

## **GENE ACTION AND HERITABILITY OF DIALLEL CROSSES IN BREAD WHEAT (*Triticum aestivum* L.) UNDER VARIOUS NUMBER OF IRRIGATIONS**

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### **ABSTRACT**

The genetic system controlling yield and its contributing i.e., flag leaf area, days to heading, number of spikes/plant, number of grains/spike, 1000-grain weight and grain yield/plant for six local and bread wheat genotypes were studied, under three treatments of irrigations (1-six irrigates 2- four irrigates 3- two irrigates). The diallel cross among parental genotypes in all possible combinations, excluding reciprocals was used. The studied genotypes were, Sakha 93, Giza 168, Sahel 1, Gemmeiza 7, Gemmeiza 9 and Sids 1. This investigation was carried out during 2005/2006 and 2006/2007 seasons at Tag El-Ezz Agricultural Research Station, Dakhlia Governorate. In the second season 2006/2007 the obtained F<sub>1</sub>'s together with the six parents were evaluated in randomized complete block design with three replicates for each treatment of irrigation. The obtained results showed that, mean squares for types of gene action according to Jones Method (1956), additive genetic variance (a) was significant for all characters under various irrigation treatments. Whereas, the dominance genetic components (b) was significant for all characters except, number of spikes/plant for first treatment of irrigation (I<sub>1</sub>). By using second degree statistic (Hayman, 1954) results indicated that, the dominance genetic variance accounted for days to heading, number of spikes/plant (I<sub>1</sub>, I<sub>2</sub> and I<sub>3</sub>), number of grains/spike (I<sub>1</sub>) and grain yield/plant (I<sub>1</sub> and I<sub>2</sub>) resulting in  $(H_1/D)^{0.5}$  which was more than unity. Both positive and negative alleles ( $H_2/4H_1$ ) were not equally distributed among parents for all characters, except number of spikes/plant (I<sub>1</sub>). Heritability in narrow sense was more than 50% for flag leaf area and days to heading (I<sub>1</sub>, I<sub>2</sub> and I<sub>3</sub>) as well as number of spikes/plant (I<sub>3</sub>).

The mean squares of general and specific combining ability were significant for all characters. The wheat cultivars, Gemmeiza 7 and Gemmeiza 9 proved to be good general combiners. Wheat crosses; Sakha 93 X Gemmeiza 9, and Gemmeiza 7 X Sids 1 could be considered promising crosses and the best crosses combinations displayed for amount of heterotic effects for grain yield/plant.

### **INTRODUCTION**

Bread wheat (*Triticum aestivum* L.) is considered the unique cereal crop in both Egyptian and world cultivation. The uniqueness of wheat crop arises from its multiple usage in human consumption. The exponential increase in Egyptian population in comparison with the limited cultivated area, enhances the responsibility of plant breeders to bridge this gap through improving and developing high yielding wheat genotypes as a quick and cheaper solution in present time. The diallel analysis provide detailed genetical information about specific genotypes before including in breeding programs. Many researches used diallel technique to obtain genetical information about yield and yield contributing characters in this respect,

Mosaad *et al.*, (1990), Alkaddoussi *et al.* (1994); Awaad (1996); Salama *et al.* (2006), these researches studied gene action and genetic that controlled yield and yield contributing characters. The importance of dominance for flag leaf area, days to heading and grain yield/plant were studied by Shehab El-Din (1997); Ab-El-Aty (2002) and Esmail (2002). The importance of additive and dominance gene action for 1000-grain weight and number of grains/spike were detected by Salama (2000), and Salama *et al.*, (2005). Many researchers used diallel cross techniques in wheat, i.e. Eissa *et al.* (1994); El-Hindi *et al.* (2005) and Sultan *et al.* (2005).

The present investigation was undertaken to obtain genetic information involving gene action of genetic system, and heritability for yield and yield contributing characters in half diallel to 6 x 6 local wheat genotypes.

## MATERIALS AND METHODS

### 1- Experimental layout and the studied materials:

Six genetically diverse wheat genotypes of local origin were crossed in half diallel, excluding reciprocals, in 1<sup>st</sup> season; 2005/2006 to obtained 15 F<sub>1</sub>'s. The pedigree and origin of the studied parents are presented in Table (1).

