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### CHARACTERIZATION OF BREAD WHEAT SEGREGATING POPULATIONS UNDER OPTIMUM IRRIGATION AND WATER STRESS CONDITIONS

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#### SUMMARY

Water scarcity is currently threatening almost every country in the arid regions. Using advanced generations, breeding can help in the development of improved bread wheat genotypes for adaptation to abiotic and biotic stresses. The pedigree selection was practiced on two bread wheat crosses (Sids 12 × Line 44) and (Line 20 × Sakha 93) during two seasons (2017-2018 and 2018-2019) under full irrigation (optimal conditions) and limited irrigation (drought-stressed) conditions at Kafer El-Hamam Agriculture Station, Agricultural Research Center, Giza, Egypt. The results indicated significant differences in two crosses of F<sub>2</sub> and F<sub>3</sub> families for all the studied traits under optimal irrigation and water stress. The estimates of phenotypic coefficients of variability (PCV) were slightly higher than those of genotypic coefficients of variability (GCV) for all the traits in two crosses of both water regimes. Broad-sense heritability  $(h_{Bs}^2)$  estimates, accompanied with high magnitudes of the genetic advance (GA), were higher under optimal irrigation than water stress in  $F_2$  and  $F_3$  generations of two crosses. A positive correlation was recorded between spikes per plant and grain yield in both water treatments of two crosses. A positive correlation (r) was revealed between offsprings ( $F_3$ ) and their parents ( $F_2$ ) in yield and its components under optimal irrigation and water stress conditions. Hence, the hybridization followed by selection under optimal and drought stress conditions have been a demand to accelerate the genetic gain of wheat grain yield.

**Keywords:** Wheat, selection parameters, heritability, genetic advance, water stress

**Key findings:** Climate change is the main threat to the wheat crop; thus, breeding strategies for drought consist of a large number of segregating wheat crosses to derive new pure lines under optimal and drought stress conditions.

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### INTRODUCTION

Wheat (*Triticum aestivum* L.) is an important cereal field crop feeding one-fifth of the world's population, and the wheat production should be doubled by 2050 to fulfill the demand for an increasing population and decreasing land for

cultivation; hence, the need to increase the yield per unit area (Alotaibi *et al.*, 2021; IPCC, 2013). Present water shortages impair nearly every country in the arid Mediterranean regions. Most significantly, water shortages in the world's arid regions continue to worsen due to sudden climatic changes, rising incomes,

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and continuous population growth (El-Hendawy et al., 2017; Swailam et al., 2021). Most of the current water-limited countries will become warmer and drier due to climate change (Darzi-Ramandi et al., 2016; Ali et al., 2021). Moreover, climate change is expected to increase the temperature by 3 °C-5 °C. reducing rainfalls by 4%-27% during the cropping season (IPCC, 2013), which will drastically affect crop production (Shauka et al. 2021). Thus, under climate changes, the demand for irrigation water in the future is generally expected to increase over time (Kamruzzaman et al., 2020). Climate change is the main threat to most crops; therefore, scientific and political contexts have gained increasing importance for this event (Sallam et al., 2019; Tembo, 2021). The drought stress effect is one of the consequences of climate change that harms field crop growth and yield (Ali, 2017; Sallam et al., 2019).

Drought reduced the wheat grain yield by 62% (Aberkane et al., 2020), and gene introgression from wild relatives pays off and can increase wheat genotype's resilience to climate change effects. Drought tolerance is a polygenic character, and genetic studies can help control drought tolerance (Gazal et al., 2016; Sallam et al., 2019). In 2030, wheat demand is expected to reach up to 40% with the continuous growth in human population (Ahmed et al., 2019a). Breeding for drought tolerance is important to know the mechanism and behavior of plants under water stress environments. The drought tolerance mechanism is complex, for reasons considering intensity, period of stress, plant growth stage, and crop species (Ahmed et al., 2019b). In that respect, there are three fundamental mechanisms, i.e., escape avoidance (tolerance) and resistance mechanisms, which a plant can acclimatize to manage drought stress (Ali et al., 2021). The selection of early flowering genotypes has been recognized as enhancing drought stress tolerance through escape (Barakat et al., 2020).

Breeding strategies for drought consist of many segregating wheat crosses to derive new pure lines, which are evaluated and compared with commercial cultivars for high yield and tolerance to biotic and/or abiotic (Al-Ashkar stresses et al., 2021). Morphological and physiological plant characters can significantly influence the productivity of wheat yield in direct and indirect pathways. Productivity is affected by environmental fluctuations when there are a high genetic environment interaction and low heritability that make the selection of genotypes more difficult (Sallam *et al.*, 2019; Mansour *et al.*, 2020; Al-Ashkar *et al.*, 2021).

Stability analysis studies revealed a differential response of wheat genotypes to various drought stress environments, and significant differences found among the bread wheat genotypes for yield components (Ali, 2017; Putri *et al.*, 2020; Al-Ashkar *et al.*, 2021). These studies reported the significance of genetic relationships, such as, selection efficiencies, genetic gain, heritability and coheritability for evaluating trait stability under stress levels and their interactions.

The selection of genetically superior wheat genotypes amongst the huge amount of segregating and recombinant progenies is an essential but complex process in plant breeding. The development of new cultivars plant demands long periods that involve cyclic hybridization and selection techniques (Ali et al., 2020). During the last two decades, crossing and selection have played an important role in wheat improvement, as well as, the transgressive segregation crossing technique using superior characters to genotype by environment interaction (GEI), where an example of this tremendous improvement in the wheat genome was the green revolution, in which the progress was slow but consistent (Alotaibi et al., 2021).

The study of heritability, genetic advance, and genotypic and phenotypic coefficient of variation helps find the magnitude and nature of variation in germplasm. Thus, crop improvement depends on genetic variability in breeding material (Sohail et al., 2018; Ahmed et al., 2019a; Arifuzzaman et al., 2020). Selection of the superior phenotype type is possible only when the genotype (G) and phenotype (Ph) are strongly correlated. A selection program can be predicted when data on heritability are available. Additionally, a high heritability coupled with a high selection estimate response value, is desired for selection (Merida-Garcia *et al.*, 2019; Ali *et al.*, 2020).

Based on the major points stated earlier, the study was conducted to explore more information on genetic variability, heritability, and selection response in bread wheat for its superior genotype development under both full irrigation and water stress conditions, by identifying characteristics having a significant direct effect on grain yield as selection criteria.

### MATERIALS AND METHODS

### Plant material

Two bread wheat crosses, i.e., (Sids  $12 \times \text{Line}$  44) and (Line  $20 \times \text{Sakha 93}$ ), were selected from 15 F<sub>1</sub>'s diallel crosses of previous work (Swelam, 2015). These crosses had the best values for yield components under full irrigation and water stress environments. These crosses were derived from four diverse parental bread wheat genotypes, i.e., two commercially adopted cultivars and two exotic genotypes (Table 1).

### Experimental site

The experiment was carried out during the two successive growing seasons, 2017/2018 and 2018/2019 at the experimental farm at the Kafer El-Hamam Agriculture Station, Agricultural Research Center, Giza, Egypt (30°37'03.3"N, 31°31'03.6"E). The soil mechanical and chemical analyses of the experimental sites are given in Table 2. The monthly weather data during two field trial seasons are shown in Figure 1.

Table 1. Pedigree and origin of the parental	I genotypes used in two bread wheat crosses.
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Genotypes		Pedigree	Origin
	name		
Cross I	Sids 12 (P1)	BUC//7C/ALD/5/MAYA74/ON//1160- 147/3/BB/GLL/4/CHAT"S"/6/MAYA/VUL//CMH74A.630/4*SX.SD7096- 4SD-1SD1SD-0SD.KS82142/2*WBLL1.	ARC, Egypt
	Line 11		CIMMVT
	Line 44 (P2)	CROC_1/AE.SQUARROSA(224)///OPATA/3/SOKOLL	CIMMYT, Mexico
Cross II	Line 20 (P1)	KS82142/2*WBLL1.	CIMMYT, Mexico
	Sakha 93 (P2)	SAKA92/TR810328.	ARC, Egypt

Properties	Sand (%)	Silt (%)	Clay (%)	Texture class	Organic matter (%)	Available (N) (ppm)	Available (P) (ppm)	Available (K) (ppm)	рН
2017/2018	18.7	34.5	46.8	Clay	1.72	44.6	9.8	378.6	8.31
2018/2019	17.5	33.6	48.9	Clay	1.59	36.71	8.4	405.5	8.22



**Figure 1.** Monthly weather data during the field trial at Kafer El-Hamam, Sharkia Governorate, Rain is precipitation (mm day-1), RH: average relative humidity at 2 meters, Tmax: average maximum temperature, Tmin: average minimum temperature, Tmean: temperature average.

### Experimental design

In the first season, the seeds of  $F_2$  for two wheat crosses and their parents were sown field water experiments—optimal under irrigation and water stress conditions-in anonreplicated plots. Each plot had 20 rows of F<sub>2</sub> population alongside eight rows for their parents ( $P_1$  and  $P_2$ ). The rows' lengths were 3 m, with row to row at 30 cm apart, and the distance between any two plants within a row was 10 cm, presenting 600 plants for each population and 120 plants for each parent. Data for earliness characters were collected on about 30 and 40 plants, and then these plants were harvested for each population under optimal irrigation and drought stress, respectively. Grain yield and its components were recorded on each plant. The height of 10 plants was selected based on their desired results for earliness, yield, and components to demonstrate the  $F_3$  generation seeds (10 families for each population under both water treatments). Also, data were recorded on10 plants for each parent for studied characters under the two environments.

