# Estimate of Combining Ability and Correlation for Some Bread Wheat Genotypes

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#### Abstract:

Seven bread wheat (Sids1, Sids12, Sids13, Misr 1, Giza 171, Gammeiza11 and Sakha93) were used as parents of a 7 x 7 half-diallel cross. The obtained 21 F<sub>1</sub> crosses and their seven parents were evaluated at Experimental Farm, Faculty of Agriculture, Al-Azhar University, Assiut Branch. Results manifested that highly significant differences were detected among genotypes, parents, crosses, parents vs. crosses, general and specific combining abilities for all studied traits.  $\sigma^2 GCA/\sigma^2 SCA$  was less than the unity for all studied traits. The F<sub>1</sub> mean performance values were increased significantly over the parental mean values for all the studied traits, except number of days to 50% blooming, where they were not differ significantly. All crosses possessed desirable significant positive heterotic effects over both mid parent and better parent for grain yield/plant, except the crosses, P<sub>3</sub> x P<sub>7</sub> and P<sub>6</sub> x P <sub>7</sub>. The cross P<sub>5</sub> x P<sub>6</sub> had significant positive desirable SCA effects for high yielding ability and number of days to 50% blooming. Significant positive correlation was detected between grain yield/plant and both of plant height, number of spikes/plant, length of spike and number of grains/plant, while it gave negative correlation with number of days to 50% blooming and 1000-kernel weight.

Keywords: Wheat, combining ability, correlation.

#### Introduction

Wheat is considered one of great important food crops in the greatest part of the world. Wheat breeders do their best to explore the genetic material in order to develop new high yielding ability of wheat genotypes to face the gap between production and consumption. They need information on heterosis as essential in hybridization programs to develop new superior genotypes. Diallel mating system is one of the most powerful breeding analysis methods to give different genetic variances information about inheritance pattern of any specific trait of specific genotypes, which give guidance to the suitable breeding system to improve and release high yielding ability (Griffing, 1956). The important genetic parameters of diallel analysis

general (GCA) and specific are (SCA) combining ability, which are essential in developing plant breeding. Therefore, the concepts of GCA and SCA defined by Sprague and Tatum (1942), which have been used extensively in breeding of several economic species. The concepts of GCA and SCA became useful for characterization of inbred lines in crosses and often have been in the description of an inbred line (Hallauer and Miranda, 1988). The variances of general and specific combining ability are related to the type of gene action effects. Variance for GCA includes additive portion, while that of SCA includes non-additive portion of total variance arising largely from dominance and epistatic deviations (Rojas and Sprague, 1952).

The objectives of the present investigation were to estimate heterosis and correlation, as well as GCA and SCA effects of the seven wheat varieties and its 21  $F_1$  cross combinations to identify the best parental general combiners and crosses to improve earliness and yield of wheat.

## Material and Methods:

This investigation was carried out during two growing seasons; 2016/2017 and 2017/2018 at the Experimental Farm, Faculty of Agriculture, Al-Azhar University, Assiut Branch. Seven varieties of bread wheat wildly different in some their agronomic traits were used as parents of a 7 x 7 half-diallel cross in this study. These varieties were Sids 1 (P<sub>1</sub>), Sids 12 (P<sub>2</sub>), Sids 13 (P<sub>3</sub>), Misr 1 (P<sub>4</sub>), Giza 171 (P<sub>5</sub>), Gammeiza 11 (P<sub>6</sub>) and Sakha 93 (P<sub>7</sub>). The pedigree and origin of these varieties are shown in Table 1.

Data were recorded for days to 50% blooming (day), plant height (cm), number of spikes/plant, length of spike (cm), number of grains/plant (g), 1000-grain weight (g) and grain yield/plant (g).

