

## GENETIC ANALYSIS OF EARLINESS AND GRAIN YIELD TRAITS IN TWO WHEAT CROSSES

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### ABSTRACT

*The six parameter model was used to determine the intra - and inter-allelic gene interactions controlling the inheritance of earliness and yield traits in two crosses of wheat ( cross I : Saunval X Line 126; cross II : Gemmeza 9 X Sids 4). Analysis of variance indicated that significant differences existed among generations i.e P<sub>1</sub>, P<sub>2</sub>, F<sub>1</sub>, F<sub>2</sub>, BC<sub>1</sub> and BC<sub>2</sub> for all studied traits in both crosses. Significant and positive additive (a) effects were exhibited in days to heading, days to maturity, spikes/plant and grain yield/plant in both crosses. This assures the enhancing effect of additive variance in the inheritance of these traits. Dominance (d) effects were significant and positive for grain filling period and 1000 - kernel weight in both crosses, days to heading, days to maturity, spikes/plant and grain yield/plant in cross I. Additive was larger in magnitude than dominance effect for days to heading in both crosses, days to maturity, spikes/plant and grain yield/plant in cross II. Digenetic epistatic gene effects were generally important in the inheritance of studied traits. Additive X additive (aa) gene effects were generally higher in magnitude than ad and dd gene effects. Significant and positive aa effects were exhibited for grain filling period in both crosses, days to heading, days to maturity and grain yield/plant in cross I. Additive X dominance type of epistasis exhibited significant and positive effects in only two cases, namely days to heading in cross I and grain filling duration in cross II. Significant and positive dominance X dominance epistatic effects were shown for days to maturity, spikes/plant and grain yield/plant in cross II. Significant and highly significant desirable percentages of heterosis relative to mid- parent were exhibited for days to heading, grain filling period, spikes / plant and grain yield / plant in both crosses. Only grain yield / plant of cross No. 1 showed highly significant desirable heterosis relative to the better parent of 43.24 %. Very high broad- sense heritability percentages (> 90 %) were obtained in all studied characters for both crosses. Narrow- sense heritability percentage was the lowest ( 21.22 % ) for days to heading, medium ( 39.12 % ) for grain filling period and the highest ( 88.02 % ) for days to heading; all in cross I. Percentage of expected genetic advance from selection was the highest (55.44 %) for number of spikes/plant in cross II, and the lowest ( 2.40 % ) for days to heading in cross I.*

Key words: *Wheat, Triticum aestivum, Earliness, Six parameter model, Epistasis, Heterosis, Heritability.*

## INTRODUCTION

Wheat is the most important cereal crop in Egypt and world wide. Increasing grain yield and improving earliness of wheat in Egypt are considered important national goals to face the growing needs of the population; therefore, it has become necessary to develop early- maturing and high-yielding wheat genotypes.

The possibility of double cropping wheat and cotton in Egypt has heightened interest in early-maturing high-yielding wheats (Menshawy 2007). Early harvest of the wheat crop is critical to allow cotton crop sufficient time to develop and to produce an adequate yield.

A better understanding of the type of gene action and inheritance of earliness and yield traits would help wheat breeders to formulate the most advantageous breeding procedures for improving such traits. Additive gene action is evidently accounted for a large amount of the variation for days to heading (Bhatt 1972 Avey *et al* 1982 and Menshawy 2000 and 2005), days to maturity (Menshawy 2000, 2005 and 2007) and grain filling duration (Beiquan and Kronstad 1994 and Menshawy 2004), but dominance also was important (Crumpacker and Allard 1962, Avey *et al* 1982 and Menshawy 2005) for earliness traits, while epistasis was reported in few studies (Amaya *et al* 1972 and Ketata *et al* 1976) for earliness traits.

The present work was undertaken to study the role of different intra- and inter-allelic gene interactions, controlling the inheritance of earliness and yield traits in two bread wheat crosses.

