises two phases: a crossing block and an evaluation of the F_1 populations of upland cotton. In April 2022, six upland cotton genotypes reached planting in a non-replicated crossing block. Each plot consisted of five rows measuring 25 m long, with plant and row spacings of 60 cm and 90 cm, respectively. Performing hand emasculation and crossing continued in a complete diallel fashion. The resulting 30 F_1 populations proceeded to plant in comparison

with parental genotypes and the check cultivar (Guliston) in April 2023 in a randomized complete block design (RCBD) with three replications.

In F₁ hybrids, the experimental plots consisted of a single row, 5.3 m in length, containing 25 plants per replication. Plant and row spacings remained at 20 cm and 90 cm, respectively. Recommended agronomic practices, including fertilization, hoeing, irrigation, and pest control, were uniform in application to all entries to minimize field variations. Harvesting ensued in October-November, with ginning performed using the eight-saw gin to obtain lint for further analysis.

Traits' measurement and statistical analysis

The data recorded comprised the randomly selected plants in each experimental unit for the number of bolls per plant and boll weight (g). The mean data incurred analysis of variance (ANOVA) following Steel et al. (1997) to test the null hypothesis of no differences among the F₁ hybrids and parental cultivars in comparison with the standard cultivar. Statistical parameters, including the Fisher criterion (F), standard deviations (S.D.), standard error (S.E.), and significance levels (P \leq 0.05*, P \leq 0.01**, and P \leq 0.001***), underwent evaluation of the reliability of the cotton genotype differences for each trait. In F₁ hybrids, the dominance coefficient determination was according to Griffing

$$hp = \frac{F_1 - MP}{P - MP}$$

Where hp = dominance coefficient; F_1 = the evaluated arithmetic mean of the hybrid; MR = the evaluated arithmetic mean of both parents; and R = the evaluated arithmetic mean of the best parents.

In F_1 hybrids, detecting the heterotic effects over mid- and better-parents followed the method of Mather and Jinks (1982):

Heterosis (H%) =
$$([F_1 - MP] / MP) \times 100$$

(1950):

Heterobeltiosis (Hb%) = ($[F_1 - BP] / BP$) × 100

Where F_1 is the mean value of the F_1 hybrid and MP is the mean value of two parental genotypes involved in the cross, while BP is the mean value (over replications) of the better parents of the particular cross.

The correlation coefficient (r), as determined among the various variables, used the approach described by Snedecor and Cochran (1981). In this analysis, the correlation value less than 0.3 indicates a weak relationship between traits, the correlation value between 0.3 and 0.7 indicates a moderate correlation, while the correlation value greater than 0.7 was considerably a strong association.

RESULTS

Bolls per plant

The evaluation of six cotton parental cultivars and their F_1 hybrids succeeded for the total number of bolls per plant, as generated through diallel crossing (Table 1). Compared with the control cultivar (Guliston), the cultivars Yuksalish and Shodlik-11 exhibited higher boll numbers (28.8 \pm 2.4 and 30.0 \pm 3.9 bolls per plant, respectively). In contrast, the cultivar AN-Boyovut-2 had the lowest boll number (18.6 \pm 1.1).

Among the F_1 hybrid combinations, the most bolls per plant resulted in hybrids Guliston \times Yuksalish, Yuksalish \times Guliston, Yuksalish \times Bukhara-102, and AN-Boyovut-2 \times Shodlik-11, ranging between 26 and 28 bolls per plant (Table 1). Conversely, the F_1 hybrids, such as Bukhara-102 \times AN-Boyovut-2, Bukhara-102 \times Kelajak, Shodlik-11 \times Guliston, Kelajak \times AN-Boyovut-2, Bukhara-102 \times Yuksalish, and AN-Boyovut-2 \times Guliston, demonstrated the fewest bolls per plant (<20).

Overdominance in bolls per plant

Overdominance in bolls per plant is a genetic phenomenon that can be beneficial in improving seed cotton yield. Overdominance is a model suggesting that the interaction of alleles at a single heterozygous locus can result in a synergistic effect surpassing both homozygous parental genotypes. For bolls per plant, overdominance was evident in the F₁ hybrids, such as Guliston × Bukhara-102, Guliston × AN-Boyovut-2, AN-Boyovut-2 × Bukhara-102, AN-Boyovut-2 × Kelajak, and Kelajak × Guliston (Table 1). Conversely, the hybrids, including Guliston × Shodlik-11, Yuksalish × Shodlik-11, Bukhara-102 × Kelajak, AN-Boyovut-2 × Guliston, AN-Boyovut-2 × Yuksalish, Shodlik-11 × Guliston, Shodlik-11 × Yuksalish, Shodlik-11 × Kelajak, and Kelajak × Shodlik-11, exhibited negative overdominance.

