Heterosis and combining ability in F_1 and F_2 of faba bean (Vicia faba L.)

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Abstract:

This investigation was carried out under the screenhouse at New Valley research station during 2009/2010 to 2011/2012 seasons, to determine heterosis and general (GCA) and specific combining ability (SCA) for yield and yield attributes in fiveparental diallel crosses. Five faba bean genotypes (Vicia faba L.) were used as parental lines. Highly significant differences among the tested entries were detected for different traits, indicating wide genetic variability for all traits. Several crosses recorded significant positive heterosis percentages relative to mid parent (MP) and better parent (BP) for number of pods/plant, 100-seed weight and seed yield/plant ranged from 6.3 to 42.4%, 2.8 to 23.0%, 3.1 to 12.8%, 3.2 to 6.6%, and 4.2 to 30.4%, 13.2 to 22.6% relative to MP and BP for each character, respectively. The ratios (GCA/SCA) exceeded the unity for all studied traits indicating that the genetic variation among these traits in the F_1 and F_2 appeared to be additive. The parental genotype, P₄ (1706B/87/2003) was found to be a good combiner for 100-seed weight and seed yield/plant and P₅

(1706A/400/2003)expressed highly significant positive GCA effect for 100-seed weight. Moreover, one parent Wadi 1was good combiners for earliness. Five F_1 's $(P_1xP_2, P_1xP_3, P_2xP_4,$ P_2xP_5 and P_4xP_5) and only one F_2 (P₁xP₃) had significant positive (SCA) effects for 100 seed weight. Five F₁'s (P₁xP₃, P₁xP₄, P_2xP_4 , P_2xP_5 and P_4xP_5) as well as three F₂'s (P₁xP₃, P₂xP₄ and P₂xP₅) had significant positive (SCA) effects for seed yield plant, indicating that the crosses may be of specific interest in hybrid faba bean breeding programe.

Introduction:

Faba bean (Vicia faba L.) is the important legume crop in the world. It is considered the popular diet in Egypt. The increasing gap between consumption and production necessitates increasing the production of faba bean in Egypt. This could be achieved through extending the cultivated area beside the productivity of unit area by adopting an efficient breeding program. Success of any plant breeding program depends largely upon a better understand of the genetic basis of the important economic traits. Information about heterosis, general and specific combining

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ability and the type of gene action may help faba bean breeder to formulate the most efficient breeding procedure for achievement of maximum genetic improvement among a particular set of genotypes. Superiority of hybrids over the mid and better parents for seed yield was found to be associated with manifestations of heterotic effects in main vield components i.e., number branches, number of pods, number of seeds plant and seed index. The effects ranged from significant positive to significant negative for different traits and were very pronounced in F₁ of faba beans especially crosses among widely divergent materials and less occurred in hybrids between local varieties (Duc. 1997: Schill et al., 1998; Bond and Crofton, 1999; Filippetti et al., 1999; Darwish et al., 2005; El-Hady et al., 2006, 2007; Gasim & Link, 2007; Alghamdi, 2009; Salem, 2009; Hossam, 2010 and Farag & Afiah, 2012). Several researchers found the significance of both general and specific combing ability effects for yield and other important traits of faba beans (Abdalla et al., 2001; Attia et al., 2002; Attia & Salem, 2006; Salem. 2009; Alghamdi, 2009; Hossam, 2010 and Farag & 2012). Salem Afiah. (2009)found that the genetic variation among number of pods plant, number of pods main stem and 100-seed weight appeared to be additive. However the predominance of non-additive gene action were found for days to flowering, plant height, number of branches plant, number of seeds plant and seed vield plant. The present investigation aimed to understand the nature of gene action and relative magnitude of heterosis and combining ability of five faba bean genotypes and their F₁ and F₂ generations using diallel cross mating design

Materials and methods

This investigation was carried out under the screenhouse at New Valley Research Station to determine the heterosis for yield and yield attributes in five—parental diallel crosses, the effective of general combining ability (GCA) and specific combining ability (SCA) to choose the most effective breeding method. Five faba bean genotypes (*Vicia faba* L.) were used as parental lines in the present study