## MATERIALS AND METHODS

The present investigation was carried out at the experimental field of Giza Research Station, Agricultural Research Center (ARC), during three successive seasons from 2003/2004 to 2005/2006. Four bread wheat genotypes namely, Gemmeiza 9, Sids 4, Line # 126 and Sunval representing a wide range of variability in earliness traits were used as parents of two crosses in this study (Table 1). In the first season (2003/2004) two crosses (Sunval X Line 126 cross I and Gemmeiza 9 X Sids 4 cross II) were made to obtain their F<sub>1</sub> seeds. In the second season (2004/2005), the hybrid seeds were sown and F<sub>1</sub> plants of each cross were backcrossed to their respective parents to produce the two backcrosses (BC<sub>1</sub>'s and BC<sub>2</sub>'s). At the same time, the F<sub>1</sub> plants were selfed to produce F<sub>2</sub> seeds. In the third season (2005/2006), the obtained seeds of these populations, i.e. P<sub>1</sub>'s, P<sub>2</sub>'s, F<sub>1</sub>'s, F<sub>2</sub>'s, BC<sub>1</sub>'s and BC<sub>2</sub>'s for the two crosses (12 entries) were evaluated in the field using a randomized complete blocks design with three replications. Rows were 4 m long, 20 cm width and the space from plant to plant was 10 cm. Each plot consisted of two rows for each P<sub>1</sub>, P<sub>2</sub>, F<sub>1</sub>, BC<sub>1</sub> and BC<sub>2</sub> and five rows for each F<sub>2</sub>.

**Table 1. Name, pedigree and origin of the four parental bread wheat genotypes used in this study.**

Genotype	Pedigree	Origin	Maturity
Line 126	BCH"S"//HORK"S"/4/7C/PATO(B)/3/LR/INI A/BB/5/CNO/GII//BB/INIA/3/NAPO//TOB66/ SPROW"S"	Egypt	Early
Sids 4	Maya"S"Mon"S"/CMH74.A592/3/sakha8*	Egypt	Early
Gemmeiza 9	ALD'S/HUAC'S//CMH74A.630/SX	Egypt	Late
Sunval	COOK*2/VPM-1//3COOK(1345)	Australia	Very late

All agricultural practices were followed according to the recommendations. Data were recorded on 20 individual guarded plants in each  $P_1$ ,  $P_2$  and  $F_1$ , 40 plants in each  $BC_1$  and  $BC_2$  and 100 plants in each  $F_2$  for number of days to 50% heading, number of days to 50% physiological maturity, grain filling period (days), number of spikes/plant, number of kernels/spike, 1000 kernel weight/(g) and grain yield/plant (g).

Heterosis (%) was calculated as the percentage increase of  $F_1$  over the mid- and better parent values. Genetic analysis of generation means and estimates of mean effect ( $\bar{m}$ ), additive (a), dominance (d), additive X additive (aa), additive X dominance (ad) and dominance X dominance (dd) effects were computed using the six-parameter model proposed by Gamble (1962). Heritability in both broad and narrow sense was calculated according to Mather (1949).

## RESULTS AND DISCUSSION

### Generation means and variances

Analysis of variance (not presented) indicated highly significant differences among the six generations for all studied traits of the two wheat crosses. Means ( $\bar{X}$ ) and variances ( $S^2$ ) of the six generations ( $P_1$ ,  $P_2$ ,  $F_1$ ,  $F_2$ ,  $BC_1$  and  $BC_2$ ) of the two studied wheat crosses for earliness and yield traits are presented in Table (2). Significant differences existed between the two parents ( $P_1$  and  $P_2$ ) of each cross for all studied traits. Such differences were more pronounced between parents for the three earliness traits and spikes/plant in both crosses and kernels/spike, 1000 kernel weight and grain yield/plant in cross II. Significant differences between means of parents of each hybrid is a prerequisite for the validity of six-parameter model to determine the magnitude of different gene effects for studied characters.

Means of both studied crosses for  $F_1$ ,  $F_2$ ,  $BC_1$  and  $BC_2$  generations for earliness and yield traits are logic as expected from the plant breeding

Table 2. Means ( $\bar{X}$ ) and variances ( $S^2$ ) of P<sub>1</sub>, P<sub>2</sub>, F<sub>1</sub>, F<sub>2</sub>, BC<sub>1</sub> and BC<sub>2</sub> populations of the two wheat crosses for earliness and yield traits.