In the second season, the field experiment was conducted to evaluate 10 families under optimal irrigation and another 10 families under water stress conditions, and compared with their parents in strip design, with a randomized complete block design (RCBD) in two replicates. Each plot consisted of 24 rows of 10 families and their parents (P<sub>1</sub> and  $P_2$ ), two rows of each population. The row length was 3 m, with row to row at 30 cm apart, and the distance between any two plants within a row was 10 cm, presenting 600 plants for each population (60 plants for each family  $\times$  10 families) and 60 plants for each parent. In each replicate, data were collected for days to heading (DH), plant height (PH), and days to maturity (DM) characters on 10 guarded and competitive plants for parents (P<sub>1</sub> and  $P_2$ ) and 100 plants for each  $F_3$  population (10 plants for each family) under the two water treatments. These plants were harvested and measured with the following traits, i.e., spikes per plant (S/P), grains per spike (G/S), 1000grain weight (TGW), and grain yield per plant (GY/P), recorded for individual plants. The means of the 10 plants were subjected to statistical and genetic analysis.

The best three families (selection intensity 30%) were selected, and the best five plants from each selected family were chosen to represent 15 families of the  $F_4$  generation,

including 15 families from each cross that will be completed in the subsequent feature work of obtaining improved yielded lines of wheat under two ways, the optimal and water stress conditions.

### Statistical procedures

Means and variances for the parents and  $F_2$  generation studied under the two irrigation conditions were statistically estimated, while data of the tested families in  $F_3$  were statistically analyzed in RCBD as described by Snedecor and Cochran (1967). Genotypic and phenotypic mean squares ( $\sigma^2$ g and  $\sigma^2$ Ph) were estimated by the method of Singh and Choudhary (1987) and Wricke and Weber (1986) as  $\sigma^2$ g = (MSg - MSI) / r and  $\sigma^2$ Ph = Vg + MSI. The genotypic and phenotypic coefficient of variations (GCV % and PCV %) were estimated according to Johnson *et al.* (1955) as follows:

$$\text{GCV } \% = \left(\frac{\sqrt{\sigma^2 g}}{\overline{X}}\right) \times 100$$

and

$$\mathsf{PCV} \ \% = \left(\frac{\sqrt{\sigma^2 \mathbf{Ph}}}{\overline{\mathbf{X}}}\right) \times 100^{\Box}$$

Where,  $\overline{\mathbf{X}}$  is grand mean. GCV and PCV values were classified according to Deshmukh et al. (1986) as follows: 0% - 10%, low; 10 % -20%, moderate, and >20%, high. Broad sense heritability  $(h_b^2)$  was described by Allard (1960) as  $h_b^2 = (\sigma^2 g / \sigma^2 P) \times 100$ . According to Singh (2001), broad-sense heritability was categorized as 0% - 40%, low; 40% - 59%, medium; 60% - 79%, moderately high, and >80%, very high. Genetic advance (GA) was estimated according to the method (Johnson et al., 1955) as GA =  $h_b^2 \times K \times \sqrt{VP}$  where K is 2.67 and 1.14 at 5% and 30% selection intensities. Realized selection differential (observed response) =  $\bar{X}_p - \bar{X}_0$  (this relates to the same generation), where,  $X_p$  = mean of selected genotypes (families) for nextgeneration and  $\mathbf{X}_{o}$  = mean of bulk population. Realized response in  $F_3$  = mean  $F_3$  - mean  $F_2$ . Correlation (r) between offspring (F<sub>3</sub>) and parents (F<sub>2</sub>) = SP<sub>xy</sub>/ $\sqrt{SS_x \times SS_y}$ . Regression (b) of offspring (F<sub>3</sub>) on parents (F<sub>2</sub>) = SP<sub>xy</sub>/SS<sub>x</sub>.

Cross	Sources of variation	d f	Optimal	irrigation					
Cross	Sources of variation	d.f	DH	DM	PH	NS/P	NG/S	TGW	GY/P
Cross I	Replications	1	$0.72^{*}$	$2.89^{*}$	$160.18^*$	0.29	33.8	5.21*	26.68
	Families	9	$0.56^{**}$	$12.27^{**}$	239.70 <sup>**</sup>	$5.65^{**}$	$63.01^{**}$	14.93**	28.49**
	Error	9	0.1	1.28	22.38	0.58	11.28	1.96	6.92
Cross II	Replications	1	0	1.06	1.68	0.29	20.69	22.35	1.1
	Families	9	3.90**	3.58**	92.60**	7.03**	34.91**	126.59**	$4.98^{*}$
	Error	9	0.3	0.41	8.34	0.26	4.56	12.49	1.07
			Water st	ress					
Cross I	Replications	1	2.24	0.11	1.8	0.18	25.54	7.42	8.1
	Families	9	2.02**	$1.75^{**}$	$178.31^{**}$	$3.23^{**}$	52.62**	$16.19^{**}$	$10.29^{*}$
	Error	9	0.48	0.32	29.24	0.69	11.62	2.95	3.08
Cross II	Replications	1	0.34	0.22	8.19	0.05	12.77	0.04	0.04
	Families	9	3.02**	$2.80^{**}$	$57.45^{*}$	2.93**	30.24*	$6.88^{*}$	$9.14^{*}$
	Error	9	0.27	0.42	13.82	0.51	5.88	1.32	2.28

**Table 3.** Analysis of variance of earliness, plant height, and grain yield components for  $F_3$  families under optimal irrigation and water stress conditions for two crosses.

DH: days to heading, DM: days to maturity, PH: plant height, S/P: spikes per plant, G/S: grains per spike, TGW: 1000-grain weight, and GY/P: grain yield per plant. Cross I is (Sids 12 × Line 44) and Cross II is (Line 20 × Sakha 93).\* P < 0.05 and \*\* P < 0.01.

Coefficient of the determinant (R) of selection response =  $r^2 \times 100$ . Microsoft Excel and SAS 9.2 (2008) computer programs for Windows were used for the statistical analysis.

## RESULTS

### Mean squares and mean performance

The analysis of variance showed that mean squares from DH, DM, PH, NS/P, NG/S, TGW, and GY/P of  $F_3$  generation families for two wheat crosses were significant to highly significant in both the optimal irrigation and water stress conditions (Table 3). The average earliness, plant height, and grain yield components in the  $F_2$  (selected on the bias of the high grain yield per plant) and  $F_3$  generations under optimal irrigation and water stress treatments for the two crosses are shown in Tables 4 and 5.

In general, wheat plants for  $F_2$  and  $F_3$ generations respond to moisture water stress with the changes in various traits for two crosses. In other words, water deficit reduced the means of two generations compared with optimal irrigation of all studied traits for two wheat crosses. Days to heading and days to maturity increased in  $F_3$  generation compared with  $F_2$  generation for all crosses under both environments. In contrast, in most families of cross II under the two water treatments, the  $F_2$ plants were taller than their offspring. In comparison, the mean behavior of the plant height fluctuated between the families in the second and third generations in cross I under optimal irrigation and water stress.

The spikes plant fluctuated among families of offspring  $(F_3)$  and their parents  $(F_2)$ in cross I under optimal irrigation and water stress, and cross II under water stress. In contrast, most F<sub>2</sub> families had more spikes per plant than F<sub>3</sub> in cross II under optimal irrigation. The grains per spike decreased in most  $F_3$  families compared with their  $F_2$  families for two crosses under both environments. On the other hand, most  $F_3$  families were greater than  $F_2$  in all crosses for 1000-grain yield under optimal irrigation and water stress. For grain yield per plant, most of the selected F<sub>2</sub> families had higher values than F<sub>3</sub> generation for two crosses in both water environments. Therefore, direct selection for grain yield is not practical in early generations for developing drought-tolerant and high-yielding varieties.