**Table (1). Pedigree and origin of the studied bread wheat genotypes.**

Serial number	Genotypes	Pedigree	Origin
1	Sakha 93	Sakha 92 / TR 810328-5887/15-25-15-05	Egypt
2	Giza 168	MRL / BUC// Seri CM 93046-8M-04-OM-2Y-OB-062	Egypt
3	Sahel 1	N.S. 732 / PIM // Veery "S"	Egypt
4	Gemmeiza 7	CM74A. 630 / Sx // Seri 82/ 3/ Agent	Egypt
5	Gemmeiza 9	ALD "S" Huac "S" // CMH 74A/6301	Egypt
6	Sids 1	HD2171/Pavon "S"//1158. 57/rIaya	Egypt

Three field experiments were conducted at Tag El-Ezz Agricultural Reaserch Station, Dakahlia governorate, Agriculture Research Center (ARC) and evaluated in three experiments for 2006/2007 grown season.

The first experiment received 6 irrigations (I<sub>1</sub>). The 2<sup>nd</sup> experiment received 4 irrigations. Meanwhile, the 3<sup>rd</sup> experiment received 2 irrigations (I<sub>3</sub>) with sowing irrigation various irrigations treatments were applied under 15 F<sub>1</sub>'s and 6 parents in randomized caplete block design with three replicates. The plot of all experiments consisted of 6 rows (2 rows to for each parent and F<sub>1</sub>'s). Row length was 2m, row to row spacing was 20 cm, plant to plant spacing was 10 cm. All agricultural recommended practices for wheat production at the area of study, except irrigation, were applied at the proper time.

The aims of this study to obtained genetic inforation under wheat various irrigations, including gene action and heritability.

## **2- Collected data:**

The following characters were measured at the proper time for parents and F<sub>1</sub>'s using sample of 10 guarded and competitive plants for F<sub>1</sub>'s and each parents:

Flag leaf area (cm<sup>2</sup>), days to heading (day), number of spikes/plant, number grains/spikes (average mean of 10 main spike). Thousand grain weight (g.), and Grain yield / plant (g.)

## **3- Statistical analysis:**

The obtained data were subjected, firstly to conventional two way analysis of variance according to Steel and Torrie (1980).

Assessment and quantifying types of gene action were computed according to Hayman (1954), Jones (1956), and Mother and Jinks (1982).

General (GCA) and specific (SCA) combining ability variances and effects were estimated using model 1 and methods 2 (Griffine, 1956). Rank correlation (rs) between mean performance and GCA effects were computed.

## **RESULTS AND DISCUSSION**

Mean squares of half diallel analysis of variance for studied characters (Table 2) indicated significant mean squares of additive "a" for all studied characters in three treatments (I<sub>1</sub>, I<sub>2</sub> and I<sub>3</sub>) of irrigations, indicating the importance of additive gene action in controlling these characters. The importance of dominance gene effects "b" in the inheritance of studied characters were shown for all studied characters, except number of spikes/plant (I<sub>1</sub>). The obtained results were in harmony with those obtained by Alkaddoussi *et al.* (1994); Salama *et al.* (2005) and Salama and Manal Salem (2006).

When the dominance component "b" was partitioned to b<sub>1</sub>, b<sub>2</sub> and b<sub>3</sub> components, results of b<sub>1</sub> indicated that dominance was unidirectional for all characters, except number of spikes/plant for I<sub>1</sub>, expressing ambidirectional. The significance of b<sub>2</sub> component for all studied characters, indicating that dominant genes were not equally distributed among parental genotypes. The specific combining ability as indicated by b<sub>3</sub> were highly significant for all studied characters except, number of spikes/plant for I<sub>1</sub> and I<sub>3</sub> and grain yield / plant for I<sub>1</sub> and I<sub>2</sub> significant b<sub>3</sub>, suggesting superiority of some specific combinations which many display a high potential of genetic variability, expected to produce high yielding wheat genotypes. Significant b<sub>1</sub>, b<sub>2</sub> and b<sub>3</sub> were detected for 1000-grain weight and grain yield/plant in bread wheat (Alkaddoussi *et al.*, 1994 and Salama 2000).

Using second degree statistics Hayman (1954), various genetic parameters were computed (Tables 3 and 4) and indicated that both additive (D) and dominance (H<sub>1</sub> and H<sub>2</sub>) effects were significant for all the studied characters, indicating the importance of additive and dominance gene effects for controlling these characters. The dominance genetic variance was higher of magnitude as compared to additive variance resulting in (H<sub>1</sub>/D)<sup>0.5</sup> exceeding more than unity for days to heading and, number of spikes/plant

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for all treatments of irrigations, number of grains /spike ( $I_1$ ) and grain yield/plant ( $I_1$  and  $I_2$ ), thus hybrid breeding method would be an effective method for improving these characters. In this respect over dominance gene effects was reported for number of spikes/plant (Dasgupta and Mondal 1988), as well as number of grains/spike, 1000 grain weight and grain yield/plant [Eissa 1989; Al-Kaddoussi *et al.* 1994, Al-Kaddoussi 1996 and Awaad 1996)]. On the other hand, the remaining characters, the additive genetic variance was more than the dominance one.  $(H_1/D)^{0.5}$  was less than unity indicating the importance of additive gene effects in controlling these characters.