Table 1: Pedigree and characteristics of some faba bean parental geno-

types used in the present study

Genotypes	Pedigree	Flower-	Seed	Seed	Hilum
		ing	size	color	
P ₁ (Giza 40)	An individual plant selected from Rebaya 40	Early	Medium	Beige	Black
P ₂ (Wadi 1)	Derived from Giza blanca × Triple white	Early	Medium	Beige	White
P ₃ (Giza 429)	An individual plant selected from Reba- ya 402	Early	Medium	Beige	Black
P ₄ (1706B/87/2003)	Derived from Giza blanca × Triple white	Late	Large	Greenish	White
P ₅ (1706A/400/2003)	Derived from Giza 674 × Giza blanca	Early	Large	Beige	Black

A diallel cross set involving the five parents were grown screenhouse in 2009/2010 season. The ten possible F₁ crosses with their respective parents were sown in 2010/2011 season. In 2011/2012 season, the five parents genotypes, $10 \, \text{F}_1$ crosses and $10 \, \text{F}_2$ populations were evaluated in a randomized complete block design with three replications. Each plot consisted of one ridge for parents and F₁ crosses and five ridges in F₂ populations of three meters in length and 60 cm. width. Hills were spaced by 20 cm with one seed per hill in one side of the ridge. Data were recorded on individual plant basis, therefore, 10 guarded plants for parent and F₁'s and 50 guarded pl ants in F₂ were randomly chosen from each plot. The following traits were recorded: days to flowering, maturity, number of branches plant, number of pods plant, number of seeds plant, 100 seed weight and seed vield

plant. The heterotic effects of F_1 crosses were estimated as percentage of mid parent and better parent. General and specific combining ability estimated as according to Griffing (1956) diallel cross analysis designated as method 2 model 1.

Results and discussions

Mean squares of genotypes in the F_1 and F_2 generations revealed highly significant differences among the tested genotypes for all characters under investigation which indicate a wide genetic variability for the studied characters and hence, the feasibility for genetic improvements in these materials (Table 2)

The mean performance of some faba bean traits presented in Table 3 showed that parent P_2 (Wadi 1) was the earliest and recorded 33.0 and 118 for days to flowering and maturity, respectively. On the other hand, the latest parent with regard to flowering was P_4 (1706B/87/2003) which recorded 50.7 days, but the

latest one for maturity was P₃ (Giza 429) which registered 139 days. With respect to number branches plant, results showed that P₄ (1706B/87/2003) followed by P₅ (1706A/400/2003) had the highest number of branches plant, 4.3 and 4.0, respectively. Meanwhile, the lowest mean value (3.5) in this respect was recorded from P2 (Wadi 1). Giza 40 produced the highest pods number/plant (31.8) while the lowest value (24.9) produced from P_5 (1706A/400/2003). With regard to number of seeds/plant, data showed that the (1706B/87/2003) followed by P₁ (Giza 40) had the highest number of seeds/plant, 89.9 and 88.5, respectively. The highest of seed index (112 g) was obtained from P₅ (1706A/400/2003) followed by P₄ (1706B/87/2003) which registered 107 g, while the lowest mean value (65 g) was obtained from P₁ (Giza 40). Also, the presented data showed

Table 2: Mean squares and combining ability for different studied

traits in the F_1 and F_2 generations.

SOV	d.f	Flowering		Mat	urity	brar	. of nch- lant	No. of pods/plant	
		F_1	F_2	F_1	F_2	F ₁	F ₂	F ₁	F ₂
Rep.	2	1.67	2.12	0.73	4.95	0.01	0.15	0.83	6.21
Geno.	14	86.14* 82.08*		85.17* *	85.20* *	0.57* 0.30		44.52*	32.38*
Par- ents	4	135.57**		230.71**		0.29		19.06	
Cross es	9	73.74*	67.37* *	23.55*	12.14*	0.32*	0.26	44.62*	39.26*
P vs C	1	0.10	0.54	57.55* *	160.61 **	3.93*	0.75	145.44 **	23.70*
G.C.V.	4	270.05	261.62	216.40	199.52 **	0.63*	0.48	76.77* *	68.03*
S.C.A.	10	12.58*	10.26*	32.68*	39.46*	0.55*	0.23	31.62*	18.11
Error	28	0.67	2.95	1.25	4.18	0.06	0.25	1.79	11.57
GCA/SCA		21.47	25.50	6.62	5.05	1.14	2.09	2.43	3.76

^{*} and ** Significant at 0.05 and 0.01 levels of probability, respectively.

Table 2: Cont.