 Table 1. The Pedigree and origin of the wheat varieties under investigation:

Name	Pedigree	Origin
Sids1	MRL/BUC/SER1	Egypt
	BUC//7C/ALD/5/MAYA74/0N//1160	
Sids 12	Egypt/47/3/BB/GLL/4/CHAT"S"/6/MAYA/VUL -	Egypt
	//CMH74A.63014*SX.SD7096-4SD-1SD-1SD-0SD.	
Sida13	ALMAZ.19=KAUZ"S"//TSI/SNB"S"IICSBW1-0375-4AP-	Egypt
Sluars	2AP-030AP-0APS-3AP-0APS-050AP-0AP-0SD	Egypt
Misr 1	OASIS/SKAUZ//4*BCN/3/2*PASTOR	Egypt
Giza 171	Sakha93 /Gemmeiza9 GZ003 – 101-1GZ -1GZ - 2 GZ -0GZ	Egypt
Gammeiza 11	BOW,,s,,/KVZ/7C/SERI82/3/GIZA168/SAKHA61	Egypt
Sakha 93	SAKHA 92/ TR 810328:	Egypt

# **Experimental layout:**

In the (2016/2017) season, the seven parents were sown in a field in two planting dates with two weeks apart to obtain enough flowers for crossing. Parents were crossed in all possible combinations, except reciprocals to produce twenty one  $F_1$  crosses.

In (2017/2018) season, the twenty one F<sub>1</sub> crosses and their seven parents were sown in a Randomized Complete Block Design (R.C.B.D) with three replications. Planting was carried out on 20<sup>th</sup> November (2017). Plants were grown on rows, 4 m long and 50 cm apart, in single seeded hill spaced at 15 cm. Each genotype was represented by three rows/plot. The agriculture practices of irrigation and fertilization were carried out as recommended for wheat production. The data were recorded on the mean of ten guarded plants/plot for each genotype.

## Statistical analysis:

An ordinary analysis of variance of RCBD for the obtained data was performed according to Snedecor and Cochran (1967) to test the significance of differences among the genotypes and treatments means were compared statistically using the test of the Lest Significant Differences (LSD). Mid and better parents heterosis was determined for each cross as percentage as follows:

Mid parents heterosis (%) =  $(F_1 - mid)$ parent/mid parent) x100.

Better parents heterosis (%) =  $(F_1 - F_1)$ better parent/better parent) x 100.

The analysis of combining ability was performed as outlined by Griffing's (1956), diallel cross analysis, method II, model I (fixed model). **Results and Discussion** 

## Analysis of variance and mean performance:

The analysis of variance (Table 2) cleared that highly significant differences were detected among genotypes for all the studied traits, indicating a wide range of diversity among the studied materials and combining ability analysis according to Griffing's (1956), method II, model I could be done. As well as, highly significant differences were detected among parents and crosses for all the studied traits, indicating great diversity among them. Also, highly significant differences between parents vs. crosses were detected for all the studied traits, indicating presence of highly heterosis response in the materials study. Both GCA and SCA mean squares were highly significant for all the studied traits, indicating both additive and non-additive gene actions are involved in the inheritance of all the studied traits. The ratio of  $\sigma^2 GCA / \sigma^2 SCA$  was less than one for all the studied traits, indicating the non-additive gene action played the major role in the inheritance of all the studied traits. These results are in agreement with those reported by Fonseca et al. (1968), Afiah et al. (2000), El-Beially and El-Sayed (2002), El-Borhamy (2004), Farooq et al. (2010) and Rahul and Kandalkar (2018).

Table 2. Mean squares of genotypes, parents, crosses, parents vs. crosses, general combining ability (GCA), specific combining ability (SCA) and their ratios. for yield and its components of the seven parents and their 21 F<sub>1</sub> crosses.