The covariance of additive and dominance gene effect in the parents revealed significant and positive "F" values for days to heading, number of grains/spike; 1000-grain weight and grain yield / plant for all treatments of irrigation, indicating that the dominant alleles were more frequent than the recessive ones in the parent for these characters, while negative "F" value for remaining characters indicated excess of recessive alleles among parents. The overall dominance effects of heterozygous loci ( $h^2$ ) indicated directional dominance for flag leaf area, number of spikes/plant and number of grains/spike for all treatments; days to heading for  $I_1$  and  $I_2$ , 1000- grain weight and grain yield / plant for  $I_1$ . The proportion of genes with positive and negative effects in the parent ( $H_2 / 4H_1$ ) were approximately equal to 0.25 for number of spikes / plant for  $I_1$  and  $I_2$  indicating equally distribution of positive and negative genes in the parental genotypes for this character. But, for the remaining characters ( $H_2 / 4H_1$ ) deviated from its maximum value (0.25). Thus, genes controlling these characters were not equally distributed among parents.

The ratio of dominance to recessive alleles (KD/KR) in the parents indicated the preponderance of dominance alleles which was  $>1$  for all characters, except flag leaf area showed an excess of decreasing alleles among parental genotypes. Narrow sense heritability ( $T_n$ ) was more than 50% for flag leaf area, days to heading in all treatments and number of spikes/plant for  $I_3$ . Heritability in broad sense ranged from 0.569-0.973 for all the studied characters under various irrigation treatments.

Mean squares for combining ability (Table 5) indicated that general (gca) and specific (sca) combining ability variances were significant for all characters, suggesting presence of additive and dominance gene effects in the inheritance of these characters. These results are in accordance with those obtained by (Eissa 1993, and Salama and Manal Salem 2006). The sca exceeded gca for number of spikes / plant and grain yield / plant (all treatments of irrigations), suggesting that non-additive gene effects accounted for the great part of the total variation for these characters. Thus, selection for these characters should be practised in to later generations.

The results in Table (6) indicated that general combining ability (gca) effects which refer to additive and additive x additive gene effects was positive and significant for flag leaf area for  $I_2$  and  $I_3$  (Gemmeiza 9), days to heading  $I_1$ ,  $I_2$  and  $I_3$  (Sahel 1) and  $I_3$  (Gemmeiza 7); number of spikes/plant  $I_1$  (Gemmeiza 7),  $I_2$  (Gemmeiza 9 and Sids 1) and  $I_3$  (Giza 168 and Gemmeiza 9), number of grains/spike  $I_1$  (Sakha 93) and  $I_3$  (Sids 1), 1000-grain weight  $I_1$  (Gemmeiza 7) and Gemmeiza 9) and  $I_2$  with  $I_3$  for Gemmeiza 9 and grain

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yield/plant for I<sub>1</sub>, I<sub>2</sub> with I<sub>2</sub> (Gemmeiza 7 and Gemmeiza 9). It could be concluded that Gemmeiza 7 and Gemmeiza 9 possessed more favorable genes for yield and its components in bread wheat. The obtained result were in harmony, with those detected by (Dasgupta and Mondel, 1988, Al-Kaddoussi, 1996 and Salama and Manal Salem, 2006).

The results of rank correlation between gca effects and performance revealed positive and significant value for number of grains/spike and 1000 - grain weight for three treatments of irrigations, indicating that parents could be judge through mean performance of these characters. Thus, when the characters is unidirectionally controlled by additive gene action, the breeder could isolate good general combiners as early as possible using mean value as quick and reliable estimate. Similar results was reported by Eissa (1993), Awaad (1996) and Hassan (1998).