Cross	Generation	Days to heading		Days to maturity		Grain filling period (days)	
		$\bar{x}$	$S^2$	$\bar{x}$	$S^2$	$\bar{x}$	$S^2$
<b>Cross I.</b>	P <sub>1</sub>	128	1.88	170	1.67	42	1.00
Sunval	P <sub>2</sub>	76	0.79	146	1.99	70	1.47
X	F <sub>1</sub>	85	1.64	154	0.34	69	1.93
Line # 126	F <sub>2</sub>	88	23.5	150	28.4	62	22.0
	BC <sub>1</sub>	110	22.0	160	20.0	50	20.0
	BC <sub>2</sub>	80	20.0	153	18.0	73	0.71
<b>Cross II.</b>	P <sub>1</sub>	108	0.92	155	1.1	47	1.70
Gemmeiza 9	P <sub>2</sub>	77	1.68	142	2.6	65	2.16
X	F <sub>1</sub>	83	1.20	147	1.5	64	1.93
Sids 4	F <sub>2</sub>	92	45.0	147	33.0	55	37.8
	BC <sub>1</sub>	100	35.0	150	25.0	50	26.7
	BC <sub>2</sub>	85	15.3	144	18.0	59	20.0
<b>LSD 0.05</b>		9		14		6	

Table 2. Cont.

Cross	Generation	Spikes/plant		Kernels/spike		1000-kernel weight(g)		Grain yield/plant(g)	
		$\bar{x}$	$S^2$	$\bar{x}$	$S^2$	$\bar{x}$	$S^2$	$\bar{x}$	$S^2$
<b>Cross I:</b>	P <sub>1</sub>	36	2.64	98	2.80	47	0.15	37	2.40
Sunval	P <sub>2</sub>	16	1.63	95	0.21	50	0.40	30	2.11
X	F <sub>1</sub>	31	1.18	75	0.01	48	0.92	53	1.98
Line #126	F <sub>2</sub>	30	29.7	80	27.0	49	22.0	47	24.0
	BC <sub>1</sub>	32	20.0	83	18.0	48	14.0	52	18.0
	BC <sub>2</sub>	29	18.0	80	19.0	52	16.0	52	16.0
<b>Cross II.</b>	P <sub>1</sub>	18	1.40	71	1.80	42	0.15	53	1.20
Gemmeiza	P <sub>2</sub>	3.2	3.48	93	1.63	54	0.15	22	1.36
9	F <sub>1</sub>	16	4.40	82	1.42	44	0.23	48	1.88
X	F <sub>2</sub>	18	60.0	56	22.0	42	12.0	44	42.0
Sids 4	BC <sub>1</sub>	18	57.0	76	15.0	40	10.0	46	22.0
	BC <sub>2</sub>	11	25.0	87	11.0	50	4.00	33	26.0
<b>LSD 0.05</b>		9		8		15		12	

point of view. Results of Table (2) showed the existence of mid-parent heterosis for days to heading and days to maturity and heterobeltiosis (better-parent heterosis) for grain filling period in both crosses and grain yield/plant in cross I.

The high means of  $BC_1$  for days to heading and days to maturity are due to lateness of the first parent ( $P_1$ ), while the low means of  $BC_2$  are due to the earliness of the second parent ( $P_2$ ) in both studied crosses. Likewise, for spikes/plant in both crosses and grain yield/plant in cross II, the higher means of  $BC_1$  are due to high means of  $P_1$  and the lower means of  $BC_2$  are due to the lower means of  $P_2$ .

The highest magnitude of variance ( $S^2$ ) was reported by the  $F_2$  generation for all studied earliness and yield traits in both crosses (Table 2) followed by that of backcross generations ( $BC_1$ 's and  $BC_2$ 's), while the lowest  $S^2$  magnitude was exhibited by  $P_1$ ,  $P_2$  and  $F_1$  populations, which is also logic from the breeding point of view, due to the homogeneity of such populations.

#### **Gene effects**

Estimates of gene effects calculated from the six-parameter model of the generation mean analysis are presented in Table (3). Significant mean effects ( $m$ ) were exhibited in all studied traits for both crosses.

The occurrence of significant and positive additive ( $a$ ) effects was in 9 out of 14 cases, namely days to heading, days to maturity, spikes/plant and grain yield/plant in both crosses and kernels/spike in cross II. This assures the enhancing effect of additive variance in the inheritance of these traits.

