

COMBINING ABILITY AND NATURE OF GENE ACTION IN OKRA (*Abelmoschus esculentus* [L.] MOENCH)

El-Gendy, Soher E. A.¹ ; H. A. Obiadalla-Ali² ; E. A. Ibrahim¹ and M. H. Z. Eldekashy³

¹ Vegetable Res. Dept., Horticultural. Res. Inst., ARC, Egypt

² Dept. of Horticulture, Fac. Agric., Sohag University, Sohag, Egypt

³ Dept. of Horticulture, Fac. Agric., Assuit University, Assuit, Egypt

ABSTRACT

Nine parental genotypes of okra were crossed in complete diallel design to study combining ability and nature of gene action for earliness and yield components. Mean squares of genotypes were found to be highly significant for all studied traits, providing evidence for presence of considerable amount of genetic variation among studied genotypes. The results showed that (P₃) and (P₄) were the best general combiners for earliness, while (P₁), (P₂), (P₅) and (P₆) were found to be good general combiners for total yield per plant. The crosses (P₂×P₁), (P₂×P₄), (P₂×P₅) and (P₄×P₆) were the earliest crosses in comparison with the other crosses. Meanwhile, the cross (P₁×P₄) had the highest mean value for fruit diameter, plant height and fruit weight. In addition, the crosses, (P₁×P₂), (P₁×P₆), (P₂×P₄) and (P₄×P₅) had the highest mean values for No. of fruit/plant and total yield /plant. Therefore, these promising crosses among F₁ hybrids and F₁ reciprocal (F_{1r}) combinations could be used for further breeding studies to improve the economic traits in okra. The results revealed that the general combining ability (GCA) and specific combining ability (SCA) mean squares were highly significant for all studied traits. Significant reciprocal effect mean squares were observed for all studied traits, indicating that these traits were controlled by extra-nuclear factors as well as nuclear factors. The results indicated that the magnitude of additive genetic variance (σ^2_A) were positive and lower than those of non additive (σ^2_D) one for most of studied traits, indicating that non additive gene action played a major role in the inheritance of these traits. The broad sense heritability estimates (H_b² %) were more than 70% and larger than their corresponding narrow sense heritability (H_n² %) for all studied traits. However, estimates of narrow sense heritability were 13.9%, 32.4, 40.0, 47.1, 76.8 for earliness, fruit length, fruit weight, plant height and fruit diameter, respectively. The estimates of narrow sense heritability ranged from 11.2 % to 17.3% for total fruit yield per plant and No. of fruit per plant, respectively. It could be concluded that the most studied traits were mainly controlled by non additive effects and cytoplasmic factors. Therefore, the genetic material used in this study could be used for hybridization for producing promising crosses to improve economic traits in okra.

Keywords: General Combining Ability, Specific combining Ability, gene action, earliness, yield, Okra

INTRODUCTION

Okra (*Abelmoschus esculentus* L.) is one of the most important vegetable crops in Egypt. Combining ability of the parents is becoming important in plant breeding, especially in hybrid production. It is useful in connection with the testing and compare the performance of the lines in hybrid combinations. Information on the general and specific combining

abilities will be helpful in the analysis and interpretation of the genetic basis of important traits. GCA and SCA provide a guideline for the nature of gene action involved in the expression of economic traits. The genetic information obtained from this method is considerable use for selecting parental lines and their crosses to develop and release new high yielding genotypes. Ramesh and Singh (1999), El-Gendy and El-Sherbeny (2005) and El-Sherbeny *et al* (2005) found that the magnitudes of additive genetic variance (σ^2_A) were larger than those of non-additive ones (σ^2_D) for most okra economic traits. On the other hand, Dhankhar and Dhankhar (2001), Prakash *et al* (2002) and Solankey and Singh (2010) stated that non additive genetic variance was higher than the additive one for days to flowering, plant height, number of branches, number of pods per plant and pod yield per plant. However, Vagish *et al* (2002), liou *et al* (2002), El-Gendy and El-Diasty (2004) and Singh *et al* (2011) indicated that both additive and non additive gene action involved in the inheritance of days to flowering, number of pods per plants and pod yield per plant.