## Genetic and selection parameters

# Populations of the cross I (Sids 12 × Line 44)

## F<sub>2</sub> Generation

Results showed that under optimal irrigation, days to heading varied from 88.0 to 92.0 with an average of 89.7 days. Days to maturity ranged from 143.0 to 147.0 with an average of 145.0 days. Plant height ranged between 90.0 and 143.0 cm with an average of 121.5 cm. Spikes per plant diverse from 2.0 to 25.0 with

Traits	Days to (days)	heading	Days to (days)	maturity	Plant he (cm)	eight	Spikes plant	5/	Gains, spike	1	1000-grain	weight (g)	Grain yie /plant(g)	
Generations	F <sub>2</sub>	F <sub>3</sub>	F <sub>2</sub>	F₃	F <sub>2</sub>	F <sub>3</sub>	F <sub>2</sub>	F₃	F <sub>2</sub>	F₃	F <sub>2</sub>	F₃	F <sub>2</sub>	F₃
Families	Optima	l irrigation												
1	90	90.8	147	153.2	135	126.4	15	14.8	60	71.8	51.7	47.25	38.1	34.17
3	89	91.6	145	150.2	143	144.9	25	15.6	84	78.5	52.4	44.96	80.5	29.84
4	88	91.3	147	155.1	101	119.6	15	14.3	86	67.1	43	46.49	43.7	32.09
5	90	92.1	145	150.5	111	130.6	6	15.8	108	80.8	47.2	51.79	36.1	28.11
6	91	91.1	146	154.1	141	143	11	15.8	73	63.6	57.5	48.03	30.5	27.27
10	89	91.8	145	157	133	148.4	16	16.8	63	68.5	57.1	52.91	42.6	25.8
17	89	91.8	145	154.9	110	145.5	20	15.3	63	74.5	44.4	46.14	44.1	28.94
19	91	90.7	145	153.2	129	146.5	23	17.6	57	70.8	43.9	48.32	46	34.08
24	89	91.4	143	156.5	110	121.7	15	16	62	66.3	53.2	44.94	38.5	27.23
30	88	92.3	145	150.5	121	137.7	17	20.2	86	66.5	45.3	45.82	49.2	37.4
Mean	89.4	91.5	145.3	153.5	123.4	136.4	16.3	16.2	74.2	70.8	49.57	47.67	44.93	30.49
SE	0.32	0.16	0.35	0.74	4.44	3.28	1.66	0.5	4.91	1.68	1.65	0.82	4.08	1.13
Sids 12 (P <sub>1</sub> )	91.8		155.4		120.4		14		86.6		57.7		29.2	
Line 44 (P <sub>2</sub> )	92.5		156.8		110.2		18		81		55		31	
Families	Water s	stress												
5	88	90.8	131	144	137	126.5	10	14.5	74	55.5	52.7	47.19	29.3	25.76
6	85	91.6	133	142.9	110	126.3	14	15.7	73	55.3	50.7	46.05	32.6	29.93
8	87	91.3	134	142.9	135	133	18	18.3	74	55.4	50	47.85	37	30.6
9	85	92.1	130	142.9	145	114.7	18	16.7	87	61.7	56.3	55.34	28.6	27.23
15	85	91.1	131	143.3	135	122.5	19	17.9	45	65.1	66.7	48.82	37.1	27.76
16	87	91.8	133	143.2	105	136	13	15.4	30	54.9	53.3	52.62	33.8	23.29
18	85	91.8	141	142.9	95	128.3	17	15.4	78	51.1	51.3	48.74	53.3	28.55
28	94	90.7	135	142.6	102	137.4	18	14.8	54	53.3	57.4	46.73	38.2	28.14
34	93	91.4	136	145.8	115	118.5	14	15.3	56	65.5	57.1	49.15	40.3	29.77
37	92	92.3	136	143.2	110	146.2	11	15.8	86	62.1	51.2	47.96	31.5	30.15
Mean	88.1	91.5	134	143.4	118.9	128.9	15.2	16	65.7	58	54.67	49.04	36.17	28.12
SE	1.07	0.16	0.97	0.28	5.24	2.83	0.97	0.38	5.61	1.54	1.51	0.85	2.15	0.68
Sids 12 (P <sub>1</sub> )	87.4		145.1		117.9		12.7		59.1		52.1		25.7	
Line 44 (P <sub>2</sub> )	85.8		135		107.2		14.6		72.7		47.1		28.1	

**Table 4.** The average earliness, plant height, and grain yield components in the  $F_2$  (selected) and  $F_3$  generations under optimal irrigation and water stress conditions for the cross I (Sids 12 × Line 44).

Traits	Days to (days)	heading	Days to (days)	maturity	Plant he	eight	Spikes plant	5/	Grains spike	5/	1000-g weight		Grain yield	/plant (g)
Generations	(uays) F2	F <sub>3</sub>	$F_2$	F <sub>3</sub>	$F_2$	F <sub>3</sub>	F <sub>2</sub>	F3	F <sub>2</sub>	F3	F <sub>2</sub>	<u>(g)</u> F₃	F <sub>2</sub>	F <sub>3</sub>
Families		l irrigation	12	13	12	13	12	13	12	13	12	13	12	13
1	86.0	87.5	147.0	156.3	130.0	122.8	10.0	14.6	71.0	71.2	56.3	70.27	44.90	27.73
2	87.0	89.4	146.0	157.7	145.0	124.2	17.0	17.6	68.0	68.1	51.5	64.24	40.50	26.94
3	85.0	88.6	146.0	158.1	125.0	122.5	16.0	12.5	73.0	66.8	50.7	45.40	37.80	28.30
4	88.0	90.0	144.0	159.7	131.0	130.8	11.0	15.1	69.0	69.8	50.7	58.68	54.10	30.06
13	85.0	91.5	147.0	158.2	135.0	137.6	22.0	16.1	69.0	64.5	47.8	54.97	62.80	30.96
19	94.0	90.8	146.0	158.7	129.0	120.0	17.0	15.0	64.0	68.8	48.4	56.12	37.80	29.05
20	95.0	87.7	145.0	155.2	125.0	117.4	20.0	11.6	70.0	75.2	40.0	50.08	76.60	29.01
26	93.0	87.9	146.0	158.0	140.0	135.8	17.0	11.9	55.0	64.2	69.1	64.05	46.00	27.05
27	92.0	88.2	146.0	158.8	110.0	122.7	16.0	14.4	92.0	74.7	43.5	50.18	42.70	31.57
28	95.0	87.9	147.0	156.6	127.0	121.7	22.0	14.4	69.0	63.2	46.4	49.30	48.00	29.82
Mean	90.0	88.9	146.0	157.7	129.7	125.6	16.8	14.3	70.0	68.6	50.44	56.33	49.12	29.05
SE	1.3	0.4	0.3	0.4	2.9	2.0	1.2	0.6	2.8	1.3	2.38	2.39	3.70	0.47
Line 20 (P <sub>1</sub> )	89.0		157.6		117.0		15.8		61.4		62.6		29.5	
Sakha 93 (P <sub>2</sub> )	87.2		150.6		110.2		16.2		63.6		61.4		28.1	
Families	Water s	stress												
1	82.0	87.3	140.0	150.2	115.0	119.7	18.0	17.2	79.0	50.8	33.7	54.03	32.80	26.37
5	83.0	87.0	139.0	148.2	125.0	126.2	18.0	17.5	66.0	48.6	31.8	53.96	30.00	23.53
9	82.0	86.0	139.0	150.3	120.0	127.0	17.0	18.9	50.0	56.7	42.0	48.85	22.50	21.82
11	81.0	88.1	141.0	148.0	125.0	110.6	17.0	16.0	59.0	56.9	28.8	50.60	29.20	22.76
17	83.0	85.7	140.0	147.9	115.0	120.7	15.0	16.9	66.0	55.3	46.3	53.95	32.90	24.51
19	86.0	85.2	139.0	146.8	125.0	128.5	13.0	17.0	64.0	48.5	40.6	51.92	31.50	18.34
28	83.0	87.7	140.0	148.4	115.0	121.1	18.0	15.8	55.0	56.8	45.5	50.65	24.20	21.47
33	83.0	85.4	140.0	146.7	130.0	121.3	17.0	16.9	68.0	50.4	36.8	53.01	29.60	22.40
34	84.0	85.3	138.0	148.3	125.0	127.7	15.0	15.2	77.0	48.8	23.4	50.61	26.40	21.22
37	83.0	88.4	142.0	148.0	120.0	125.0	19.0	18.9	67.0	47.8	26.3	50.26	32.70	23.24
Mean	83.0	86.6	139.8	148.3	121.5	122.8	16.7	17.0	65.1	52.1	35.52	51.78	29.18	22.56
SE	0.4	0.4	0.3	0.4	1.6	1.6	0.6	0.4	2.7	1.2	2.41	0.56	1.11	0.64
Line 20 (P <sub>1</sub> )	86.0		147.6		101.0		15.1		58.6		54.4		23.5	
Sakha 93 (P <sub>2</sub> )	82.8		147.3		109.8		14.7		51.1		57.8		21.7	

**Table 5.** The average of earliness, plant height, and grain yield components in the  $F_2$  (selected) and  $F_3$  generations under optimal irrigation and water stress conditions for the cross II (Line 20 × Sakha 93).

an average of 11.9 spikes. Grains per spike varied between 51.0 and 108.0 with an average of 71.3 grains. Thousand-grain yield ranged from 38.8 to 116.5 g with an average of 53.7 g and grain yield per plant varied from 12.8 to 60.5 g with an average yield of 31.2 g (Table 6).

On the other side, under water stress, the traits ranged as follows: from 84.0 to 94.0 with an average of 88.7 days for days to heading; days to maturity - from 130.0 to 142.0 with an average of 134.8 days; plant height - from 99.0 to 140.0 cm with an average of 117.0 cm; spikes per plant - from 3.0 to 19.0 with an average of 10.6; grains per spike - from 30.0 to 87.0 with an average of 61.6 grains; 1000-grain weight - from 35.0 to 66.7 g with an average of 52.9 g, and grain yield per plant from 5.2 to 45.3 g with an average of 23.2 g.

The values of genotypic coefficients of variability (GCV) in F2 under optimal irrigation and water stress were 1.01% and 2.51% for days to heading, 0.63% and 1.49% for days to maturity, 8.25% and 7.19% for plant height, 40.65% and 34.55% for spikes per plant, 15.91% and 19.82% for grains per spike, 21.85% and 9.34% for 1000-grain weight, and 23.88% and 17.70% for grain yield per plant, respectively. As for the values of phenotypic coefficient of variability (PCV), these were 1.28% and 3.25% for days to heading, 0.71% and 1.91% for days to maturity, 9.51% and 8.58% for plant height, 45.32% and 38.88% for spikes per plant, 20.53% and 22.25% for grains per spike, 26.35% and 13.74% for 1000-grain weight, and 34.93% and 36.79% for grain yield per plant under optimal irrigation and water stress, respectively.