As seen in Table (7) specific combining ability effects (sca) indicated that the best crosses displayed positive and significant sca effects for flag leaf area I<sub>2</sub> and I<sub>3</sub> (Sakha 93 x Sahel 1; I<sub>1</sub> (Sakha 93 x Sids 1), I<sub>2</sub> and I<sub>3</sub> (Giza 168 x Gemmeiza 9); days to heading for I<sub>1</sub> and I<sub>2</sub> (Sakha 93 x Giza 168); I<sub>2</sub> (Sahel 1 x Gemmiza 7), I<sub>3</sub> Sakha 93 x Gemmeiza 7, Giza 168 x Sids 1, Sahel 1 x Gemmeiza 9, and Gemmeiza 7 x Gemmeiza 9); number of spikes/plant for I<sub>1</sub>, I<sub>2</sub> and I<sub>3</sub> (Sakha 93 x Sids a and Gemmeiza 7 x Sids 1); number of grains / spike for I<sub>1</sub>, I<sub>2</sub> and I<sub>3</sub> (Sakha 93 x Sids 1, Giza 168 x Gemmeiza 9; and Gemmeiza 7 x Gemmeiza 9); 1000-grain weight for I<sub>2</sub>, I<sub>2</sub> and I<sub>3</sub> (Giza 168 x Sahel 1) and Sahel 1 x Sids 1) and grain yield / plant for I<sub>1</sub>, I<sub>2</sub> and I<sub>3</sub> (Sakha 93 x Gemmeiza 9 and Gemmeiza 7 x Sids 1). It could be concluded that the cross combinations; (Sakha 93 x Gemmiza 9 and Gemmeiza 7 X Sids 1) displayed sca effects for, yield and its contributing characters, suggesting their importance in wheat breeding programs. It could be seen that less environmental influence (Stable crosses) enhancing its valuable as promising one for improving grain yield in bread wheat under various irrigations treatments.





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### الفعل الجيني وكفاءة التوريث لهجن الدياليل في قمح الخبز تحت عدد مختلف من الريات سليمان محمد جمعة سلامة

المعمل المركزى لبحوث التصميم والتحليل الإحصائي – مركز البحوث الزراعية – الجيزة - مصر

- أجريت هذه الدراسة في الموسم الشتوى لأعوام ٢٠٠٦/٢٠٠٥ و ٢٠٠٦ / ٢٠٠٧ وذلك بغرض دراسة تأثير ثلاثة معاملات من الري وهي ٦ ريات، ٤ ريات، وريتين على الفعل الجيني وكفاءة التوريث لسته أصناف من قمح الخبز المصرى وهي: سخا ٩٣، جيزة ١٦٨، ساحل ١، جيزة ٧، جيزة ٩، سدس ١.
- وتم دراسة الصفات التالية مساحة ورقة العلم، عدد الأيام حتى طرد السنابل، وعدد السنابل للنبات وعدد حبوب السنبل ووزن الألف حبة، ومحصول الحبوب للنبات. وذلك بتجهين الستة أصناف بطريقة الدياليل مستبعداً الهجن العكسية وتم تقييم الجيل الأول والاباء بزراعتها بمحطة البحوث الزراعية بتاج العز دقهلية في ثلاث معاملات من الري كل على حدة في تجربة منفصلة (ثلاث تجارب) في ١٠ نوفمبر ٢٠٠٦ باستخدام تصميم القطاعات العشوائية الكاملة في ثلاث مكررات وتم تحليل البيانات إحصائياً ووراثياً باستخدام نماذج الدياليل جونز ١٩٥٦ وهابمان ١٩٥٤ وجرفنج ١٩٥٦، وقد أوضحت النتائج الآتى:
- ١- أظهر تحليل التباين بطريقة جونز ١٩٥٦ أن التباين الراجع للفعل الجيني المضيف كان معنوياً لجميع الصفات بينما كان التباين الراجع للفعل السيدى (b) معنوياً لجميع الصفات ماعدا عدد السنابل للنبات معاملة الري الأولى (٦ ريات).
  - ٢- أظهر التحليل بطريقة هابمان ١٩٥٤ أن المكون السيدى هو المكون الأعظم في وراثة عدد الأيام حتى طرد السنابل وعدد السنابل/نبات للثلاث معاملات من الري وعدد حبوب السنبل في معاملة الري الأولى (٦ ريات) ومحصول الحبوب للنبات في معاملي الري الأولى والثانية (٦ ريات و ٤ ريات) وكانت قيم متوسط درجة السيادة أعلى من الوحدة لهذه الصفات.
  - ٣- توزعت كل من الجينات الموجبة والسالبة توزيعاً غير متماثل بين الآباء لجميع الصفات ما عدا صفة عدد السنابل للنبات في معاملة الري الأولى (٦ ريات).
  - ٤- كانت كفاءة التوريث بالمعنى المحدود أكبر من ٥٠% لمساحة ورقة العلم وعدد الأيام حتى طرد السنابل للثلاث معاملات من الري وعدد السنابل للمعاملة الثالثة. بينما كانت أقل من ٥٠% لباقي الصفات.
  - ٥- وأوضحت النتائج أن التباين الراجع للقدرة العامة والخاصة على التآلف كان معنوياً لجميع الصفات المدروسة.
  - ٦- أظهرت النتائج أن الصنفين جيزة ٧ وجيزة ٩ كانت أفضل الآباء للقدرة العامة على التآلف لصفة محصول الحبوب ومعظم الصفات المرتبطة وأظهرت الهجن سخا ٩٣ × جيزة ٩ وجيزة ٧ × سدس ١ قدرة خاصة مرغوبة وعالية المعنوية ويمكن إستخدام هذه الهجن لزيادة محصول الحبوب للنبات (تحت ظروف الريات). وتؤكد النتائج أهمية إستخدام هذه الأصناف في إنتاج وعزل تراكيب وراثية متفوقة في المحصول ومكوناته تحت ظروف الريات المختلفة.