S.O.V	d.f	Number of days to 50% blooming	Plant height	Number of spikes/plant	0	Number of grains/plant	0	Grain yield/plant
Replicates	2	3.76	0.32	4.12	6.99	0.77	33.67	21.98
Genotypes	27	5.68**	78.89**	8.26**	6.38**	74.32**	54.83**	23.32**
Parents (P)	6	234.80**	152.40**	101.21**	65.47**	107.60**	80.59**	102.03**
Crosses (C)	20	89.33**	77.98**	43.65**	40.33**	50.22**	46.43**	76.28**
P vs C	1	53.32**	33.76**		18.98**	23.42**	16.88**	25.45**
GCA	6	8.31**	112.60**	11.21**	5.47**	137.70**	88.69**	12.33**
SCA	21	9.91**	122.70**		6.64**	155.45**	96.96**	25.08**
Error	54	0.87	0.79	1.00	0.37	2.25	1.99	1.04
∑gij2/∑sij2		0.84	0.92	0.95	0.64	0.86	0.92	0.49

\*\* Indicate to significant at 0.01 level of probability, respectively.

#### **Mean performance:**

Mean performance values of the seven parents and their twenty one  $F_1$ crosses are presented in Table 3. The obtained results exhibited that mean performance of the parents was wide extended with a range of 81.41 -84.88 days  $(P_5 - P_7)$ , 91.74 106.31 cm  $(P_3 - P_6)$ , 11.11 - 14.44  $(P_6 - P_1)$ , 13.18 - 14.78  $(P_3 - P_4)$ , 67.54 - 84.68  $(P_5 - P_6 \text{ and } P_7), 43.21 - 52.68 (P_6 - P_6)$  $P_5$ ) and 25.08-29.58 ( $P_7 - P_5$ ) for number of days to 50% blooming, plant height, number of spikes/plant, spike length, number of grains/plant, 1000-grain weight and grain vield/plant, respectively. Meanwhile, mean performance values of the  $F_1$ crosses were extended with a range of  $74.14 - 85.08 \text{ day } (P_5 \times P_6) - (P_2 \times P_6)$  $P_3$ ), 101.81 - 110.71 cm ( $P_4 \times P_7$ ) - ( $P_2$ x P<sub>6</sub>), 12.78 - 17.61 (P<sub>2</sub> x P<sub>5</sub>) - (P<sub>1</sub> x  $P_3$ ), 15.08 - 18.38 cm ( $P_1 \times P_3$ ) - ( $P_4 \times P_3$ )  $P_6$ ), 78.04 - 93.71 ( $P_1 \times P_5$ ) - ( $P_2 \times P_6$ ),  $48.28 - 56.41 \text{ g} (P_1 \times P_6) - (P_2 \times P_5)$ and P<sub>5</sub> x P<sub>7</sub>) and 25.08 - 36.38 (P<sub>6</sub> x  $P_7$ ) - ( $P_2 \times P_7$ ) for the abovementioned traits, respectively. The  $F_1$ mean performance values were surpassed significantly over the parental mean values for all the studied traits, except number of days to 50% blooming, where they were not differ significantly. Apparently, the different means among the seven parents and their  $F_1$  seemed to be valuable in improving the studied traits in bread wheat breeding programs.

Table 3. Mean performance values	of the 7 parents a	and their 21 F	1 crosses for all
the studied traits.			