Estimates of broad-sense heritability  $(h_{Bs}^2)$  in F<sub>2</sub> were 62.14% and 59.87% for days to heading, 78.15% and 60.78% for days to maturity, 75.27% and 70.39% for plant height, 80.45% and 79.0% for spikes per plant, 60.07% and 79.35% for grains per spike, 68.73% and 46.22% for 1000-grain weight, and 46.73% and 23.15% for grain yield per plant for optimal irrigation and water stress conditions, respectively.

The magnitudes of the genetic advance (GA) in  $F_2$  were 0.80% and 2.19% for days to heading, 0.90% and 1.99% for days to maturity, 9.74% and 8.97% for plant height, 4.89% and 4.14% for spikes per plant, 9.85% and 13.82% for grains per spike, 10.90% and 4.27% for 1000-grain weight, and 5.71% and 2.51% for grain yield per plant under optimal irrigation and water stress, respectively.

### *F*<sub>3</sub> Generation

Results for  $F_3$  generation in Table 6 showed the following data under optimal irrigation. Days to heading varied from 90.7 to 92.3 with an average of 91.5 days. Days to maturity ranged from 150.2 to 157.0 with an average of 153.5 days. Plant height ranged between 119.6 and 148.5 cm with an average of 136.4 cm. Spikes per plant diverse from 14.3 to 20.2 with an average of 16.2 spikes. Grains per spike varied between 63.6 and 80.8 with an average of 70.8 grains. Thousand-grain weight ranged from 44.9 to 52.9 with an average of 47.7 g. Grain yield per plant varied from 25.8 to 37.4 with an average of 30.5 g.

With water stress condition, the studied traits varied as follows: Days to heading from 84.6 to 87.2 with an average of 85.9 days; Days to maturity varied from 142.6 to 145.8 with an average of 143.4 days; Plant height ranged from 114.7 to 146.2 cm with an average of 128.9 cm; Spikes per plant varied from 14.5 to 18.3 with an average of 15.9 spikes; Grains per spike varied from 51.1 to 65.5 with an average of 57.9 grains; 1000-grain weight varied from 46.1 to 55.3 g with an average of 49.0 g; and Grain yield per plant varied from 23.3 to 30.6 g with an average of 28.1 g.

The GCV values in F<sub>3</sub> under both optimal irrigation and water stress for all the traits were 0.53% and 1.02% for days to heading, 1.53% and 0.59% for days to maturity, 7.64% and 6.70% for plant height, 9.82% and 7.05% for spikes per plant, 7.18% and 7.81% for grains per spike, 5.34% and 5.25% for 1000-grain weight, and 10.77% and 6.70% for grain yield per plant, respectively. The PCV values, on the other hand, were 0.63% and 1.30% for days to heading, 1.70% and 0.71% for days to maturity, 8.39% and 7.90% for plant height, 10.88% and 8.77 % for spikes per plant, 8.60% and 9.77% for grains per spike, 6.10% and 6.31% for 1000grain weight, and 13.80% and 9.23% for grain yield per plant, for optimal irrigation and water stress, respectively.

Estimates of broad-sense heritability  $(h_{Bs}^2)$  in F<sub>3</sub> under optimal irrigation and water stress treatments were 69.85% and 61.68% for days to heading, 81.05% and 69.38% for days to maturity, 82.92% and 71.83% for plant height, 81.47% and 64.73% for spikes per plant, 69.63% and 63.83% for grains per spike, 76.76% and 69.20% for 1000-grain weight and 60.93% and 57.77% for grain yield per plant, respectively.

Environments			Optir	nal irriga	tion					Water s	stress			
Traits	DH	DM	PH	NS/P	NG/S	TGW	GY/P	DH	DM	PH	NS/P	NG/S	TGW	GY/P
F <sub>2</sub>	Genetic	paramete	rs											
Range	(88.0	(143.0	(90.0 -	(2.0 -	(51.0	(38.8	(12.8	(84.0	(130.0	(99.0 -	(3.0 -	(30.0	(35.0	(5.2 -
	-	-			-	-	-	-	-			-	-	
	92.0)	147.0)	143.0)	25.0)	108.0)	116.5)	60.5)	94.0)	142.0)	140.0)	19.0)	87.0)	66.7)	45.3)
Mean	89.7	145	121.5	11.9	71.3	53.7	31.2	88.7	134.8	117	10.6	61.6	52.9	23.20
VE	0.5	0.23	33.03	5.75	85.6	62.69	63.44	3.33	2.61	29.81	3.58	38.83	28.42	56.20
VG	0.82	0.83	100.53	23.66	128.7	137.7	55.65	4.97	4.04	70.86	13.48	149.2	24.42	16.93
					7	7						5		
G.C.V.%	1.01	0.63	8.25	40.65	15.91	21.85	23.88	2.51	1.49	7.19	34.55	19.82	9.34	17.70
P.C.V.%	1.28	0.71	9.51	45.32	20.53	26.35	34.93	3.25	1.91	8.58	38.88	22.25	13.74	36.79
h <sup>2</sup> <sub>Bs</sub>	62.14	78.15	75.27	80.45	60.07	68.73	46.73	59.87	60.78	70.39	79	79.35	46.22	23.15
GA	1.47	1.66	17.92	8.99	18.12	20.05	10.5	3.55	3.23	14.55	6.72	22.42	6.92	4.08
Realized selection	-0.3	0.27	1.87	4.33	2.87	-4.16	11.69	-0.63	-1.35	2.4	4.58	4.05	1.77	12.13
differential														
F <sub>3</sub>	I. Gene	tic parame	ters											
Range	(90.7-	(150.2-	(119.6	(14.3	(63.6-	(44.9-	(25.8-	(84.6-	(142.6-	(114.7	(14.5	(51.1-	(42.1-	(23.3
-	•		-	-	•	•	•		•	_	-	•	•	-
	92.3)	157.0)	148.5)	20.2)	80.8)	52.9)	37.4)	87.2)	145.8)	146.2)	18.3)	65.5)	55.3)	30.6)
Mean	91.50	153.50	136.40	16.20	70.80	47.70	30.50	85.90	143.40	128.90	15.90	57.90	46.00	28.10
VE	0.10	1.28	22.38	0.58	11.28	1.96	6.92	0.48	0.32	29.24	0.69	11.62	2.95	3.18
VG	0.23	5.49	108.66	2.54	25.86	6.48	10.79	0.77	0.72	74.54	1.27	20.50	6.62	3.55
VP	0.33	6.77	131.04	3.11	37.14	8.44	17.70	1.25	1.04	103.77	1.96	32.12	9.57	6.73
V F3	0.56	12.27	239.70	5.65	63.01	14.93	28.49	2.02	1.75	178.31	3.23	52.62	16.19	10.29
G.C.V.%	0.53	1.53	7.64	9.82	7.18	5.34	10.77	1.02	0.59	6.70	7.05	7.81	5.25	6.70
P.C.V.%	0.63	1.70	8.39	10.88	8.60	6.10	13.80	1.30	0.71	7.90	8.77	9.77	6.31	9.23
h² <sub>Bs</sub>	69.85	81.05	82.92	81.47	69.63	76.76	60.93	61.68	69.38	71.83	64.73	63.83	69.20	52.77
Realized selection	-0.22	-0.52	-3.88	0.51	-1.79	-0.69	3.94	0.87	0.32	2.06	0.30	1.59	-1.29	2.00
differential														
	II. Sele	ction parar	neters											
Realized response in F3 (rR)	1.64	8.62	17.67	6.42	1.94	-8.14	6.08	-2.98	8.23	13.17	6.88	-2.31	-3.25	9.52
Percentage rR of F3 (rR %)	1.79	5.61	12.95	39.58	2.74	-17.09	19.95	-3.46	5.74	10.22	43.04	-3.98	-6.63	33.85
GA	0.51	2.75	12.45	1.88	5.36	2.88	3.16	0.85	0.89	9.30	1.13	4.49	2.70	1.64
Predicted selection	0.84	3.92	17.34	2.66	8.89	4.33	5.98	1.80	1.68	16.96	2.28	9.21	5.11	4.07
differential			-								-			-
Predicted generalized in F4	2.19	2.46	1.14	2.70	0.24	-2.11	1.14	-2.10	6.22	0.99	3.83	-0.32	-0.81	2.97
R	-0.60	-0.11	0.59	0.17	0.43	0.25	0.18	-0.23	0.05	-0.52	0.60	-0.05	0.17	0.24
В	-0.29	-0.05	0.79	0.56	1.24	0.50	0.65	-0.03	0.16	-0.96	1.53	-0.16	0.31	0.74
R	-0.29 35.60	1.22	0.79 34.54	2.91	1.24	6.32	3.29	-0.03 5.48	0.18	26.82	36.22	-0.16 0.20	3.06	5.56

**Table 6.** Genetic and selection parameters for the cross I (Sids  $12 \times \text{Line 44}$ ) of studied traits in the F<sub>2</sub> and F<sub>3</sub> generations under optimal irrigation and water stress conditions.