Table (2): Mean squares of half diallel analysis of variance for studied characters under three treatments of irrigations in bread wheat Jones 1956.

Character s	S.O.V. D.F	Flag leaf area (Cm) <sup>2</sup>			Days to heading (day)			Number of spikes/plant			Number of grains/spike			1000-grain weight (gm)			Grain yield /plant (gm)		
		I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>
a	5	68.916**	71.362**	53.27**	92.532**	111.291**	107.32**	0.823**	0.714**	0.652**	13.910**	14.622**	11.880**	16.522**	13.401**	15.834**	2.932*	1.106*	2.470**
b	15	51.326**	42.381**	36.281**	19.88**	20.232**	22.471**	0.521	0.823**	0.621*	19.141**	16.522**	17.362**	14.831**	12.693**	110.852**	1.902*	1.821**	1.793**
b <sub>1</sub>	1	50.312**	32.163**	29.852**	16.481**	15.94**	17.503**	0.636	0.922**	0.722**	18.411**	17.932**	16.523**	20.414**	28.513**	26.142**	3.173**	3.252**	1.873**
b <sub>2</sub>	5	21.522**	23.689**	27.112**	19.351**	18.420**	19.934**	0.962*	0.848**	0.751*	19.260**	19.253**	20.622**	21.413**	24.892**	14.858**	4.169**	3.810**	2.461**
b <sub>3</sub>	9	67.99**	53.900**	42.090**	20.55**	21.715**	24.432**	0.263	0.798**	0.520	19.156**	14.848**	15.644**	10.500**	4.602**	5.261**	0.506	0.557	1.413**
Error	40	2.611	3.292	3.848	5.170	0.362	6.473	0.403	0.259	0.310	4.823	3.031	4.312	3.94	2.33	2.82	0.907	0.423	0.692

I<sub>1</sub>, I<sub>2</sub>, and I<sub>3</sub> refers to 2, 4 and 6 irrigations.

Table (3): Additive (D), dominance (H) and environmental (E) genetic components together with derived parameters for flag area, days to heading and number of spikes/plant under three treatments of irrigation (I) of half 6 x 6 diallel cross in bread wheat.

Characters Parameters	Flag leaf area (Cm) <sup>2</sup>			Days to heading (day)			Number of spikes/plant		
	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>
D	10.300**+0.716	12.210**+0.652	11.74**+0.842	8.09**+0.978	9.362**+1.031	11.157**+1.079	0.819**+0.305	0.710**+0.285	0.780**+0.264
H <sub>1</sub>	6.431**+1.818	7.090**+2.052	7.21**+2.141	22.176**+2.482	25.868**+2.619	28.202**+2.741	2.266**+0.775	1.420**+0.725	2.439**+0.671
H <sub>2</sub>	5.662**+1.625	6.56**+1.833	6.94**+1.913	6.122**+2.218	7.604**+2.341	10.332**+2.449	2.253**+0.692	1.860**+0.648	1.298**+0.600
F	-3.300**+1.731	-1.127**+1.954	-3.51**+2.038	15.721**+2.363	18.321**+2.494	21.808**+2.610	0.049**+0.738	0.253**+0.690	0.170**+0.639
h <sup>2</sup>	3.652**+1.093	4.272**+1.233	5.62**+1.287	6.960**+1.492	0.419**+0.955	31.571**+1.648	19.06**+0.466	16.901**+0.436	10.614**+0.403
E	3.390**+0.270	3.681**+0.305	2.41**+0.318	0.179**+0.368	0.178**+0.389	0.174**+0.407	0.153**+0.115	0.095**+0.107	0.081**+0.100
Derived parameters									
(H <sub>1</sub> /D) <sup>0.5</sup>	0.790	0.762	0.591	1.655	1.662	1.589	1.659	1.644	3.126
H <sub>2</sub> /4H <sub>1</sub>	0.220	0.231	0.241	0.069	0.073	0.091	0.249	0.242	0.133
KD/KR	0.663	0.886	0.679	2.421	6.00	4.192	1.018	1.243	1.131
T(n)	0.599	0.566	0.651	0.711	0.691	0.567	0.350	0.315	0.683
T(b)	0.717	0.700	0.797	0.969	0.973	0.972	0.861	0.884	0.936

Table (4): Additive (D), dominance (H) and environmental (E) genetic components together with derived parameters for number of grains/spike, 1000, grain weight and grain yield/plant under three treatments of irrigation (I) of half 6 x 6 diallel cross in bread wheat.