unc	e studied tra	aits.					
Traits	Number of days to 50% blooming (day)	Plant height (cm)	Number of spikes/plant	Length of spike (cm)	Number of grains/plant	1000- grain weight (g)	Grain yield/plant (g)
P <sub>1</sub>	82.44	102.91	14.44	13.24	75.83	45.38	28.81
P <sub>2</sub>	84.54	95.41	11.84	14.04	80.54	47.54	28.18
P <sub>3</sub>	84.71	91.74	14.41	13.18	81.75	45.01	27.65
P <sub>4</sub>	83.01	95.88	12.44	14.78	77.58	47.54	28.24
P <sub>5</sub>	81.41	104.81	12.58	14.08	67.54	52.68	29.58
P <sub>6</sub>	82.6	106.31	11.11	13.91	84.68	43.21	26.94
P <sub>7</sub>	84.88	99.31	13.84	14.04	84.68	50.31	25.08
Mean	83.37	99.48	12.95	13.90	78.94	47.38	27.78
$P_1 x P_2$	81.24	106.71	15.28	16.24	82.28	50.34	31.94
$P_1 x P_3$	82.61	105.91	17.61	15.08	85.68	49.44	31.41
$P_1 x P_4$	82.41	106.01	15.58	17.64	80.18	52.04	31.64
$P_1 x P_5$	81.58	109.41	15.41	15.74	78.04	54.98	33.18
$P_1 x P_6$	80.64	108.31	15.08	16.41	85.74	48.28	31.41
$P_1 x P_7$	84.64	106.41	15.44	16.51	89.61	54.34	32.58
$P_2 x P_3$	85.08	102.31	15.38	15.78	86.61	51.04	30.88
$P_2 x P_4$	82.14	102.01	13.21	17.52	86.64	52.68	32.88
P <sub>2</sub> xP <sub>5</sub>	82.48	108.41	12.78	16.74	85.04	56.41	33.68
$P_2 x P_6$	82.94	110.71	16.41	16.74	93.71	50.78	29.78
$P_2 x P_7$	82.68	103.51	17.31	16.48	87.24	54.31	36.38
P <sub>3</sub> xP <sub>4</sub>	82.81	109.01	15.64	17.74	85.51	50.91	30.71
P <sub>3</sub> xP <sub>5</sub>	83.64	106.61	15.98	16.98	82.38	54.71	32.38
P <sub>3</sub> xP <sub>6</sub>	82.98	108.61	15.61	15.81	85.68	49.44	30.98
P <sub>3</sub> xP <sub>7</sub>	83.21	102.21	15.04	16.38	91.81	53.44	29.51
P <sub>4</sub> xP <sub>5</sub>	82.64	107.01	13.78	17.81	80.54	55.64	32.51
P <sub>4</sub> xP <sub>6</sub>	83.48	108.61	15.08	18.38	88.54	51.68	30.61
P <sub>4</sub> xP <sub>7</sub>	82.54	101.81	15.31	17.58	85.21	53.51	32.91
P <sub>5</sub> xP <sub>6</sub>	74.14	108.91	15.64	16.74	89.18	56.11	34.01
P <sub>5</sub> xP <sub>7</sub>	83.41	107.71	15.51	16.61	86.98	56.41	34.94
P <sub>6</sub> xP <sub>7</sub>	83.41	109.51	15.51	16.61	86.98	53.01	25.08
Mean	82.41	106.65	15.36	16.74	85.88	52.83	31.88
LSD <sub>0.05</sub>	1.54	1.47	1.65	1.00	2.48	2.33	1.68
LSD <sub>0.01</sub>	2.06	1.96	2.21	1.34	3.31	3.11	2.25

#### **Heterosis:**

Data in Table 4 showed that there were significant values for the heterosis over mid and better parent for all the studied traits, indicating that heterosis played an important role in the inheritance of these traits. For number of days to 50% blooming only cross  $P_5 \times P_6$  possessed significant negative desirable standard heterotic effect over both mid parent and better parent.