DH: days to heading, DM: days to maturity, PH: plant height, S/P: spikes per plant, G/S: grains per spike, TGW: 1000-grain weight, and GY/P: grain yield per plant. GA: genetic advance,  $h_{BS}^2$ : heritability in the broad sense, r: correlation between offspring (F<sub>3</sub>) and parents (F<sub>2</sub>), b: the regression of offspring (F<sub>3</sub>) on parents (F<sub>2</sub>), and R: coefficient of the determinant of selection response.

The realized (actual) response to selection in  $F_3$  (rR) values were 1.64 and -2.98 days to heading, 8.62 and 8.23 days to maturity, 17.67 and 13.17 cm for plant height, 6.42 and 6.88 spikes per plant, 1.94 and -2.31 grain for grains per spike, -8.14 and -3.25 g for 1000-grain weight, and 6.08 and 9.52 g for grain yield per plant under optimal irrigation and water stress, respectively.

On the other hand, the predicted selection differential in F<sub>3</sub> values were 0.84 and 1.80 days to heading, 3.92 and 1.68 days to maturity, 17.34 and 16.96 cm for plant height, 2.66 and 2.28 spikes per plant, 8.89 and 9.21 grain for grains per spike, 4.33 and 5.11 g for 1000-grain weight, and 5.98 and 4.07 g for grain yield per plant under optimal irrigation and water stress, respectively. Meanwhile, the magnitudes of the GA in  $\ensuremath{\mathsf{F}}_3$  under optimal irrigation and water stress conditions were 0.51% and 0.85% for days to heading, 2.75% and 0.89% for days to maturity, 12.45% and 9.30% for plant height, 1.88% and 1.13% for spikes per plant, 5.36% and 4.49% for grains per spike, 2.88% and 2.70% for 1000-grain weight, and 3.16% and 1.64% for grain yield per plant, respectively (Table 6).

Moreover, the predicted generalized in  $F_4$  was 2.19 and -2.10 days to heading, 2.46 and 6.22 days to maturity, 1.14 and 0.99 cm for plant height, 2.70 and 3.83 spikes per plant, 0.24 and -0.32 grain for grains per spike, -2.11 and -0.81 g for 1000-grain weight, and 1.14 and 2.97 g for grain yield per plant under optimal irrigation and water stress, respectively.

Correlation (r) between offspring  $(F_3)$ and parents  $(F_2)$  showed positive values for plant height (0.59), spikes per plant (0.17), grains per spike (0.43), 1000-grain weight (0.25), and grain yield per plant (0.18) under optimal irrigation. Likewise, under water stress condition, positive values were noted for days to maturity (0.05), spikes per plant (0.60), 1000-grain weight (0.17), and grain yield per plant (0.24). Conversely, negative correlation values were recorded for days to heading (-0.6) and days to maturity (-0.11) under optimal irrigation, while under water stress, these were also for days to heading (-0.23), as well as, for plant height (-0.52), and grains per spike (-0.05).

Regression (b) of offspring ( $F_3$ ) on parents ( $F_2$ ) estimates were positive for plant height (0.79), spikes per plant (0.56), grains per spike (1.24), 1000-grain weight (0.5), and grain yield per plant (0.65) under optimal irrigation, as well as, for days to maturity (0.16), spikes per plant (1.53), 1000-grain weight (0.31), and grain yield per plant (0.74) under water stress. The negative regression values recorded under optimal irrigation were for days to heading (-0.29) and days to maturity (-0.05), while days to heading (-0.03), plant height (-0.96), and grains per spike (-0.16) were noted for the water stress conditon.

### Populations of cross II (Line 20 × Sakha 93)

Genetic and selection parameters for the cross II (Line 20 × Sakha 93) on days to heading, days to maturity, plant height, spikes per plant, grains per spike, 1000-grain weight, and grain yield per plant in the  $F_2$  and  $F_3$  generations under optimal irrigation and water stress conditions are shown in Table 7.

# F<sub>2</sub> Generation

Under optimal irrigation, the following were the results. Days to heading varied from 85.0 to 95.0 with an average of 89.6 days. Days to maturity ranged from 143.0 to 149.0 with an average of 146.2 days. Plant height ranged between 108.0 and 145.0 cm with an average of 128.1 cm. Spikes per plant diverse from 9.0 to 22.0 with an average of 14.5 spikes. Grains per spike varied between 44.0 and 92.0 with an average of 66.1 grains. The 1000-grain weight ranged from 24.2 to 69.1 with an average of 47.0 g. Lastly, grain yield per plant varied from 10.6 to 62.8 with an average of 34.5 g.

Under water stress, the following were the results. Days to heading ranged from 81.0 to 87.0 with an average of 84.4 days. Days to maturity varied from 136.0 to 144.0 with an average of 139.6 days. Plant height ranged from 105.0 to 135.0 cm with an average of 118.8 cm. Spikes per plant varied from 5.0 to 19.0 with an average of 12.3 spikes. Grains per spike ranged between 50.0 and 79.0 with an average of 65.0 grain. The 1000-grain weight varied from 17.1 to 47.8 g with an average of 33.6 g, and grain yield per plant varied from 10.1 to 37.2 g with an average of 21.3 g.

The values of GCV in  $F_2$  of cross II were 3.37% and 1.55% for days to heading, 0.72% and 0.80% for days to maturity, 6.61% and 4.32% for plant height, 17.53% and 24.93% for spikes per plant, 10.08% and 6.36% for grains per spike, 17.36% and 18.47% for 1000-grain weight, and 24.17% and 19.23% for grain yield per plant under optimal irrigation and water stress,