Heterotic effects for bolls per plant

By assessing the heterotic effects of F₁ hybrids relative to their parental lines for the bolls per plant, significant and moderate heterotic effects appeared (Table 1). Hybrids, such as Guliston × AN-Boyovut-2 and AN-Boyovut-2 × Bukhara-102, displayed notable heterosis for the number of bolls per plant. Based on the analysis, the F_1 hybrids Guliston \times Bukhara-102, Guliston × AN-Boyovut-2, AN-Boyovut-2 × Bukhara-102, AN-Boyovut-2 × Kelajak, and Kelajak × Guliston occurred with superior genetic and breeding values. These hybrids exhibited the higher boll number, dominance, and remarkable heterotic effects, arising as promising cross combinations for further breeding programs.

Boll weight

Among the parental cotton cultivars, the heaviest boll weight recording was in cultivar Bukhara-102, where the boll weight per boll exceeded 8.0 g (Table 2). Conversely, the lightest boll weight manifested in cultivar Yuksalish, with an average of 6.2 ± 0.1 g boll weight. In F₁ hybrids Guliston × Shodlik-11, Guliston × Kelajak, Yuksalish × AN-Boyovut-2, Yuksalish × Shodlik-11, Yuksalish × Kelajak, Shodlik-11 × Guliston, and Shodlik-11 × Yuksalish, the boll weight ranged from 6.0 to 7.0 g. In the remaining F₁ hybrids, boll weight 7.0 g, indicating performance in these cross combinations.

Table 1. Degree of dominance and heterotic effects for bolls per plant in parental genotypes and F_1 hybrids of upland cotton.

Parental genotypes and F ₁ hybrids	M±SE	S.D.	Dominance Coefficient	Heterosis	Heterobeltiosis
Guliston (control cultivar)	20.3±1.4	5.8	-	-	-
AN-Boyovut-2	18.6±1.1	4.1	-	-	-
Buxoro-102	20.7±2.0	6.8	-	-	-
Yuksalish	28.8±2.4	14.6	-	-	-
Shodlik-11	30.0±3.9	12.9	-	-	-
Kelajak	23.2±1.4	4.8	-	-	-
Guliston x Yuksalish	26.6±2.5	10.5	0.5	8.4	-7.6
Guliston x Buxoro-102	21.3±1.3	6.9	4.0	3.9	2.9
Guliston x AN-Boyovut-2	22.6±1.4	6.6	3.7	16.2	11.3
Guliston x Shodlik-11	22.1±1.4	8.0	-1.4	-12.1	-26.3
Guliston x Kelajak	23.0±1.3	7.3	0.9	5.7	-0.9
Yuksalish x Guliston	26.8±1.8	6.9	0.5	9.2	-6.9
Yuksalish x Buxoro-102	26.4±1.4	5.8	0.4	6.7	-8.3
Yuksalish x AN-Boyovut-2	21.7±2.0	10.4	-0.4	-8.4	-24.7
Yuksalish x Shodlik-11	21.5±1.0	5.3	-13.2	-26.9	-28.3
Yuksalish x Kelajak	25.4±1.2	6.3	-0.2	-2.3	-11.8
Buxoro-102 x Guliston	20.5±1.0	5.8	0.0	0.0	-1.0
Buxoro-102 x Yuksalish	19.8±0.9	5.5	-1.2	-20.0	-31.3
Buxoro-102 x AN-Boyovut-2	18.3±0.8	4.9	-0.1	-6.9	-11.6
Buxoro-102 x Shodlik-11	21.2±1.3	7.4	-0.1	-16.4	-29.3
Buxoro-102 x Kelajak	18.5±0.9	5.2	-2.8	-15.7	-20.3
AN-Boyovut-2 x Guliston	18.5±0.8	4.4	-1.1	-4.9	-8.9
AN-Boyovut-2 x Yuksalish	21.3±1.0	5.3	-5.8	-10.1	-26.0
AN-Boyovut-2 x Buxoro-102	23.6±1.3	7.0	2.9	20.1	14.0
AN-Boyovut-2 x Shodlik-11	28.1±1.7	6.8	0.7	15.6	-6.3
AN-Boyovut-2 x Kelajak	23.6±2.1	9.1	1.2	12.9	1.7
Shodlik-11 x Guliston	19.9±0.7	3.8	-1.0	-20.9	-33.7
Shodlik-11 x Yuksalish	20.5±1.2	7.2	-14.8	-30.3	-31.7
Shodlik-11 x Buxoro-102	22.6±1.4	6.5	-0.6	-10.8	-24.7
Shodlik-11 x AN-Boyovut-2	21.4±1.1	6.3	-0.5	-11.9	-28.7
Shodlik-11 x Kelajak	20.5±1.2	6.7	-1.8	-22.9	-31.7
Kelajak x Guliston	24.2±1.2	4.6	1.7	11.3	4.3
Kelajak x Yuksalish	24.8±1.4	6.7	-0.4	-4.6	-13.9
Kelajak x Buxoro-102	21.6±1.2	5.6	-0.3	-1.6	-6.9
Kelajak x AN-Boyovut-2	19.7±1.0	5.8	-0.5	-5.7	-15.1
Kelajak x Shodlik-11	23.1±1.2	6.3	-1.0	-13.2	-23.0

Table 2. Degree of dominance and heterotic effects for boll weight in parental genotypes and their F_1 hybrids in upland cotton.