the F <sub>1</sub> crosses for all the studied traits.														
	Number of days			height	Numb		Length of spike Number of					-grain	Grain	
Traits.	to 50% k	8	•		spikes/plant		-	-	grains/plant		weight		yield/plant	
	M.P	B.P	M.P	B.P	M.P	B.P	M.P	B.P	M.P	B.P	M.P	B.P	M.P	B.P
P1 X P2	-2.55**	-0.21	7.27**	0.38	17.99**	5.82**	16.83**	9.88**	4.23**	-2.83*	6.25**	-4.44**	14.97**	7.98**
P1 X P3	-0.91	1.47	6.46**	-0.38	35.98**	21.95**	8.49**	2.03**	8.54**	1.18	4.35**	-6.15**	13.07**	6.19**
P1 X P4	-1.15	1.23	6.56**	-0.28	20.31**	7.89***	26.91**	19.35**	1.57	-5.31**	9.84**	-1.21	13.89**	6.96**
P1 X P5	-2.15**	0.21	9.98**	2.92**	19.00**	6.72**	13.24**	6.50**	-1.14	-7.84**	16.04**	4.37**	19.44**	12.17**
P1 X P6	-3.27**	-0.95	8.88**	1.88*	16.45**	4.43**	18.06**	11.03**	8.61**	1.25	1.90	-8.35**	13.07**	6.19**
P1 X P7	1.52	3.97**	6.97**	0.09	19.23**	6.93**	18.78**	11.71**	13.52**	5.82**	14.69**	3.15**	17.28**	10.14**
P2 X P3	2.05*	4.51**	2.84**	-3.76**	18.76**	6.51**	13.53**	6.77**	9.72**	2.28	7.72**	-3.11**	11.16**	4.39**
P2 X P4	-1.48	0.90	2.54**	-4.04**	2.01*	-8.52**	26.04**	18.54**	9.75**	2.31	11.19**	0.00	18.36**	11.16**
P2 X P5	-1.07	1.31	8.98**	1.98**	-1.31	-11.50**	20.43**	13.26**	7.73**	0.43	19.06**	7.08**	21.24**	13.86**
P2 X P6	-0.52	$1.88^{*}$	11.29**	4.14**	26.72**	13.64**	20.43**	13.26**	18.71**	10.66**	7.18**	-3.61**	7.20**	0.68
P2 X P7	-0.83	1.56*	4.05**	-2.63**	33.67**	19.88**	18.56**	11.50**	10.51**	3.02*	14.63**	3.09**	30.96**	22.99**
P3 X P4	-0.67	$1.72^{*}$	9.58**	2.54**	20.77**	8.31**	27.63**	20.03**	8.32**	0.98	7.45**	-3.36**	10.55**	3.82**
P3 X P5	0.32	2.74**	7.17**	0.28	23.40**	10.66**	22.16**	14.88**	4.36**	<b>-</b> 2.72 <sup>*</sup>	15.47**	3.85**	16.56**	9.47**
P3 X P6	-0.47	1.93*	9.18**	2.16**	20.54**	8.10**	13.74**	6.97**	8.54**	1.18	4.35**	-6.15**	11.52**	4.73**
P3 x P7	-0.19	2.21**	2.74**	-3.86**	16.14**	4.16**	17.84**	10.83**	16.30**	8.42**	12.79**	1.44	6.23**	-0.24
P4 X P5	-0.88	1.51	7.57**	0.66	6.41**	-4.57**	28.13**	20.50**	2.03	-4.89**	17.43**	5.62**	17.03**	9.91**
P4 X P6	0.13	2.54**	9.18**	2.16**	16.45**	4.43**	32.23**	24.36**	12.16**	4.56**	9.08**	-1.90	10.19**	3.48**
P4 X P7	-1.00	1.39	2.34**	-4.23**	18.22**	6.02**	26.47**	18.94**	7.94**	0.63	12.94**	1.58	18.47**	11.26**
P5 X P6	-11.07**	-8.93**	9.48**	2.45**	20.77**	8.31**	20.43**	13.26**	12.97**	5.31**	18.43**	6.51**	22.43**	14.98**
P5 X P7	0.05	2.46**	8.27**	1.32	19.77**	7.41**	19.50**	12.38**	10.18**	2.72*	19.06**	7.08**	25.77**	18.12**
P6 X P7	0.05	2.46**	10.08**	3.01**	19.77**	7.41**	19.50**	12.38**	10.18**	2.72*	11.88**	0.63	-9.72**	-15.21**

Table 4. Heterosis as percentage of mid-parents (M.P) and better parent (B.P) in the  $F_1$  crosses for all the studied traits.

and \*\* indicate to significant at 0.05 and 0.01 levels of probability, respectively.