$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	Environments			Opti	mal irriga	ation						Water stre	ess		
Range (85.0- (143.0- (108.0- (9.0- (44.0- (24.2 (10.6- (81.0 (136.0 (105.0- (5.0 (17.1- (10.1-   Mean 95.0) 149.0) 145.0) 22.0) 92.0) 62.8) 87.0) 144.0) 135.0) 19.0) 79.0) 47.80) 37.20)   WE 3.35 0.55 20.85 7.7 39.55 32.88 78.74 0.64 0.81 20.64 5.06 33.74 21.03 0.47   VG 9.11 1.11 71.56 6.49 44.43 66.6 69.57 1.71 1.24 26.32 9.56 33.64 1.82 1.02 5.77 30.85 10.96 22.95 32.24   P_G.V.% 3.31 1.78 15.33 3.55 9.99 13.76 11.77 2.29 1.79 7.91 5.14 4.94 10.3 5.04   GA 0.4 -0.17 1.63 2.77 3.67 1.76 <	Traits	DH	DM	PH	NS/P	NG/S	TGW	GY/P	DH	DM	PH	NS/P	NG/S	TGW	GY/P
Range (85.0- (143.0- (108.0- (9.0- (44.0- (24.2 (10.6- (81.0 (136.0 (105.0- (5.0 (17.1- (10.1-   Mean 95.0) 149.0) 145.0) 22.0) 92.0) 62.8) 87.0) 144.0) 135.0) 19.0) 79.0) 47.80) 37.20)   WE 3.35 0.55 20.85 7.7 39.55 32.88 78.74 0.64 0.81 20.64 5.06 33.74 21.03 0.47   VG 9.11 1.11 71.56 6.49 44.43 66.6 69.57 1.71 1.24 26.32 9.56 33.64 1.82 1.02 5.77 30.85 10.96 22.95 32.24   P_G.V.% 3.31 1.78 15.33 3.55 9.99 13.76 11.77 2.29 1.79 7.91 5.14 4.94 10.3 5.04   GA 0.4 -0.17 1.63 2.77 3.67 1.76 <	F <sub>2</sub>	Genetic	c paramete	ers											
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$		(85.0-	(143.0-	(108.0-	(9.0 -	(44.0-	(24.2	(10.6 -	(81.0	(136.0	(105.0	(5.0 -	(50.0-	(17.1 -	(10.1 -
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	-						-		-	-	-				
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$ \begin{array}{cccccccccccccccccccccccccccccccccccc$			0.55										33.74		
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	VG														
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	G.C.V.%	3.37	0.72		17.53	10.08	17.36		1.55	0.8	4.32	24.93	6.36		
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	P.C.V.%	3.94	0.88	7.51	25.92	13.86	21.22	35.29	1.82	1.02	5.77	30.85	10.96	22.95	32.24
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	h <sup>2</sup> <sub>Bs</sub>	73.1	66.89		45.73	52.91	66.95	46.91	72.62	60.63	56.04	65.3	33.64		35.58
$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	GA	5.31	1.78	15.33	3.55	9.99	13.76	11.77	2.29	1.79	7.91	5.14	4.94	10.3	5.04
Range(87.5-(155.2-(117.4-(11.6-(63.2-(45.4-(26.94)(85.2(146.7(11.0.6(10.20)(47.84)(48.85)(18.34)91.45159.70137.6017.6075.2070.27)31.57)88.4)150.3)128.5)18.9)56.9)54.03)26.37)Mean88.95157.73125.5514.3268.6456.3329.0586.61148.28122.7813.0352.0551.7822.56VE0.30.418.340.264.5612.491.070.270.4213.820.515.881.322.28VG1.81.5942.133.3915.1857.051.961.381.1921.811.2112.182.783.43VP2.1250.473.6519.7369.543.021.651.6135.641.7218.064.15.71VF33.93.5892.67.0334.91126.54.983.022.857.452.9330.246.889.14G.C.V.%1.630.95.6613.346.4714.85.981.480.864.867.718.163.9110.59h^8_{B6}85.8879.583.4792.8476.9182.0564.7483.7873.6261.2170.2767.4467.8559.99Realized selection differential1.040.220.91.060.28-0.73	Realized selection differential	0.4	-0.17	1.63	2.27	3.87	3.44	13.03	-1.38	0.2	2.68	4.33	0.05	1.88	7.85
Mean91.45159.70137.6017.6075.2070.27)31.57)88.4)150.3)128.5)18.9)56.9)54.03)26.37)Mean88.95157.73125.5514.3268.6456.3329.0586.61148.28122.7813.0352.0551.7822.56VE0.30.418.340.264.5612.491.070.270.4213.820.515.881.322.28VG1.81.5942.133.3915.1857.051.961.381.1921.811.2112.182.783.43VP2.1250.473.6519.7369.543.021.651.6135.641.7218.064.15.71V F33.93.5892.67.0334.91126.54.983.022.857.452.9330.246.889.14G.C.V.%1.610.85.1712.855.6813.694.821.360.733.86.466.73.228.2P.C.V.%1.630.95.6613.346.4714.85.981.480.864.867.718.163.9110.59h <sup>2</sup> BSRealized selection differential1.040.220.91.060.28-0.730.720.490.290.120.6-1.061.271.85II. Selection parametersII. Selection parameters<	F <sub>3</sub>						]	[. Genetic	paramet	ters					
Mean 88.95 157.73 125.55 14.32 68.64 56.33 29.05 86.61 148.28 122.78 13.03 52.05 51.78 22.56   VE 0.3 0.41 8.34 0.26 4.56 12.49 1.07 0.27 0.42 13.82 0.51 5.88 1.32 2.26   VE 0.3 0.41 8.34 0.26 4.56 12.49 1.07 0.27 0.42 13.82 0.51 5.88 1.32 2.26   VG 1.8 1.59 42.13 3.39 15.18 57.05 1.96 1.38 1.19 21.81 1.21 12.18 2.78 3.43   VF3 3.9 3.58 92.6 7.03 34.91 126.5 4.98 3.02 2.8 57.45 2.93 30.24 6.88 9.14   G.C.V.% 1.63 0.9 5.66 13.36 6.47 14.8 5.98 1.48 0.86 4.86 7.71	Range	(87.5-	(155.2-	(117.4-	(11.6-	(63.2-	(45.4-	(26.94	(85.2	(146.7	(110.6	(10.20	(47.84	(48.85	(18.34
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V F3 $3.9$ $3.58$ $92.6$ $7.03$ $34.91$ $126.5$ $4.98$ $3.02$ $2.8$ $57.45$ $2.93$ $30.24$ $6.88$ $9.14$ G.C.V.% $1.51$ $0.8$ $5.17$ $12.85$ $5.68$ $13.69$ $4.82$ $1.36$ $0.73$ $3.8$ $6.46$ $6.7$ $3.22$ $8.2$ P.C.V.% $1.63$ $0.9$ $5.66$ $13.34$ $6.47$ $14.8$ $5.98$ $1.48$ $0.86$ $4.86$ $7.71$ $8.16$ $3.91$ $10.59$ $h^2_{Bs}$ $85.88$ $79.5$ $83.47$ $92.84$ $76.91$ $82.05$ $64.74$ $83.78$ $73.62$ $61.21$ $70.27$ $67.44$ $67.85$ $59.99$ Realized selection differential $1.04$ $0.22$ $0.9$ $1.06$ $0.28$ $-0.73$ $0.72$ $0.49$ $0.29$ $0.12$ $0.6$ $-1.06$ $1.27$ $1.85$ Percentage rR of F3 (rR) $-0.45$ $11.48$ $-1.7$ $0.92$ $4.44$ $11.05$ $1.09$ $1.78$ $8.74$ $4.85$ $6.1$ $-12.98$ $18.77$ $3.85$ Percentage rR of F3 (rR %) $-0.51$ $7.28$ $-1.35$ $6.42$ $6.47$ $19.61$ $3.74$ $2.05$ $5.9$ $3.95$ $35.8$ $-24.94$ $36.24$ $17.05$ GA $1.64$ $1.46$ $7.79$ $2.39$ $4.41$ $8.95$ $1.4$ $1.41$ $1.19$ $4.5$ $1.17$ $3.6$ $1.73$ $1.76$ Predicted selection $2.21$ $2.12$ $10.78$ <t< td=""><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></t<>															
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	VP			50.47	3.65	19.73	69.54	3.02	1.65	1.61	35.64	1.72	18.06		
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	V F3	3.9	3.58	92.6	7.03	34.91		4.98	3.02	2.8	57.45	2.93	30.24	6.88	9.14
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$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	G.C.V.%														
Realized selection differential 1.04 0.22 0.9 1.06 0.28 -0.73 0.72 0.49 0.29 0.12 0.6 -1.06 1.27 1.85   III. Selection parameters   Realized response in F3 (rR) -0.45 11.48 -1.7 0.92 4.44 11.05 1.09 1.78 8.74 4.85 6.1 -12.98 18.77 3.85   Percentage rR of F3 (rR %) -0.51 7.28 -1.35 6.42 6.47 19.61 3.74 2.05 5.9 3.95 35.8 -24.94 36.24 17.05   GA 1.64 1.46 7.79 2.39 4.41 8.95 1.4 1.41 1.19 4.5 1.17 3.6 1.73 1.76   Predicted selection 2.21 2.12 10.78 2.97 6.62 12.6 2.5 2.21 2.18 6.98 3.33 3.84	P.C.V.%			5.66	13.34	6.47	14.8	5.98	1.48	0.86	4.86	7.71	8.16		10.59
II. Selection parameters   Realized response in F3 (rR) -0.45 11.48 -1.7 0.92 4.44 11.05 1.09 1.78 8.74 4.85 6.1 -12.98 18.77 3.85   Percentage rR of F3 (rR %) -0.51 7.28 -1.35 6.42 6.47 19.61 3.74 2.05 5.9 3.95 35.8 -24.94 36.24 17.05   GA 1.64 1.46 7.79 2.39 4.41 8.95 1.4 1.41 1.19 4.5 1.17 3.6 1.73 1.76   Predicted selection 2.21 2.12 10.78 2.97 6.62 12.6 2.5 2.21 2.18 6.98 3.33 3.84	h <sup>2</sup> <sub>Bs</sub>	85.88	79.5	83.47	92.84	76.91	82.05	64.74	83.78	73.62	61.21	70.27	67.44	67.85	59.99
Realized response in F3 (rR) Percentage rR of F3 (rR %)-0.4511.48-1.70.924.4411.051.091.788.744.856.1-12.9818.773.85GA Predicted selection1.641.467.792.394.418.951.41.411.194.51.173.61.731.76QA Predicted selection2.212.1210.782.976.6212.62.52.212.129.632.186.983.333.84	Realized selection differential	1.04	0.22	0.9	1.06	0.28	-0.73	0.72	0.49	0.29	0.12	0.6	-1.06	1.27	1.85
Percentage rR of F3 (rR %)-0.517.28-1.356.426.4719.613.742.055.93.9535.8-24.9436.2417.05GA1.641.467.792.394.418.951.41.411.194.51.173.61.731.76Predicted selection2.212.1210.782.976.6212.62.52.212.129.632.186.983.333.84								. Selectio	n param	eters					
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GA 1.64 1.46 7.79 2.39 4.41 8.95 1.4 1.41 1.19 4.5 1.17 3.6 1.73 1.76   Predicted selection 2.21 2.12 10.78 2.97 6.62 12.6 2.5 2.21 2.12 9.63 2.18 6.98 3.33 3.84	Percentage rR of F3 (rR %)	-0.51	7.28	-1.35	6.42	6.47	19.61	3.74	2.05	5.9	3.95	35.8	-24.94	36.24	17.05
		1.64	1.46	7.79	2.39	4.41	8.95	1.4	1.41	1.19	4.5	1.17	3.6	1.73	1.76
		2.21	2.12	10.78	2.97	6.62	12.6	2.5	2.21	2.12	9.63	2.18	6.98	3.33	3.84
	differential														
Predicted generalized in F4 -0.23 6.07 -0.18 0.35 0.75 0.98 0.49 1.02 5.23 0.64 3.56 -2.36 7.15 1.27		-0.23	6.07	-0.18	0.35	0.75	0.98	0.49	1.02	5.23	0.64	3.56	-2.36	7.15	1.27
R -0.33 -0.31 0.48 -0.06 0.59 0.65* 0.29 -0.58 -0.11 0.11 0.36 -0.68* 0.14 0.4															
B -0.11 -0.22 0.67 -0.12 1.3 0.65* 2.3 -0.54 -0.11 0.11 0.54 -1.57* 0.62 0.69															
R 10.62 9.5 22.87 0.31 34.92 42.79 8.67 33.97 1.25 1.27 13.01 46.81 2.07 16.07		-													

**Table 7.** Genetic and selection parameters for the cross II (Line  $20 \times \text{Sakha 93}$ ) of studied traits in the  $F_2$  and  $F_3$  generations under optimal irrigation and water stress conditions.