SOV	d.f	No. of see	eds/plant	Seed yie	eld/plant	100 seed weight		
30 v		F_1	F_2	F_1	F_2	F_1	F_2	
Rep.	2	7.29 44.34		44.34 1.29 3.71		10.46	37.98	
Geno.	14	279.61*	205.82*	794.98* *	683.66* *	927.79*	814.76* *	
Par- ents	4	163.10**		780.54**		1428.83**		
Cross es	9	233.19*	181.77*	741.81*	620.90*	793.33* *	631.16*	
P vs	1	1663.38	593.11*	1331.33	860.94*	133.74*	10.82**	

С		**	*	**	*	*	
G.C.V.	4	451.08* *	375.17*	2078.79	1627.37 **	2909.68	2538.55 **
S.C.A.	10	211.02*	138.08*	281.45*	306.17*	135.02*	125.25*
Error	28	8.18	25.28	4.89	18.96	3.57	11.38
GCA/SC A		2.14	2.72	7.39	5.32	21.55	20.27

^{*} and ** Significant at 0.05 and 0.01 levels of probability, respectively.

Table 3:Mean performance of five faba bean parental genotypes and their F_1 and F_2 generations for yield and some attributes.

and then 11 and 12 generations for yield and some attributes.															
Traits									Num				Seed yield		
	Flov	wer-	Maturi-			er	C	of	0	f	Weight		Plant		
Geno-	ir	ıg	t	y	of		pods		seed						
types						branch		Plant		nt					
					e	S									
					pla	ant									
P_1	36	5.3	13	35	3.	.7	31	.8	88	88.5		5.0	57.0		
P_2	33	3.0	11	18	3.	.5	26	5.9	77	.6	94	0	73.0		
P ₃	37	7.7	13	39	3.	.7	28	3.1	81	.1	69	0.0	56.0		
P_4	50).7	13	37	4.	.3	28	3.4	89	.9	10'	7.0	93	0.0	
P_5	38	3.7	13	37	4.	.0	24	1.9	72	.4	112	2.0	81	.0	
Crosses	F_1	F_2	F_1	F_2	F_1	F_2	F_1	F_2	F_1	F_2	F_1	F_2	F_1	F_2	
$P_1 \times P_2$	35.	35.	13	13	4.2	4.1	38.	36.	98.7	96.	98.8	81.4	80.2	75.4	
	3	3	0	4			8	7		6					
$P_1 \times P_3$	33.	33.	13	13	4.0	3.9	34.	31.	104.	97.	75.4	74.1	73.2	70.5	
	3	7	5	9			8	4	7	0					
$P_1 \times P_4$	43.	44.	13	13	4.7	3.9	34.	31.	100.	95.	90.0	80.6	93.7	77.3	
	7	7	6	7			5	2	2	8					
$P_1 \times P_5$	37.	38.	13	13	4.2	3.9	30.	29.	96.2	93.	91.8	78.3	81.1	73.0	
	3	3	4	9			4	7		4					
$P_2 \times P_3$	34.	35.	13	13	4.6	4.0	30.	27.	84.7	83.	82.2	80.0	71.3	67.4	
	7	7	6	8			1	9		5					
$P_2 \times P_4$	47.	44.	13	13	4.7	4.5	35.	32.	94.8	93.	111.	108.	109.	106.	
	0	3	5	5			0	1		3	1	6	0	3	
$P_2 \times P_5$	38.	37.	13	13	4.7	4.3	25.	23.	76.1	73.	118.	111.	96.8	92.5	
	7	0	6	7			5	1		7	1	5			
$P_3 \times P_4$	45.	45.	14	14	4.6	4.1	30.	28.	86.8	83.	83.1	76.8	75.1	64.1	
	3	0	0	0			1	7		5					
$P_3 \times P_5$	35.	33.	13	13	4.0	3.7	28.	27.	86.5	84.	85.6	79.9	74.3	68.1	
	3	3	7	8			9	5		9					
$P_4 \times P_5$	43.	43.	14	14	5.0	4.6	30.	26.	98.4	94.	117.	108.	107.	101.	
	0	0	0	0			2	8		6	7	2	6	3	
L.S.D.															
0.05	1.4		1.	.9	0.	.4	2	.2	4.	8	3.	.1	3.	.7	
for P+F ₁															
L.S.D.								_							
0.05	2	.9	3.	.4	0.6		5	.7	8.	8.4		5.6		7.3	
for P+F ₂]														