Parental genotypes and F ₁ hybrids	M±SE	S.D.	Dominance Coefficient	Heterosis	Heterobeltiosis
Guliston (control cultivar)	7.1±0.26	0.46	=	-	-
AN-Boyovut-2	7.1±0.15	0.26	-	_	-
Buxoro-102	8.1±0.12	0.21	-	_	-
Yuksalish	6.2 ± 0.1	0.17	-	-	-
Shodlik-11	7.2±0.09	0.15	-	-	-
Kelajak	7.5±0.23	0.40	-	_	-
Guliston x Yuksalish	7.5±0.28	0.49	2.0	12.8	5.6
Guliston x Buxoro-102	7.7 ± 0.1	0.17	0.1	1.3	-4.9
Guliston x AN-Boyovut-2	7.6 ± 0.18	0.31	54.0	7.0	7.0
Guliston x Shodlik-11	6.4±0.38	0.66	-18.2	-10.4	-11.1
Guliston x Kelajak	6.9±0.35	0.60	-1.7	-5.5	-8.0
Yuksalish x Guliston	7.2±0.27	0.47	1.2	8.3	1.4
Yuksalish x Buxoro-102	7.1 ± 0.07	0.12	0.0	-0.7	-12.3
Yuksalish x AN-Boyovut-2	6.8±0.45	0.78	0.4	2.3	-4.2
Yuksalish x Shodlik-11	6.9±0.29	0.50	0.4	3.0	-4.2
Yuksalish x Kelajak	6.8±0.3	0.51	-0.1	-0.7	-9.3
Buxoro-102 x Guliston	7.7±0.38	0.66	0.2	1.3	-4.9
Buxoro-102 x Yuksalish	7.2±0.18	0.31	0.1	0.7	-11.1
Buxoro-102 x AN-Boyovut-2	7.7±0.29	0.50	0.4	1.3	-4.9
Buxoro-102 x Shodlik-11	7.9±0.5	0.87	0.6	3.3	-2.5
Buxoro-102 x Kelajak	8.3±0.1	0.17	1.8	6.4	0.0
AN-Boyovut-2 x Guliston	7.3±0.27	0.47	19.0	2.8	1.4
AN-Boyovut-2 x Yuksalish	7.1±0.52	0.90	1.0	6.8	0.0
AN-Boyovut-2 x Buxoro-102	7.1±0.32	0.55	-0.8	-6.6	-12.3
AN-Boyovut-2 x Shodlik-11	7.1±0.1	0.17	-0.4	-0.7	-1.4
AN-Boyovut-2 x Kelajak	7.2±0.17	0.29	-0.2	-1.4	-4.0
Shodlik-11 x Guliston	6.5±0.13	0.23	-14.5	-9.1	-13.3
Shodlik-11 x Yuksalish	6.4±0.34	0.59	-0.5	-4.5	-11.1
Shodlik-11 x Buxoro-102	7±0.12	0.20	-1.4	-8.5	-13.6
Shodlik-11 x AN-Boyovut-2	7.5±0.03	0.06	7.0	4.9	4.2
Shodlik-11 x Kelajak	7.6±0.09	0.15	1.6	3.4	1.3
Kelajak x Guliston	7.7±0.59	1.03	2.1	5.5	2.7
Kelajak x Yuksalish	7.4±0.09	0.15	1.9	8.0	-1.3
Kelajak x Buxoro-102	8.2±0.35	0.61	1.0	5.1	1.2
Kelajak x AN-Boyovut-2	7.5±0.4	0.70	-0.1	2.7	0.0
Kelajak x Shodlik-11	7.5±0.85	1.47	1.1	2.0	0.0

Overdominance in boll weight

For boll weight, positive overdominance resulted in the F_1 hybrids Guliston \times Yuksalish, Guliston \times AN-Boyovut-2, Yuksalish \times Guliston, Bukhara-102 \times Kelajak, AN-Boyovut-2 \times Guliston, AN-Boyovut-2 \times Yuksalish, Shodlik-11 \times AN-Boyovut-2, Shodlik-11 \times Kelajak, Kelajak \times Guliston, Kelajak \times Yuksalish, Kelajak \times Bukhara-102, and Kelajak \times Shodlik-11 (Table 2). However, the negative overdominance was notable in the F_1 hybrids, such as Guliston \times Shodlik-11, Guliston \times

Kelajak, Shodlik-11 \times Guliston, and Shodlik-11 \times Bukhara-102.