DH: days to heading, DM: days to maturity, PH: plant height, S/P: spikes per plant, G/S: grains per spike, TGW: 1000-grain weight, and GY/P: grain yield per plant. GA: genetic advance,  $h_{Bs}^2$ : heritability in the broad sense, r: correlation between offspring (F<sub>3</sub>) and parents (F<sub>2</sub>), b: regression of offspring (F<sub>3</sub>) on parents (F<sub>2</sub>), and R: coefficient of the determinant of selection response.

respectively. For the PCV values, results were 3.94% and 1.82% for days to heading, 0.88% and 1.02% for days to maturity, 7.51% and 5.77% for plant height, 25.92% and 30.85% for spikes per plant, 13.86% and 10.96% for grains per spike, 21.22% and 22.95% for 1000-grain weight, and 35.29% and 32.24% for grain yield per plant under optimal irrigation and water stress, respectively.

Estimates of  $h_{Bs}^2$  in F<sub>2</sub> were 73.10% and 72.62% for days to heading, 66.89% and 60.63% for days to maturity, 77.44% and 56.04% for plant height, 45.73% and 65.30% for spikes per plant, 52.91% and 33.64% for grains per spike, 66.95% and 64.73% for 1000-grain weight, and 46.91% and 35.58% for grain yield per plant under optimal irrigation and water stress, respectively. The GA estimates in  $F_2$  were 2.89% and 1.41% for days to heading, 0.97% and 1.10% for days to maturity, 8.34% and 4.88% for plant height, 1.93% and 3.17% for spikes per plant, 5.43% and 3.05% for grains per spike, 7.48% and 6.35% for 1000-grain weight, and 6.40% and 3.11% for grain yield per plant under optimal irrigation and water stress, respectively.

## F<sub>3</sub> Generation

Mean performance under optimal irrigation of  $F_3$  generation for cross Line 20 × Sakha 93 in Table 7 showed that days to heading varied from 87.5 to 91.5. Days to maturity ranged from 155.2 to 159.7. Plant height ranged between 117.4 and 137.6 cm. Spikes per plant ranged from 11.6 to 17.6. Grains per spike varied between 63.2 and 75.2. The 1000-grain weight diverse from 45.4 to 27.3. Grain yield per plant varied from 26.9 to 31.6 with an average of 29.0 g. Moreover, under water stress, these traits ranged between 85.2-88.4 (days to heading), 146.7-150.3 (days to maturity), 110.6-128.5 cm (plant height), 15.2-18.9 (spikes per plant), 47.8-56.9 (grains per spike), 48.8-54.0 g (1000-grain weight), and 18.3-26.4 g (grain yield per plant).

The magnitude of GCV values in  $F_3$  of cross II were 1.51%, 0.80%, 5.17%, 12.8%, 5.68%, 13.0%, and 4.82% under optimal irrigation, and 1.36%, 0.73%, 3.80%, 6.46%, 6.70%, 3.22%, and 8.20% under water stress, for days to heading, days to maturity, plant height, spikes per plant, grains per spike, 1000-grain weight, and grain yield per plant traits, respectively. The values of PCV were 1.63%, 0.90%, 5.66%, 13.34%, 6.47%, 14.80%, and 5.98% under optimal irrigation, and 1.48%, 0.86%, 4.86%, 7.71%, 8.16%,

3.91%, and 10.59% under water stress for enumerated traits, respectively.

Estimates of  $h_{Bs}^2$  in  $F_3$  were 85.88%, 79.50%, 83.47%, 92.84%, 76.91%, 82.05%, and 64.74% under optimal irrigation, and 83.78%, 73.62%, 61.21%, 70.27%, 67.44%, 67.85%, and 59.99% under water stress for days to heading, days to maturity, plant height, spikes per plant, grains per spike, 1000-grain weight and grain yield per plant traits, respectively.

## Selection parameters

The realized (actual) response to selection in F<sub>3</sub> (rR) values of cross II were -0.45 day, 11.48 day, -1.70 cm, 0.92 spike, 4.44 grains, 11.05 g, and 1.09 g under optimal irrigation, and 1.78 days, 8.74 days, 4.85 cm, 6.10 spikes, -12.98 grain, 18.77 g, and 3.85 g under water stress for days to heading, days to maturity, plant height, spikes per plant, grains per spike, 1000-grain weight, and grain yield per plant traits, respectively. On the other hand, the predicted selection differential in F<sub>3</sub> values were 2.21 days, 2.12 days, 10.78 cm, 2.97 spikes, 2.62 grains, 12.60 g, and 2.50 g under optimal irrigation, and 2.21 days, 2.12 days, 9.63 cm, 2.18 spikes, 6.98 grains, 3.33 g, and 3.84 g under water stress, for enumerated traits, respectively (Table 7).

Meanwhile, the GA values in F<sub>3</sub> of cross II were 1.64%, 1.46%, 7.79%, 2.39%, 4.41%, 8.95%, and 1.40% under optimal irrigation, and 1.41%, 1.19%, 4.50%, 1.17%, 3.60%, 1.73%, and 1.76% under water stress, for days to heading, days to maturity, plant height, spikes per plant, grains per spike, 1000-grain weight, and grain yield per plant traits, respectively (Table 7). Moreover, the predicted generalized in  $F_4$  was -0.23 and 1.02 days to heading, 6.07 and 5.23 days to maturity, -0.18 and 0.64 cm for plant height, 0.35 and 3.56 spikes per plant, 0.75 and -2.36 grain for grains per spike, 0.98 and 7.15 g for 1000-grain weight, and 0.49 and 1.27 g for grain yield per plant under optimal irrigation and water stress, respectively.

Positive correlation (r) values were recorded between offsprings ( $F_3$ ) and parents ( $F_2$ ) under optimal irrigation (0.48, 0.59, 0.65, and 0.29) and drought stress (0.11, 0.36, 0.14, and 0.40) for plant height, grains per spike, 1000-grain weight, and grain yield per plant, respectively. However, days to heading and days to maturity showed negative correlation values under optimal irrigation (-0.33 and -0.31) and water stress (-0.58 and -0.11), as well as, in spikes per plant (-0.06) under optimal irrigation and grains per spike (-0.68) under water stress condition.

Positive regression (b) estimates of offspring ( $F_3$ ) on parents ( $F_2$ ) were observed for plant height, grains per spike, 1000-grain weight, and grain yield per plant under optimal irrigation (0.67, 1.30, 0.65, and 2.30) and water stress (0.11, 0.54, 0.62, and 0.69). But for negative regression, values were recorded for days to heading and days to maturity under optimal irrigation at -0.11 and -0.22 and water stress at -0.54 and -0.11, as well as, for spikes per plant and grains per spike under optimal irrigation and water stress (-.12 and -1.57), respectively.

# DISCUSSION

All studied traits of  $F_2$  and  $F_3$  families had significant differences in the two crosses under two water regimes. These confirm considerable genetic variation between  $F_2$  plants and  $F_3$ families and the measured traits and possibility of selection for drought tolerance and optimal irrigation. Genetic variation was found in wheat for earliness traits and yield components (Mahdy *et al.*, 2015; Aziz *et al.*, 2018; Darwish *et al.*, 2018), where the efficiency of selection and heterosis are mainly dependent upon the magnitude of genetic variability existing in the wheat populations (Al-Naggar *et al.*, 2020).

Water shortage stress reduced the means of  $F_2$  and  $F_3$  generations for all studied traits for two wheat crosses compared with the optimal irrigation. Wheat traits of cross I (Sids12 × Line 44) were reduced in  $F_2$ generation by 1.11%, 7.03%, 3.7%, 10.92%, 13.6%, 1.49%, and 25.64%, as well as, in  $F_3$ by 6.12%, 6.58%, 5.5%, 1.85%, 18.22%, 3.56%, and 7.87% for DH, DM, PH, S/P, G/S, TGW, and GY/P, respectively. Likewise, traits of cross II (Line 20 × Sakha 93) were reduced in F<sub>2</sub> by 5.83%, 4.49%, 7.21%, 14.8%, 1.63%, 28.43%, and 38.19%, and in  $F_3$  by 2.63%, 5.99%, 2.21%, 9.01%, 24.17%, 8.08%, and 22.34% for DH, DM, PH, S/P, G/S, TGW, and GY/P, respectively. The different behaviors of genotypes under drought stress and full irrigation conditions reflect the effect of genetic influences and the impact of environments, similar to results observed by wheat scientists (Ahmed et al., 2019a). Furthermore, grain yield losses under water stress were associated with an insufficient supply of assimilates to support the wheat reproductive stages and grain growth. Aberkane et al. (2020) reported that the wheat grain yield was reduced by 62% under drought and found high variation for phenology and agronomic traits, with days to heading explaining 16% of GY under drought.

In addition, drought decreased wheat grain yield by 47% in Australia in 2006 (Rauf et al., 2016). Forward-looking, grain yield losses by 2050 are predicted to reach 10%-30% associated with the expected increase in drought and heat stresses (Kumar et al., 2013). Thus, for each 1 °C increase in earth temperature, the wheat grain yields worldwide are expected to reduce by 6% (Asseng et al., 2015). Therefore, breeding programs for drought tolerance in the arid and semi-arid area are very useful for developing elite wheat germplasm that will respond to the requirements of farmers and consumers and adapt to climate change effects. It is worthy to note that transgressive segregants were observed for GY/P and its related traits occurred in many selected recombination in F2 and F<sub>3</sub>; these recombinants exceeded the highest or lesser parent, suggesting that all selected parents had associated alleles with high values of these traits. Moreover, the highyielding segregants plants would be used as a germplasm source to improve wheat productivity under optimal irrigation and water stress conditions, as explained by Jan et al. (2015); Saleem et al. (2016); and Channa (2022). Breeding water for deficit environments is challenging because drought tolerance is a complex trait; its mechanism is generally environment-specific, and intense in interaction of genotype  $\times$  environment, which reduces the efficiency of selection (Mickelbart et al., 2015). Consequently, breeding for drought stress requires methodologies in plant science and the integration of multiple disciplines (Mwadzingeni et al., 2016).