Characters Parameters	Number of grains/spike			1000-grain weight (gm)			Grain yield/plant (gm.)		
	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>
D	5.589 <sup>***</sup> ±0.916	11.649 <sup>**</sup> ±3.141	10.999 <sup>**</sup> ±1.085	17.009 <sup>**</sup> ±1.063	13.11 <sup>**</sup> ±1.103	10.70 <sup>**</sup> ±0.978	8.32 <sup>**</sup> ±0.709	8.940 <sup>**</sup> ±0.74	10.28 <sup>**</sup> ±0.763
H <sub>1</sub>	17.674 <sup>**</sup> ±2.325	7.535±2.703	10.489 <sup>**</sup> ±2.755	15.138 <sup>**</sup> ±2.70	11.38 <sup>**</sup> ±2.615	7.700 <sup>**</sup> ±2.482	10.05 <sup>**</sup> ±1.802	9.69 <sup>**</sup> ±1.921	9.100 <sup>**</sup> ±1.94
H <sub>2</sub>	11.136 <sup>**</sup> ±2.013	5.568 <sup>**</sup> ±2.415	5.396 <sup>**</sup> ±2.460	10.418 <sup>**</sup> ±2.412	9.616 <sup>**</sup> ±2.337	6.08 <sup>**</sup> ±2.218	6.98 <sup>**</sup> ±1.616	6.50 <sup>**</sup> ±1.716	8.48 <sup>**</sup> ±1.732
F	6.874 <sup>**</sup> ±2.215	8.326 <sup>**</sup> ±2.573	14.280 <sup>**</sup> ±2.623	19.838 <sup>**</sup> ±2.571	12.077 <sup>**</sup> ±2.490	7.586 <sup>**</sup> ±2.363	9.78 <sup>**</sup> ±1.716	8.94 <sup>**</sup> ±0.757	10.05 <sup>**</sup> ±1.840
h <sup>2</sup>	29.32 <sup>**</sup> ±1.399	10.24 <sup>**</sup> ±1.625	4.00 <sup>**</sup> ±1.656	2.161±1.623	3.162±1.572	3.614 <sup>**</sup> ±1.492	1.63±1.083	2.232±1.155	4.322 <sup>**</sup> ±1.165
E	1.231 <sup>**</sup> ±0.346	1.314 <sup>**</sup> ±0.401	1.702 <sup>**</sup> ±0.409	2.311 <sup>**</sup> ±0.401	1.62 <sup>**</sup> ±0.388	1.240 <sup>**</sup> ±0.368	0.990 <sup>**</sup> ±0.268	1.05 <sup>**</sup> ±0.285	1.340 <sup>**</sup> ±0.288
Derived param.									
(H <sub>1</sub> /D) <sup>0.5</sup>	1.778	0.804	0.976	0.943	0.931	0.848	1.099	1.041	0.941
H <sub>2</sub> /4H <sub>1</sub>	0.161	0.184	0.128	0.172	0.211	0.214	0.173	0.168	0.233
KD/KR	2.057	2.599	4.925	4.217	2.955	2.435	3.302	3.729	3.162
T(n)	0.395	0.402	0.227	0.161	0.259	0.383	0.227	0.206	0.109
T(b)	0.814	0.754	0.569	0.605	0.701	0.758	0.720	0.688	0.653

Table (5): Mean squares of general (gca) and specific combining ability (sca) for studied character under three treatment of irrigation (I) of half 6x6 diallel cross in bread wheat.