For plant height, number of grains/plant and 1000-grain weight; nine, eight and eight crosses out of the twenty one possessed desirable significant positive heterotic effects over both mid parent and better parent, respectively. For number of spikes/plant, length of spike/plant and grain yield/plant all crosses possessed desirable significant positive heterotic effects over both mid parent and bet-

ter parent, except the three crosses;  $P_2$  x  $P_4$ ,  $P_2$  x  $P_5$  and  $P_4$  x  $P_5$  for number of spikes/plant and the two crosses;  $P_3$  x  $P_7$  and  $P_6$  x  $P_7$  for grain yield/plant. These results are supported with the findings of Ashoush *et al.* (2001), Jahanzeb and Ihsan (2004), El-Sayed and Moshref (2005), Abdel-Moneam (2009), Peng *et al.* (2009), Kundan *et al.* (2010), Ahmad (2010) and Zaazaa *et al.* (2012).

## A- General combining ability:

Data in table 5 revealed that  $P_1$ ,  $P_2$  and  $P_3$  recorded significant desirable GCA effects for plant height.  $P_4$  had significant positive desirable GCA effects for both plant height and increasing length of spike/plant.  $P_5$  possessed significant desirable GCA effects for number of days to 50%

blooming, 1000-grain weight and grain yield/plant.  $P_6$  possessed significant desirable GCA effects for number of days to 50% blooming, plant heightens and number of grains/plant.  $P_7$  possessed significant desirable GCA effects for plant shortness, number of grains/plant and 1000-grain weight.

 Table 5. General combining ability effects of the seven wheat varieties for all the studied traits.

Traits	Number of days to 50% blooming	Plant height	Number of spikes/plant	Length of spike	Number of grains/plant	1000-grain weight	Grain yield/plant
<b>P</b> <sub>1</sub>	-0.12	$0.78^{**}$	0.19	-0.15	-0.74*	-0.43	0.11
<b>P</b> <sub>2</sub>	0.16	1.06**	-0.15	-0.03	0.35	-0.04	0.19
P <sub>3</sub>	0.32*	1.21**	0.22	-0.15	0.30	-0.47*	-0.21
P <sub>4</sub>	0.03	0.93**	-0.17	$0.30^{*}$	-0.42	-0.01	0.03
P <sub>5</sub>	-0.39**	0.51	-0.14	0.02	-1.33**	1.03**	$0.48^{*}$
P <sub>6</sub>	-0.31**	0.59*	-0.09	0.01	0.96**	-0.59*	-0.41*
<b>P</b> <sub>7</sub>	0.31*	-5.08**	0.14	0.001	0.89**	$0.51^{*}$	-0.19
LSE 0.05	0.29	0.55	0.31	0.19	0.46	0.44	0.31
LSE 0.01	0.48	0.74	0.52	0.32	0.77	0.69	0.52

\* and \*\* indicate to significant at 0.05 and 0.01 levels of probability, respectively.

## **B-Specific combining ability:**

Data in Table 6 illustrated that the cross  $P_5 \times P_6$  only possessed significant negative desirable SCA effects for number of days to 50% blooming. The six crosses;  $P_1 \times P_7$ ,  $P_2 \times P_7$ ,  $P_3 \times P_7$ ,  $P_4 \times P_7$ ,  $P_5 \times P_7$  and  $P_6 \times P_7$  possessed significant positive desirable SCA effects for plant heightens, on the other hand only the cross  $P_5 \times P_6$  possessed significant negative desirable SCA effects for plant shortness. None of the crosses possessed significant desirable SCA effects for number of spikes/plant, length of spike and 1000-grain weight. The five crosses;  $P_1 \times P_7$ ,  $P_2$  $x P_6$ ,  $P_3 x P_7$ ,  $P_5 x P_6$  and  $P_5 x P_7$  possessed significant positive desirable effects for number SCA of grains/plant. The three crosses;  $P_2 x$  $P_7$ ,  $P_5 \times P_6$  and  $P_5 \times P_7$  possessed significant positive desirable SCA effects for high yielding ability and among them the cross  $P_5 \times P_6$ , also possessed significant desirable SCA effects for number of days to 50% blooming, therefore this cross could be considered as good combination for high yielding ability with earliness.