These results indicated that the potentiality of improving the performance of these traits using pedigree selection program may be affective as reported by Abul-Naas *et al* (1993).

The estimates of dominance ( $d$ ) effects were significant and positive in 9 out of 14 cases, namely grain filling period and 1000-kernel weight in both crosses, days to heading, days to maturity, spikes/plant and grain yield/plant in cross I and kernels/spike in cross II.

Additive was larger in magnitude than dominance effect in 6 cases, namely days to heading in both crosses, days to maturity, spikes / plant and grain yield/plant in cross II and kernels/spike in cross I. On the contrary, dominance was larger than additive variance in 7 cases, namely grain filling period and 1000-kernal weight in both crosses, days to maturity and grain yield/plant in cross I and kernels/spike in cross II. Similar conclusions were reported by Bhatt (1972), Avey *et al* (1982) and Menshawy (2005) for days to heading. However, Beiquan and Kronstad (1994) and Menshawy (2004) reported that additive was more important than dominance effect for grain filling period. Differences in conclusions between this study and others and between cross I and cross II with regard to the relative

**Table 3. Mean estimates of the six gene effects for studied traits in two wheat crosses.**

Trait	Cross	Gene action parameter					
		m	a	d	aa	ad	dd
Days to heading	I	88**	30**	15**	28**	4.30**	-33**
	II	92**	15**	-7.5	2.0*	-0.50	-21**
Days to maturity	I	150**	7**	10.65**	26**	-4.0**	-27**
	II	147**	6**	-7.50**	0.00	-0.5	3.0*
Grain filling period	I	49**	-23**	35.50**	22**	-9.0**	-19**
	II	48**	-2*	20**	12**	7.0*	-4.0*
Spikes /plant	I	30**	3.0*	3.0*	3.1*	1.5	-12.0**
	II	18**	7.0*	-8.6*	-14.2**	-0.4	9.20**
Kernels /spike	I	80**	3.0*	-15**	6.0*	1.5	11.0**
	II	56**	-11.0**	102**	102**	0.00	-10.0**
1000 kernel weight	I	49**	-4.0*	3.5*	4.0*	-2.50	-11.0**
	II	42**	-9.0**	6.7*	10.6**	-3.30*	-7.89*
Grain yield /plant	I	47**	5.0*	29.5**	10.0**	1.50	-35.0**
	II	44**	13.0**	-7.95**	-18.0**	-2.95*	31.9**

\* & \*\* indicate significance at 0.05 and 0.01 levels of probability, respectively.  
m = mean effect, a= additive effect and d= dominance effect.

importance of either additive or dominance effects may be due to the differences of the genetic background of materials used in these studies.

Significant digenic epistatic gene effects (Table 3) were exhibited in most studied cases for all the three types of epistasis (aa, ad, and dd). This indicates that epistatic gene effects were generally important in the inheritance of studied traits. Additive X additive (aa) gene effects were generally higher in magnitude than ad and dd gene effects. Significant and positive aa effects were exhibited in 7 cases, i.e grain filling period in both crosses days to heading, days to maturity and grain yield/plant in cross I and

kernels/spike and 1000-kernel weight in cross II. This result suggests an enhancing effect of additive X additive type of epistasis for inheritance of these traits. This component (aa) plus the additive (a) one are amenable types of gene action for more efficient selection.

The additive X dominance type of epistasis exhibited significant and positive effects in only two cases, namely days to heading in cross I and grain filling duration in cross II (Table 3).

Significant and positive dominance X dominance epistatic gene effects were shown in 4 cases, namely days to maturity, spikes/plant and grain yield/plant in cross II and kernels/spike in cross I.

### **Heterosis, heritability and genetic advance**

Favorable (desirable) heterosis percentages in the present study were considered those with negative sign for days to heading and days to maturity and those of positive one for grain filling period and all studied yield traits.

Significant and highly significant desirable percentages of heterosis relative to mid- parent were exhibited for days to heading, grain filling period, spikes/plant and grain yield/plant in both crosses (Table 4). Only grain yield/plant of cross No. 1 showed highly significant desirable heterosis relative to the better parent (heterobeltosis) of 43.24 %. Mid- parent heterosis of grain yield/plant was 58.2 % for cross I and 28.0 % for cross II.