that the highest seed yield plant (93g) was obtained from P₄ (1076 B/87/2003) followed by P₅ (1706A/400/2003) which yielded 81 g. On the other hand, the lowest value in this respect was obtained from

parental genotype P₃ (Giza 429) with no significant difference from the value registered from parental genotype P₁ (Giza 40) which gained 57g plant. The performance of the crosses indicated that none of

the crosses was earliest than the earliest parent for days to flowering and maturity in both F₁ and F₂ generations. However, six $(P_1xP_4, P_2xP_3, P_2xP_4,$ P₂xP₅, P₃xP₄ and P₄xP₅), four $(P_1 \times P_2,$ P_1xP_3 P_1xP_4 and P_2xP_4), six (P_1xP_2 , P_1xP_3 , P_2xP_4 P_1xP_4 $P_1 \times P_5$ and P_4xP_5), two $(P_2xP_5 \text{ and } P_4xP_5)$ and four $(P_1xP_4, P_2xP_4, P_2xP_5)$ and P₄xP₅) F₁ crosses. While, F_2 generations six $(P_1 x P_2,$ P_1xP_3 , P_2xP_3 , P_2xP_4 , P_2xP_5 and P_4xP_5), two $(P_1xP_2 \text{ and } P_2xP_4)$, $six (P_1xP_2, P_1xP_3, P_1xP_4,$ P_1xP_5 , P_2xP_4 and P_4xP_5), two (P_2xP_4, P_2xP_5) and five $(P_1xP_2,$ P_1xP_3 , P_2xP_4 , P_2xP_5 and P_4xP_5) populations exceed the highest parental genotypes for number of branches plant, number of pods plant, number of seeds plant, 100 seed weight and seed yield plant, respectively. It could be concluded that the above mentioned parents and crosses would prospect in faba bean breeding and therefore may be valuable for improving seed yield via its component characters.

Heterosis effect

Values of heterosis percentage relative to mid parent (MP) and Better parents (BP) were significantly differed between crosses (Table 4). Heterosis percentage relative mid parent (MP) were ranged from -9.9 to 12.3, -1.7 to 6.2, 0.8 to 27.8, -1.4 to 32.4. 1.4 to 23.4. -29.3 to 17.8 and -1.2 to 30.4 for days to flowering, days to maturity, number of branches, pods and seeds per plant, 100 seed weight and seed yield per plant, respectively. However, heterosis percentages relative to the better parent (BP) ranged from -8.3 to 42.4, -1.2 to 15.6. 0.8 to 23.2. -5.1 to 23.0. -3.5 to 18.3, -29.3 to 6.6 and -20.7 to 22.6 for days to flowering, days to maturity, number of branches, pods and seeds per plant, 100-seed weight and seed vield/plant. respectively. The highest values of heterosis percentage relative to the mid parent (MP) in the F₁ crosses were registered from crosses (P₂xP₄), (P_2xP_5) , (P_2xP_3) , (P_1xP_2) , (P_1xP_3) , (P_1xP_2) and (P_2xP_4) with regard to days to flowering, days to maturity, number of branches, pods and seeds per plant, 100-seed weight and seed vield per plant traits, respectively. While, the highest values of heterosis percentage relative to the better parent (BP) in the F₁ generations were registered from crosses (P_2xP_4) , (P_2xP_3) , (P_2xP_3) , (P_2xP_4) , (P_1xP_3) , (P_1xP_3) and (P_1xP_3) for the same previous traits in the same order. It could be suggested that the heterotic effects for seed vield was associated with other components

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Furthermore, various cross combinations exhibited different degrees of F₁ superiority in some traits based on the genes in parental combinations that may contribute directly or indirectly to the traits. Varied values of heterosis might be due to the genetic diversity of the parents with nonallelic interaction which increase or decrease the expression of heterosis (Hayman, 1958). The heterosis estimates for the majority of the traits showed that there was sufficient genetic divergence among the parents assessed, resulting in a favorable situation for breeding (Barelli et al., 2000). Even in the absence of epistasis, multiple alleles at a locus lead to either positive or negative heterosis (Cress, 1966). Pronounced and favourable heterosis have been obtained by several researchers for faba bean traits which varied according to the cross combinations and traits (Duc, 1997; Stelling, 1997; Schill et al., 1998; Abdulmula et al., 1999; Bond and Crofton, 1999; Filippetti et al., 1999; Abdalla et al., 2001; Attia et al., 2002; Darwish et al., 2005; Attia & Salem, 2006; El-Hady et al., 2006; Gasim and Link, 2007; Elghamdi, 2009; Salem, 2009; Hossam, 2010 and Farga & Afiah, 2012).