Crosses	Number of days to 50% blooming	Plant height	Number of spikes/plant	Length of spike	grains/plant		Grain yield/plant
$P_1XP_2$	-0.52	-1.30	0.53	0.25	-0.23	0.09	0.06
$P_1XP_3$	-0.21	-1.00	0.25	-0.01	0.96	0.22	0.29
$P_1XP_4$	0.01	-0.78	0.17	0.39	-0.16	0.63	0.12
$P_1XP_5$	0.15	-0.63	0.01	0.04	0.04	0.57	0.18
$P_1XP_6$	-0.24	-1.03	-0.10	0.27	0.31	-0.04	0.49
$P_1XP_7$	0.47	5.97**	0.13	0.31	1.67*	0.87	0.66
$P_2XP_3$	0.33	-0.46	-0.20	0.09	0.18	0.37	0.03
P <sub>2</sub> XP <sub>4</sub>	-0.36	-1.15	-0.37	0.22	0.90	0.45	0.45
P <sub>2</sub> XP <sub>5</sub>	0.17	-0.62	0.79	0.24	1.28	0.66	0.27
P <sub>2</sub> XP <sub>6</sub>	0.24	-0.54	0.86	0.25	1.88*	0.41	-0.14
P <sub>2</sub> XP <sub>7</sub>	-0.47	5.03**	0.24	0.17	-0.21	0.47	1.85**
P <sub>3</sub> XP <sub>4</sub>	-0.29	-1.08	0.33	0.43	0.58	0.29	0.13
P <sub>3</sub> XP <sub>5</sub>	0.40	-0.38	0.15	0.45	0.45	0.52	0.24
P <sub>3</sub> XP <sub>6</sub>	0.11	-0.68	-0.27	0.07	-0.75	0.39	0.66
P <sub>3</sub> xP <sub>7</sub>	-0.44	5.06**	-0.02	0.27	1.37*	0.62	-0.04
P <sub>4</sub> XP <sub>5</sub>	0.35	-0.43	0.37	0.28	0.55	0.37	0.04
P <sub>4</sub> XP <sub>6</sub>	0.56	-0.23	0.22	0.48	0.92	0.67	0.30
P <sub>4</sub> XP <sub>7</sub>	-0.38	5.12**	0.53	0.22	-0.11	0.17	0.85
P <sub>5</sub> XP <sub>6</sub>	-2.14**	-2.92**	0.25	0.21	2.05**	1.11	0.98*
P <sub>5</sub> XP <sub>7</sub>	0.33	5.83**	0.21	0.17	1.39*	0.10	$1.07^{*}$
P <sub>6</sub> XP <sub>7</sub>	0.25	5.75**	-4.08**	0.18	-0.91	0.59	-1.32*
LSE 0.05	0.84	1.61	0.90	0.55	1.35	1.27	0.92
LSE 0.01	1.39	2.16	1.34	0.93	2.15	2.05	1.47

Table 6. Specific combining ability effects of the 21 F<sub>1</sub> bread wheat crosses for all the studied traits.

\* and \*\* indicate to significant at 0.05 and 0.01 levels of probability, respectively.

#### **Correlation:**

The correlation values between each pairs of studied traits were shown in Table 7. The obtained results manifested that significant positive correlation was detected between grain yield/plant and both of plant height, number of spikes/plant, length of spike and number of grains/plant, while it gave negative correlation with number of days to 50% blooming and 1000-kernel weight. These obtained results are in the same line with those obtained by El-Sayed and Mostafa (2005), Khaled and Abd Eldayem (2014) and Samier and Ismail (2015).