Characters		Flag leaf area (Cm) <sup>2</sup>			Days to heading (day)			Number of spikes/plant			Number of grains/spike			1000-grain weight (gm)			Grain yield /plant (gm)		
S.O.V.	D.F	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>
g.c.a	5	68.916 <sup>**</sup>	71.362 <sup>**</sup>	53.07 <sup>**</sup>	92.532 <sup>**</sup>	111.291 <sup>**</sup>	107.32 <sup>**</sup>	0.823 <sup>**</sup>	0.714 <sup>**</sup>	0.652 <sup>**</sup>	13.910 <sup>**</sup>	14.622 <sup>**</sup>	11.880 <sup>**</sup>	16.522 <sup>**</sup>	13.401 <sup>**</sup>	15.834 <sup>**</sup>	2.932 <sup>**</sup>	1.106 <sup>**</sup>	2.470 <sup>**</sup>
S.c.a	15	54.231 <sup>**</sup>	44.262 <sup>**</sup>	42.513 <sup>**</sup>	91.143 <sup>**</sup>	82.367 <sup>**</sup>	58.40 <sup>**</sup>	1.529 <sup>**</sup>	1.913 <sup>**</sup>	1.422 <sup>**</sup>	13.183 <sup>**</sup>	12.622 <sup>**</sup>	11.198 <sup>**</sup>	13.940 <sup>**</sup>	6.879 <sup>**</sup>	9.262 <sup>**</sup>	3.315 <sup>**</sup>	2.987 <sup>**</sup>	3.038 <sup>**</sup>
Error	40	2.611	3.292	3.848	5.170	6.362	6.473	0.403	0.259	0.310	4.823	3.031	4.312	3.94	2.33	2.82	0.907	0.423	0.692
$\frac{\sigma^2_{gca}}{\sigma^2_{sca}}$		1.270	1.612	1.248	1.015	1.221	1.837	0.538	0.373	0.458	1.055	1.158	1.061	1.185	1.948	1.709	0.884	0.370	0.813

**Table (6): Mean performance, general combining ability effects and rank correlation for studied characters under three treatments of irrigation of half 6 x 6 diallel cross in bread wheat.**

Charac.	Flag leaf area (cm) <sup>2</sup>			Days to heading (day)			Number of spikes/plant			Number of grains/spike			1000-grain weight (gm)			Grain yield/plant (gm)			
	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	
<b>Parents</b>																			
Sakha 93	GCA	0.646	0.261	0.710	1.263	-0.851	0.829	-0.260	-0.283	-0.534**	2.123**	-0.340	0.567	-0.241	-1.503*	-1.269*	-1.793**	-0.839**	-0.513
	X	38.61	36.25	30.22	93.15	90.00	87.33	11.23	9.72	7.16	65.60	49.71	45.16	52.97	48.66	40.83	13.69	11.82	9.87
Giza 168	GCA	1.031	0.924	0.011	0.071	0.614	0.630	-0.514**	-0.930**	0.721**	-3.152**	-0.341	-2.821**	-0.262	-0.265	-0.707	-0.071	-0.837**	0.001
	X	36.66	34.18	33.20	99.62	95.66	92.15	12.29	9.87	6.63	52.66	45.88	44.16	52.36	43.21	40.16	12.58	11.01	9.63
Sahel 1	GCA	-0.373	-2.202**	-1.948**	2.531**	1.940*	2.824**	-0.232	0.025	-0.172	-0.070	-0.526	-2.142**	-3.610**	-0.259	-0.524	0.267	0.253	-0.282
	X	37.58	35.14	32.99	100.15	97.66	96.33	13.17	11.80	9.62	53.88	47.77	46.19	48.82	46.34	41.91	11.13	9.92	8.56
Gemmeiza 7	GCA	-1.242*	-0.312	-0.613	-2.21**	0.003	2.176**	0.674**	0.243	-0.614**	-0.282	0.894	0.941	2.147**	0.914	0.369	0.482*	0.491*	0.653*
	X	42.43	41.15	39.90	99.47	95.33	92.66	9.52	7.85	6.36	51.30	50.37	49.90	58.29	52.32	47.77	14.63	12.33	10.74
Gemmeiza 9	GCA	0.661	1.562**	1.503**	0.372*	-0.829	-1.797**	0.011	0.523**	0.542**	1.236	0.180	0.001	1.892**	1.858**	1.742**	0.810**	0.753**	0.652*
	X	43.28	40.72	37.91	103.33	101.33	98.15	8.95	7.62	5.38	64.30	60.16	47.22	59.11	52.39	49.76	13.85	12.07	10.82
Sids 1	GCA	-0.669	-0.233	0.337	-2.027**	-0.877	-4.662**	0.321	0.422**	-0.299	0.145	0.133	3.454**	0.074	-0.745	0.389	0.305	0.179	-0.511
	X	44.32	42.78	41.17	98.15	96.00	94.00	11.24	9.08	6.13	61.11	49.81	51.00	53.84	44.60	42.99	14.83	12.92	10.77
S.E (g)		0.521	0.585	0.632	0.733	0.813	0.600	0.204	0.164	0.179	0.708	0.561	0.669	0.640	0.492	0.541	0.229	0.210	0.268
Rank corr.		-0.486	-0.142	0.029	0.029	0.200	-0.142	-0.486	-0.771	-0.086	0.943**	0.885*	0.857*	0.943**	0.828*	0.885*	0.543	0.257	0.543

Table (7): Specific combining ability (sca) for studied character under three treatments of irrigations in half 6 x 6 diallel crosses of bread wheat.