Combining ability

Mean squares of both GCA and SCA estimates were highly significant in both generations for all the studied traits except number of branches/plant in the F₂ generation (Table 2). Also, the ratio of GCA/SCA estimates exceeded the unity for all studied

traits in both F₁ and F₂ generations. This indicates that most of the genetic variation among the investigated genotypes for the mentioned traits appears to be under additive gene actions. A direct selection could thus be useful for improving these traits. Comparisons between GCA effects associated with each parent (Table 5), revealed that the parent P₁, P₂ and P₃ showed significant negative effects for days to flowering and could be considered as sources for earliness in breeding program, whereas, P₄ parent showed highly significant positive effect in this respect. P₂ parent had a highly negative effect for days to maturity and could be considered as a source for earliness in breeding program. These findings are highly important for breeding early faba bean cultivars. Concerning number of pods and seeds plant traits, P₁ parent showed significant effect in this respect. On the other hand, the tow parents $(P_4 \text{ and } P_5)$ had a highly significant positive effect on 100 seed weight and also P₂ parent showed significant positive effect in this respect. P₄ parent showed a highly significant effect on seed yield plant, while P₅ parent showed significant positive effect in this respect. These results suggest that the mentioned parental genotypes were good combiners for improving most studied traits. The significant relation between combining ability results and the mean performance of parental genotypes indicates the efficiency of phenotypic performance for

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Table 5: General combining ability effects in F_1 and F_2 generations for various traits.

Parents		P_1	P_2	P_3	P_4	P_5	S.E
Days to	F_1	-1.95*	-2.05*	-1.71*	6.33**	-0.62	0.82
Flowering	F_2	-1.43	-2.33	-1.67	6.24**	-0.81	1.72
Days to	F_1	-0.47	-5.40**	2.35*	1.99	1.53	1.12
Maturity	F_2	0.34	-5.38**	2.15	1.12	1.75	2.05
Number of	F_1	-0.15	-0.06	-0.12	0.28	0.05	0.24
branches plant	F_2	-0.12	-0.03	-0.14	0.23	0.06	0.50
Number of pods	F_1	2.68*	-0.04	-0.45	0.46	-2.65*	1.34
Plant	F_2	2.61	-0.05	-0.03	0.21	-2.44	3.40
Number of seed	F_1	6.01*	-3.58	-1.38	3.63	-4.68	2.86
Plant	F_2	5.35	-2.87	-1.58	3.53	-4.43	5.03
100 seed	F_1	-11.18**	4.80*	-13.99**	9.04**	11.34**	1.89
Weight	F_2	-12.78**	5.29	-10.90**	7.79*	10.61**	3.37
Seed yield	F_1	-6.50**	1.38	-12.56**	13.17**	4.51*	2.21
plant	F_2	-7.64	4.14	-11.00*	9.26*	5.24	4.35

^{*, **} significant at 0.05 and 0.01 level of probability, respectively

detecting the potentiality of parents for inclusion in cross breeding programs. Three F₁'s (P₁xP₃, P₃xP₅ and P₄xP₅) had significant negative effect for days to flowering (Table 6). Moreover tow F_1 's $(P_1xP_2 \text{ and } P_2xP_4)$ showed highly significant positive effects for number of pods plant. With respect to number of seeds plant, four F_1 's $(P_1xP_3, P_1xP_5, P_2xP_4)$ and P₄xP₅) had significant positive effects. Five F_1 's $(P_1 \times P_2,$ P_1xP_3 , P_2xP_4 , P_2xP_5 and P_4xP_5) and only one F₂ (P₁xP₃) had significant positive effects for 100 seed weight. Five F₁'s (P₁xP₃, P_1xP_4 , P_2xP_4 , P_2xP_5 and P_4xP_5) as well as three F₂'s (P₁xP₃, P₂xP₄ and P₂xP₅) had significant positive effects for seed vield plant. Thus SCA for seed yield per plant seemed to be influenced by SCA for yield components.