Traits	Plant height	Number of days to 50%	Number	Length of	Number of grains/plant	1000-grain weight	Grain yield/plant
Plant height	-	-0.58**	0.25	0.59**	-0.46*	0. 50*	0.50*
Number of days to 50% blooming		-	-0.47*	-0.44*	-0.47*	-0.56**	-0.53**
Number of spikes/plant			-	0.41*	0.53**	-0.51**	0.66**
Length of spike				-	0.42*	$0.47^{*}$	0.57**
Number of grains/plant					-	-0.47*	0.59**
1000-grain weight						-	-0.58**
Grain yield/plant							-

Table 7. Correlation among studied characters for genotype.

and \*\* indicate to significant at 0.05 and 0.01 levels of probability, respectively.

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Zaazaa, E.I., M.A. Hager and E.F. El-Hashash (2012). Genetical analysis of some quantitative traits in wheat using six parameters genetic model. American-Eurasian J. Agric. & Environ. Sci., 12 (4): 456-462. تقدير القدرة على التالف والارتباط لبعض التراكيب الوراثية من قمح الخبز مختار حسن هريدي، ابراهيم نجاح عبد الظاهر، أحمد يوسف مهدي قسم المحاصيل ، كلية الزراعة ، جامعة الأز هر باسيوط

## الملخص

أجريت هذه الدراسة بالمزرعة التجريبية بقسم المحاصيل بكلية الزراعة جامعة الأزهر (فرع أسيوط) خلال موسمي ٢٠١٧/٢٠١٦ و ٢٠١٧/٢٠١٢. تم أختيار سبعة آباء من قمح الخبز لهذه الدراسة وهي سدس ١ وسدس ١٢ وسدس ١٣ ومصر ١ وجيزة ١٧١ وجميزة ١١ وسخا ٩٣. في موسم ٢٠١٧/٢٠١٦ تم إجراء التهجين بين الـ ٧ اباء بكل الطرق الممكنة بدون إجراء الهجن العكسية لإنتاج حبوب الـ ٢١ هجين فردى. في موسم ٢٠١٨/٢٠١٢ ، تم تحليل البيانات وراثيا لمحصول الحبوب/نبات وبعض الصفات ذات الصلة للاباء والهجن تبعا للطريقة الثانية، الموديل الأول. أظهر تحليل التباين وجود اختلافات عالية المعنوية بين التراكيب الوراثية والآباء والهجن والأباء مقابل الهجن (تأثير قوة الهجين). كان التباين الراجع إلى القدرة العامة والخاصة على التالف عالي المعنوية لجميع الصفات المدروسة، مما يشير إلى أهمية كلا من فعل الجين المضيف وغير المضيف في وراثتها. كانت نسبة تباين القدرة العامة على الجيني غير الخاصة على التآلف أقل من واحد لجميع الصفات المدروسة، مما يشير إلى أمية من فعل الجين المنيف وغير المصيف في وراثتها. كانت نسبة تباين القدرة العامة على الجيني غير الناصة على التالف الله من واحد لجميع الصفات المدروسة، مما يشير إلى أن الفعل الجيني غير المضيف وغير المضيف في وراثتها. كانت نسبة تباين القدرة العامة على التالف الي تباين القدرة المضيف على التالف الم من واحد لجميع الصفات المدروسة، مما يشير إلى أن الفعل الجيني غير المناسة على التالف أقل من واحد لجميع الصفات المدروسة، مما يشير الى أن الفعل الجيني غير الماميف لعب الدور الأهم في توريثها. كان هناك ارتباط موجب ومعنوى بين محصول حبوب النبات وكلا من طول النبات وعدد السنابل/نبات وطول السنبلة وعدد حبوب السنبله بينما كان