Characters	Flag leaf area (cm <sup>2</sup> )			Days to heading (day)			Number of spikes/plant			Number of grains / spike			1000-grain weight (gm)			Grain yield / plant (gm)			
	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	I <sub>1</sub>	I <sub>2</sub>	I <sub>3</sub>	
<b>Crosses</b>																			
Sakha93 x Giza168	0.261	0.111	1.263	4.161**	5.310**	3.141	0.261	0.016	0.542	0.672	0.822	0.175	0.426	-0.534	0.172	0.111	0.362	0.157	
Sakha 93 x Sahel 1	0.072	3236**	3.251**	0.023	-0.101	-0.252	-0.022	0.104	0.152	-0.693	-1.812	0.410	0.184	0.170	0.293	0.623	-0.510	0.232	
Sakha93 x Gemmiza 7	-0.581	0.142	-0.651	1.111	2.322	4.621**	-0.132	-0.253	0.162	0.536	0.811	-0.936	0.460	0.508	0.272	-0.242	-0.163	-0.511	
Sakha 93 x Gemmiza 9	0.718	-0.160	-0.651	-0.562	-0.211	-3.622	0.429	0.151	0.201	4.262**	5.36**	5.292**	-4.616**	-5.152**	-6.143**	1.678**	1.325**	1.724**	
Sakha 93 x Sids 1	4.264**	2.152	1.122	-0.361	-0.822	0.811	1.413**	1.042**	1.262**	0.143	5.361	-6.264**	0.151	0.182	0.173	0.622	0.143	0.122	
Giza168 x Sahel 1	0.102	0.682	2.829	0.032	0.013	0.024	0.581	-0.322	-0.178	0.541	0.006	0.178	5.81**	5.362**	7.263**	0.726	0.801	0.653	
Giza168 x Gemmiza 7	-2.141	-0.142	-1.781	0.012	0.966	2.173	0.142	0.116	0.153	0.892	0.761	0.111	1.829	1.101	2.001	-0.013	-0.174	0.152	
Giza168 x Gemmiza 9	-1.563	1.240	3.412**	0.172	-0.123	0.152	0.556	0.531	0.111	3.948**	4.751**	6.293**	0.152	0.143	-0.111	0.840	0.925	1.013	
Giza 168 x Sids 1	-0.921	-0.523	-2.152	2.652	-1.826	-4.736**	-0.622	-0.132	0.107	-5.322**	-5.461**	-0.652	0.891	0.793	1.288	-0.814	-0.652	-0.163	
Sahel 1 x Gemmiza 7	-0.822	0.222	-0.828	2.763	3.916	2.363	-1.612**	-1.480**	-1.222**	0.613	0.142	0.136	-1.793	1.429	1.653	-1.610**	-1.442**	-1.253**	
Sahel 1 x Gemmiza 9	0.173	-0.006	0.712	2.93	0.781	4.610**	0.050	0.104	-0.006	0.858	0.006	0.963	-1.262	-1.422	0.819	-0.180	-0.114	0.126	
Sahel 1 x Sids 1	-5.022**	-0.053	0.146	-2.654	1.282	0.101	-0.053	0.156	0.174	0.943	0.988	1.293	6.140**	4.812**	3.880**	0.593	0.114	0.801	
Gemmiza 7 x Gemmiza 9	0.033	0.442	2.361	2.531	0.882	4.671**	0.236	0.482	-0.713	1.144	1.826	1.263	0.510	0.824	0.761	-0.101	-0.142	-0.123	
Gemmiza 7 x Sids 1	0.042	-3.122**	-0.802	0.513	-0.142	0.072	1.411**	0.998**	0.923**	9.176**	8.411**	7.652**	-0.531	-0.172	0.822	1.931**	1.409**	1.873**	
Gemmiza 9 x Sids 1	0.516	0.143	-0.011	-0.910	0.821	0.154	-0.154	-0.021	0.142	-5.361**	-4.252**	-5.130**	-0.717	0.163	0.183	-0.070	-0.182	-0.173	
S.E sgi	1.183	1.328	1.436	1.436	1.664	1.847	0.464	0.378	0.367	1.456	1.274	1.520	1.453	1.117	1.229	0.521	0.476	0.609	