Under optimal irrigation for the cross I in F<sub>2</sub> and F<sub>3</sub> generations, family three was early with high NS/P and GY/P, making it a potential family for earliness and high grain yield. In addition, family 19 had high NS/P, NG/S, TGW, and GY/P. As for the water stress condition, the family 16 was early with high TGW, making it a potential family for earliness and high grain-filling rate. Family 34 exhibited high NG/S, TGW, and GY/P and short PH, as well as, families six, eight, 18, and 37 showing high GY/P under drought. Family four was early with high TGW and GY/P for cross II under optimal irrigation, where family 27 had high NG/S, TGW, and GY/P, and family 13 showed high NS/P and GY/P. On the other side, under water stress for cross II, family 17 was early with high NG/S, TGW, and GY/P, while family one had high NS/P, TGW, and GY/P and short PH.

Crosses	Cross I (S	Sids 12 × Li	ne 44)		Cross II (Line 20 × Sakha 93)							
Generations		F <sub>2</sub>	F	3		F <sub>2</sub>	F <sub>3</sub>					
Environments	OI	WS	OI	WS	OI	WS	OI	WS				
Days to heading	-0.35	-0.02	-0.11	0.10	0.18	0.16	0.33	0.42				
Days to maturity	-0.12	0.84**	-0.49*	0.06	-0.28	0.43	0.31	0.48*				
Plant height (cm)	0.32	-0.57	-0.14	0.05	-0.02	-0.03	0.07	-0.44				
Spikes plant <sup>-1</sup>	0.75**	0.39	0.46*	0.33	0.36	-0.08	0.13	0.19				
Grains spike <sup>-1</sup>	0.15	-0.10	-0.09	0.20	-0.06	0.60*	0.20	0.10				
1000-grain weight (g)	-0.09	0.00	-0.39	-0.50*	-0.39	-0.18	-0.56*	0.49*				

**Table 8.** Correlation between grain yield per plant and other studied traits under optimal irrigation (OI) and water stress (WS).

This finding highlights the ability to include such families in the selective breeding programs to obtain new pure-lines and improve specific adaptive characters to drought stresses. Therefore, these bread wheat genotypes should be selected for the following breeding programs to improve yield under well irrigation and water stress environments. Similar findings have also been reported by Aberkane *et al.* (2020) and Ali *et al.* (2021) for yield related traits in bread and durum wheat.

Selecting early wheat lines with a long grain filling period (GFP) might lead to selecting genotypes with good levels of drought tolerance (Aberkane *et al.*, 2020). Li *et al.* (2021) reported that under severe water shortages, the old hexaploid genotypes with drought evading characteristics exhibited lower reduction for grain yield and aboveground biomass than modern wheat genotypes, whereas under moderate and mild water deficits modern genotypes had higher aboveground biomass and yields.

The correlation among studied traits and grain yield showed a positive correlation between NS/P and GY/P at two water regimes in two crosses (Table 8). In two generations, days to maturity correlated positively with TGW and GY/P under drought. The grains per spike positively correlated with GY/P under water stress in two generations for cross II. Thousand-grain weight was positively valued with GY/P under water stress in F<sub>3</sub> for cross II. Several studies reported the contribution of NG/S to higher yields under drought stress (Sardouei-Nasab *et al.*, 2019; Arifuzzaman *et al.*, 2020; Ali *et al.*, 2021). Wheat breeders strongly associated TGW and GY/P during drought (Aberkane *et al.*, 2020).

The selection of superior genetic genotypes between the enormous amount of recombinant and segregating offspring is a necessary but complex process in plant breeding. The development of wheat cultivars implicates cyclic crossing and selection trials

over long times (Ali et al., 2020). Usually, wheat scientists relied on phenotypic selection to define the genetic potential of families or individuals in the field and select the best one simultaneously displayed that multiple desirable characters. Phenotypic selection serves as a well-organized strategy to improve complex characters by continuously increasing the number of favorable alleles (Li et al., 2018; Al-Naggar et al., 2020). Conventional breeding and modern techniques had pushed the yearly genetic gain for wheat grain yield from ~0.7% to ~1.2%, but the current annual genetic yield of yield in main food crops counting wheat is not enough to meet future demands (Li et al., 2018). Hence, the hybridization followed by selection under optimal and drought stress conditions has been a demand to accelerate the genetic gain of wheat grain yield (Ali et al., 2021).

The estimates of phenotypic coefficients of variability (PCV) were a little higher than those of genotypic coefficients of variability (GCV) for all traits in two crosses under both water regimes, indicating the low effect of the environment on the studied characters. The ranges, means, genetic variances, PCV, and GCV estimates for most studied traits were higher under optimal irrigation than corresponding estimates under water stress condition in two crosses for F<sub>2</sub> and  $F_3$  populations. These present results agree with the findings of investigators Mahdy et al. (2015) and Al-Naggar et al. (2020), who reported higher GCV and PCV under full irrigation conditions than under water stress environments. On the opposite, the GCV and PCV values were higher under water stress than corresponding estimates under optimal irrigation conditions for DH and DM in cross I, and DM and TGW in cross II. Wheat scientists reported similar results for DM and TGW (Al-Naggar et al., 2020), where they recorded higher values for GCV and PCV under drought stress.

The study's results showed that the genotypic variances, GCV, and PCV values were higher in the  $F_2$  generation than corresponding estimates in the  $F_3$  generation under the two water treatments for all studied traits, except DM in two crosses, indicating the higher genetic variations between the plants of  $F_2$ . The highest estimates of GCV and PCV were exhibited by NG/S, GY/P, and TGW, whereas DH and DM showed the lowest ones.

The coefficients of heritability  $(h^2_{Bs})$  for all studied traits were higher under optimal irrigation than corresponding estimates under water stress conditions in the  $F_2$  and  $F_3$ generations for two wheat crosses, except NG/S for the cross I and NS/P for cross II in the  $F_2$  population. Moreover, it is higher in  $F_3$ than F<sub>2</sub> generations for all studied traits under two water regimes in two crosses. The  $h_{Bs}^2$ estimates were more than 50% for all traits in  $F_3$  generation for all cases. Significant estimates of  $h^2_{\,\text{Bs}}$  are associated with higher genetic variability, higher selective accuracy, and better possibilities for achievement in selecting wheat genotypes with greater grain yield productivity (Attri et al., 2021; Channa, 2022). Al-Naggar et al. (2020) investigated heritability estimates are higher under full watering than those under drought for DH, S/P, G/S, and GY/P.

In the same trend, the genetic advance (GA) estimates were higher under optimal irrigation than corresponding estimates under water stress condition for most studied traits in the  $F_2$  and  $F_3$  generations for the two wheat crosses. In comparison, GA values in the  $F_2$ generation showed higher corresponding estimates than in the  $F_3$  generation for all studied traits under both water conditions for the two wheat crosses. Therefore, in general, estimates of heritability  $(h^2_{Bs})$  and genetic advance (GA) were higher under optimal irrigation than under water stress conditions for the most of studied traits (Al-Naggar et al., 2020), indicating the effects of genotype  $\times$ environment interaction. Also, high heritability coupled with high to moderate genetic advance is a percent of the mean for DH, NS/P, PH, SL, GY/P, and 1000-grain weight (Attri et al., 2021; Channa, 2022). The genetic advance was the highest for GY/P followed by the number of productive tillers per plant and 1000-grain weight (Al-Naggar et al., 2020).

The maximum values of response to selection in the first cross were 17.67 and 13.17 cm for plant height and 11.05 and 18.77 g for 1000-grain weight under optimal irrigation and water stress, respectively. Positive values of correlation (r) between offsprings ( $F_3$ ) and parents ( $F_2$ ) in yield and its components were noted under optimal irrigation and water stress conditions in the two crosses. The regression results showed a high positive estimation in grains per spike under optimal irrigation and spikes per plant under water stress in the first cross, and grain yield per plant in both the optimal irrigation and water stress conditions in the second cross. Similar results have also been reported by Aziz *et al.* (2018), Attri *et al.* (2021), and Channa (2022).

## CONCLUSIONS

The outperforming wheat genotypes under drought stress and optimal irrigation conditions can be helpful in future wheat breeding programs. The Cross-I families three and 19, and Cross-II families one, four, 13, and 27 showed the best desirable values under optimal irrigation for yield traits. Cross-I families six, eight, 18, and 37, and Cross-II families one and 17 exhibited high GY/P under drought stress. Early selection for the earliness and yield component characters will effectively develop drought-tolerant and high-yielding wheat varieties. At the same time, direct selection for grain yield is not effective in early generations for developing drought-tolerant and high-yielding varieties. This information could help future wheat breeders' selection for improve water stress tolerance in bread wheat.

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