The obtained results clear that some yield components are more important for yield expression than others. GCA effects seemed to provide appropriate criterion for detecting the validity of a line in hybrid combination (or synthetic variety) but SCA effects may be related to heterosis (Peng and Virmani, 1999). In a cross showing high SCA, it might include only one good combiner. such combinations would show desirable transgressive segregations, providing that the additive genetic system present in the crosses are acting in the same direction to reduce underisible plant characteristics and maximize the characters in view (Abdalla et al., 1999). Therefore,

most of the earlier crosses may be of importance in traditional breeding programs. These results are in full agreement with Abdalla *et al.* (2001), Attia *et al.* (2002), Zeid (2003), Darwish *et al.* (2005), Attia & Salem (2006), El-Hady *et al.* (2006), El-Hady *et al.* (2007), Elghamdi (2009), Salem (2009), Hossam (2010) and Farag & Afiah (2012).

Conclusion

The obtained results are highly promising to breed faba bean cultivars, hybrids or synthetics possessing genetic factors for earliness and high yield potential.

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قوة الهجين والقدرة علي الائتلاف في الجيل الأول والثاني في الفول البلدي خالد مجد عجد يماني خالد عجد عجد يماني

قسم بحوث المحاصيل البقولية – معهد بحوث المحاصيل الحقلية – مركز البحوث الزراعية – الجيزة

أجري هذا البحث تحت الصوبة السلكية في محطة بحوث الوادي الجديد خلال المواسم ٢٠١٠/٢٠١٩ و ٢٠١٠/٢٠١١ بهدف دراسة قوة الهجين والقدرة علي الائتلاف في الجيل الأول والثاني في الفول البلدي لصفات المحصول ومكوناته وقد أستخدم لهذا الغرض خمسة تراكيب وراثية من الفول البلدي ذات أصول وراثية متباعدة وأجريت الهجن التبادلية بين الآباء. تم تقييم سلوك الآباء والهجن في الجيل الأول والثاني بالإضافة لتقدير تأثيرات قوة الهجين علي أساس متوسط الأبوين والأب الأعلى والقدرة العامة والخاصة على الائتلاف. وكانت النتائج المتحصل عليها كالآتي:

- كان التباين الراجع للتراكيب الوراثية والآباء والهجن عالي المعنوية لكل من الجيل الأول والثاني لكل الصفات.
- سجلت بعض الهجن قيماً معنوية موجبة لقوة الهجين سواء المحسوبة علي أساس متوسط الأبوين أو الأب الأعلى محصولاً لصفات عدد القرون علي النبات ووزن الـ ١٠٠ بذرة ووزن محصول البذور للنبات وتراوحت القيم بين 7.7-3.7% و 7.7-3.7% و 7.7-3.7% و 7.7-7.7% و 7.7-3.7% و 7.7-3.7% بالنسبة لمتوسط الأبوين أو الأب الأعلى محصولاً لكل صفة على التوالى.
- كانت النسبة بين التباين الراجع للقدرة العامة والخاصة على التآلف ذات قيم عالية تفوق الوحدة لكل الصفات في الجيل الأول والثاني.
- أظهرت السلالة الأبوية (1703B/87/2003) قدرة عامة علي الائتلاف معنوية لصفتي وزن الـ ۱۰۰ بذرة ومحصول البذور النبات. كما أعطت السلالة الأبوية (1706A/400/2003) أعلي القيم المعنوية الموجبة لتأثيرات القدرة العامة علي التآلف لصفة وزن الـ ۱۰۰ بذرة. من ناحية ثانية أعطت ثلاثة آباء جيزه ۶۰ ووادي ۱ وجيزه ۲۶ قدرة عامة للتبكير.
- أظهرت خمسة هجن في الجيل الأول $(P_1xP_2, P_1xP_3, P_2xP_4, P_2xP_5, P_4xP_5)$ وهجن واحد فقط في الجيل الثاني (P_1xP_3) تأثيرات موجبة وعالية لقدرة الخاصة لصفة وزن الـ ۱۰۰ بذرة . كما أعطت خمسة هجن في الجيل الأول $(P_1xP_3, P_1xP_4, P_2xP_4, P_2xP_5, P_4xP_5)$ وثلاثة هجن في الجيل الثاني $(P_1xP_3, P_1xP_4, P_2xP_4, P_2xP_4, P_2xP_5)$ تأثيرات موجبة وعالية للقدرة الخاصة علي التآلف لصفة محصول البذور للنبات وهذه يمكن الاستفادة منها في برامج التربية لتحسين محصول البلاي.