Significant favorable heterosis was also reported by many investigators (El- Borhamy 2000 Salgotra *et al* 2002 and Al- Naggat *et al* 2007 for days to heading and to maturity, Essa *et al* 1994 and El- Maghraby 1998 for spikes / plant and Walia *et al* 1993, El- Sherbeny *et al* 2000 Awaad 2002 and Al-Naggat *et al* 2007 for grain yield and its components).

Very high broad- sense heritability percentages (> 90 %) were obtained in all studied characters for both crosses (Table 4), indicating that genetic variance was accounted for most of the phenotypic variance. However, narrow- sense heritability percentage was the lowest ( 21.22 % ) for days to heading, medium (39.12 %) for grain filling period and the highest (88.02 %) for days to heading; all in cross I (Table 4). The difference in magnitude between broad- and narrow- sense heritability is attributed to the non- additive genetic effects, i.e dominance, additive X dominance and dominance X dominance effects since narrow- sense heritability is based on additive effects which include additive and additive X additive heritability variance. Such differences were very high for days to heading and grain filling period in the first cross, suggesting that non-additive components accounted for the largest part of genetic variance for these cases. Heterosis breeding is a recommended procedure to utilize such non - additive components. On the contrary, in the second cross, days to

**Table 4. The heterobeltosis %, heritability % and genetic advance from selection (in absolute units and percentage) for studied traits in two wheat crosses**

Trait	Cross	Heterosis %		Heritability %		Genetic absolute	advance %
		BP	MP	Broad	Narrow		
Days to heading	I	11.82*	-16.7*	93.87	21.22	2.11	2.40
	II	7.79	-10.3*	97.18	88.02	12.16	13.20
Days to maturity	I	5.80	-2.5	96.47	66.09	7.25	4.83
	II	3.52	-1.0	94.77	69.69	8.24	8.24
Grain filling period	I	-1.40	10.7*	93.34	39.12	4.07	8.32
	II	-1.50	14.3*	94.90	76.45	9.69	20.19
Spikes /plant	I	-13.89*	19.2*	93.89	72.22	8.11	27.04
	II	-11.11*	52.4**	94.80	62.54	9.97	55.44
kernels /spike	I	-23.47**	-22.3*	92.27	62.96	6.74	8.42
	II	-11.83*	0.0	92.64	81.81	7.90	14.12
1000 kernel weight	I	-4.00	1.0	97.77	63.63	6.14	12.54
	II	18.33*	8.3	98.53	83.33	5.94	13.92
Grain yield/plant	I	43.24**	58.2**	90.98	58.33	5.88	12.52
	II	-9.43	28.0**	96.47	85.71	11.44	26.00

\* & \*\* indicate significance at 0.05 and 0.01 levels of probability levels, respectively.

heading, kernels/spike, 1000- kernel weight and grain yield/plant traits, which showed heritability in narrow-sense of above 80 %, indicating that selection in the segregating generations could be considered the method of choice.

Percentage of expected genetic advance from selection (Tale 4) was the highest (55.44 %) for number of spikes / plant in cross II, followed by (27.04 %) for spikes/plant in cross I and (26.00 %) for grain yield in cross II and the lowest (2.40 %) for days to heading in cross I. High percentage of expected genetic advance would help breeder in improving the trait of interest via a few cycles of selection. This conclusion was supported by many investigators (Shehab E- Din 1997, Abd El- Aty and Katta 2002, Menshawy 2005 and Al- Naggar *et al* 2007). This is applicable for number of spikes/plant in both crosses, grain filling period and grain yield/plant in the second cross.