Heterotic effects for boll weight

In F_1 hybrids, the heterotic effect for boll weight, compared with their parental lines, revealed significant and moderate heterosis (up to 10%) (Table 2). The promising F_1 hybrids exhibited heterotic effects, including Guliston \times Yuksalish, Guliston \times AN-Boyovut-2, Yuksalish \times Guliston, AN-Boyovut-2 \times Guliston, Shodlik-11 \times AN-Boyovut-2, Shodlik-

11 × Kelajak, Kelajak × Guliston, and Kelajak × Bukhara-102. Based on the analysis, the F_1 hybrids Guliston × Yuksalish, Shodlik-11 × AN-Boyovut-2, Shodlik-11 × Kelajak, Kelajak × Guliston, and Kelajak × Bukhara-102 came out with better genetic potential.

DISCUSSION

The F_1 hybrids, obtained through the crosses Guliston \times Yuksalish, Shodlik-11 \times AN-Boyovut-2, Shodlik-11 \times Kelajak, Kelajak \times Guliston, and Kelajak \times Bukhara-102, demonstrated the maximum boll weight, averaging 7.0 g. Boll weight is a crucial yield component, and many assume it positively influences the seed cotton yield if the number of bolls per plant remains constant. These findings align with previous research, which suggested that significant heterotic effects for boll weight may be due to both the additive and non-additive gene effects in upland cotton (Khan and Qasim, 2012).

The results also indicate the parental cultivars Kelajak and Shodlik-11, along with the testers Guliston and AN-Boyovut-2, were the best general combiners, making them highly suitable for hybridization and breeding programs. An increase in the number of bolls per plant in cotton significantly enhances the overall seed cotton yield, demonstrating a strong positive correlation between the number of bolls per plant and seed cotton yield. In terms of heterotic effects, the F₁ hybrids Guliston × Yuksalish, Shodlik-11 × AN-Boyovut-2, Shodlik-11 × Kelajak, Kelajak × Guliston, and Kelajak × Bukhara-102 exhibited higher relative heterosis and heterobeltiosis than the other hybrids.

These hybrids expressed superior hybrid vigor for the number of bolls per plant, making them promising candidates for hybrid crop development. Past studies also reported the highest heterosis over the better parent and check cultivars for boll formation, with the said trait found remarkably associated with seed cotton yield in upland cotton (Azimov *et al.*, 2024c). In F₁ hybrids, the heterotic effects for bolls per plant ranged from moderate to high. Dong *et al.* (2007) mentioned the

significant heterosis for boll size and boll number compared with the standard cultivar, while Basal and Turgut (2003) reported considerable heterosis for boll weight in upland cotton.

Since boll weight is a major yieldcontributing factor and has a positive association with seed cotton yield, boll numbers remain constant. Similarly, Yuan et al. (2002) observed average heterosis (13.3%) for boll weight over the mid-parent. However, boll weight differences appeared minimal by comparing with the better parent. Khan et al. (2007, 2009) stated the topmost heterosis over better parents for boll weight, which could mean an improved seed cotton yield. The study results were also consistent with past findings of Kumar et al. (2014), who also noted significant heterotic effects in F₁ and F₂ populations for bolls per plant.

For bolls per plant, the F₁ hybrids Guliston × AN-Boyovut-2 and AN-Boyovut-2 × Bukhara-102 delivered the maximum relative heterosis and heterobeltiosis. Moreover, the F₁ hybrids Guliston × Yuksalish, Guliston × AN-Boyovut-2, and Shodlik-11 × AN-Boyovut-2 showed moderate heterosis over mid and better parents for boll weight. Previous studies have consistently reported the significant heterosis over the mid and better parents for yield-contributing traits, which further supported this study's present results (Komal et al., 2014; Baloch et al., 2015).

CONCLUSIONS

The significance of heterotic effects was a revelation of the substantial mean square differences among the parental genotypes and their F₁ hybrids. The F₁ hybrids outperformed their parental lines for key yield traits due to the expression of heterotic effects. Guliston \times AN-Boyovut-2 and AN-Boyovut-2 × Bukhara-102 produced the higher number of bolls per plant among the F₁ hybrids. For boll weight, the best-performing F₁ hybrids were Guliston × Yuksalish, Guliston × AN-Boyovut-2, and Shodlik-11 × AN-Boyovut-2. Overall, the cross combination Guliston × AN-Boyovut-2 exhibited an excellent performance in the F₁ generation, receiving authentication as a promising candidate genotype for further breeding programs.

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