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## التحليل الوراثى لصفات التبكير و صفات محصول الحبوب فى هجينين من القمح

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٢- قسم المحاصيل ، كلية الزراعة ، جامعة القاهرة ، الجيزة

٣- قسم بحوث القمح ، معهد بحوث المحاصيل الحقلية ، مركز البحوث الزراعية ، الجيزة

استخدم نموذج المعلمات الستة لتقدير التفاعلات بين اليلات الجين الواحد وبين اليلات الجينات المختلفه التى تتحكم فى وراثه صفات التبكير ومحصول الحبوب فى هجينين من قمح الخبز هما : الهجين الاول : سنفال X سلالة رقم ١٢٦ و الهجين الثانى : جميزة ٩ X سدس ٤. اشار تحليل الاختلاف الى وجود فروق معنويه بين الاجيال الستة ( الاب الاول ، الاب الثانى ، الجيل الهجينى الاول ، الجيل الهجينى الثانى ،الجيل الرجعى لسلاب الاول ،الجيل الرجعى لسلاب الثانى ) لكل الصفات المدروسة فى كلا الهجينين. ظهرت تأثيرات معنويه و موجبه للتباين المضيف فى صفات عدد الايام حتى طرد السنابل ، عدد الايام حتى النضج ، عدد سنابل النبات ومحصول حبوب النبات فى كلا الهجينين ، ويؤكد ذلك التأثير التعظيمى للتباين المضيف فى وراثه هذه الصفات . كانت تأثيرات السيادة معنويه وموجبه لفترة امتلاء الحبوب ووزن الـ ١٠٠٠ حبه فى كلا الهجينين وعدد الايام حتى الطرد وعدد الايام حتى النضج وعدد سنابل النبات ومحصول حبوب النبات فى الهجين الاول . كان مقدار التباين

المضيف اكبر من تباين السيادة بالنسبة لعدد الايام حتى الطرد فى كلا الهجينين ، عدد الايام حتى النضج ، عدد سنابل النبات ومحصول حبوب النبات فى الهجين الثانى ، كانت تأثيرات التفاعل بين اليلات الجينات المختلفة هلمه بصفه علمه فى وراثه الصفات تحت الدراسه ، كانت تأثيرات التفاعل من نوع التباين المضيف اعلى بصفه عامه من التفاعلات للتفوقيه من نوع ( التباين المضيف X تباين السيادة ) ، و( تباين السيادة X تباين السيادة ) ، ظهرت تأثيرات معنويه وموجبه لتفاعل التفوق من النوع ( المضيف X المضيف ) فى صفه فتره امتلاء الحبوب فى كلا الهجينين وعدد الايام حتى الطرد وحتى النضج ومحصول حبوب النبات فى الهجين الاول ، وظهرت تأثيرات معنويه وموجبه للتفاعل من النوع ( المضيف / السيادة ) فى حالتين فقط هما عدد الايام حتى الطرد فى الهجين الاول وفتره امتلاء الحبوب فى الهجين الثانى ، كما ظهرت تأثيرات معنويه وموجبه للتفاعل من نوع ( السيادة X السيادة ) فى صفات عدد الايام حتى النضج وعدد سنابل النبات ومحصول حبوب النبات فى الهجين الثانى . اشارت النتائج الى وجود قيم معنويه وعاليه المعنويه مرغوبه لقوه الهجين بالنسبه لمتوسط الابوين فى صفات عدد الايام حتى الطرد ، فتره امتلاء الحبوب ، وعدد سنابل النبات ومحصول حبوب النبات فى كلا الهجينين بينما ظهرت قوه هجين عاليه المعنويه مرغوبه بالنسبه للاب الاصن فى حاله واحد فقط وهى محصول حبوب النبات فى الهجين الاول وقدرها ٤٣.٢٤% ، تم الحصول على نسب عاليه جدا لكفاءه التوريت العلمه (تزيد عن ٩٠%) فى كل الصفات المدروسه لكلا الهجينين ، كانت قيم كفاءه التوريت الخاصه اقل ما يمكن (٢١.٢٢%) لعدد الايام حتى الطرد ومتوسطه (٣٩.١٢%) لفتره امتلاء الحبوب ، واعلى ما يمكن (٨٨.٠٢%) لعدد الايام حتى الطرد فى الهجين الاول ، كان التقدم الوراثى المتوقع بالانتخاب اعلى ما يمكن (٥٥.٤٤%) لعدد سنابل النبات فى الهجين الثانى واقل ما يمكن (٢٠.٤٠%) لعدد الايام حتى الطرد فى الهجين الاول .

المجلة المصرية لتربية النبات ١٣ : ٣٧١-٣٨١ (٢٠٠٩)