Hence, the objective of this study was to assess the combing ability of nine genetically divergent lines in a complete diallel analysis to choose suitable breeding program for improving economic traits in okra.

MATERIALS AND METHODS

Nine genetically divergent parent lines of okra were previously created and developed by Soher El-Gendy in 2009 (Elgendy, Soher 2012). These genotypes are: line 1 (P_1), line 2 (P_2), line 3 (P_3), line 4 (P_4), line 5 (P_5), line 6 (P_6), line 7 (P_7), line 8 (P_8) and line 9 (P_9). The present study was conducted at El-Baramoon research Station, Horticulture Research Institute, ARC, Egypt during the summer seasons of 2010 and 2011. In the summer season of 2010, the seeds of nine inbred lines were sown on April and all possible combinations among them were made according to a complete diallel mating design to produce 36 F_1 and 36 F_1 reciprocal (F_{1r}) hybrids. In the summer season of 2011, 72 F_1 hybrids were evaluated in a randomized block design with three replications. Each block contains 72 plots. Each plot was 3 rows, 3.0 m. long and 60 cm. wide. Hills were spaced 30 cm. Apart. All other agricultural practices were applied as recommended for okra production.

Data were recorded on 10 plants chosen at random from each plot for the following traits: Number of days to 50% flowering (No. of DF) ; Plant height (PH cm); Number of fruit/plant (No. of F/P); Fruit Diameter (FD cm); Fruit Length (FL cm); Fruit weight (FW gm) and Total yield per plant (TY/P gm).

Data were subjected to the analysis of variance in order to test the significance of the differences among the 72 F_1 and F_{1r} reciprocal hybrids according to Cochran and Cox (1957).

Sum squares of studied genotypes was partitioned according to Griffing's (1956) as method 3 into sources of variations due to GCA and SCA.

The variances of GCA (σ^2_g) and SCA (σ^2_s) were obtained on the basis of the expected mean squares for all studied traits. Additive (σ^2_A) and non-additive (σ^2_D) genetic variances were estimated according to Matzinger and Kempthorne (1966) as follows:

$$\sigma^2_A = \frac{1}{2} \sigma^2_g$$

$$\sigma^2_D = \sigma^2_s$$

Estimates of heritability in both broad and narrow sense were calculated according to the following equations:

$$h^2_b\% = [(\sigma^2_A + \sigma^2_D) / (\sigma^2_A + \sigma^2_D + \sigma^2_e)] \times 100$$

$$h^2_n\% = [\sigma^2_A / (\sigma^2_A + \sigma^2_D + \sigma^2_e)] \times 100$$

RESULTS AND DISCUSSION

Genotypic variations

Analyses of variance for all genotypes are presented in Table 1 for all studied traits. Mean squares of genotypes were found to be highly significant for all studied traits. This provides evidence for presence of considerable amount of genetic variation among studied genotypes. These results are in harmony with those previously obtained by El-Sherbeny *et al* (2000) and Abdelmageed (2010).

Table 1: Analysis of variance and mean squares of all genotypes for studied traits

SV	DF	No. of DF	PH (cm)	No. of F/P	FD (cm)	FL (cm)	FW (gm)	TY/P (gm)
Reps	2	32,21	12,0	102,0	0,40	1,78	4,84	110030,8**
Geno.	71	20,96**	2937,1**	3032,9**	0,02**	1,13**	1,03**	103112,9**
Error	142	1,40	72,1	34,3	0,00	0,20	0,27	6649,9

Mean performance

Mean performance of the 26 F₁ hybrids for all studied traits are shown in Table 2. The results showed considerable variation were obtained among all F₁ hybrids for all studied traits. The crosses (P₂ × P₁), (P₂ × P₃) and (P₂ × P₄) were the earliest crosses in comparison with the other crosses. Meanwhile, the cross (P₁ × P₃) had the highest mean value for fruit diameter, plant height and fruit weight. In addition, the crosses, (P₁ × P₁), (P₁ × P₂) and (P₂ × P₃) had the highest mean values for No. of fruit/plant and total yield /plant.

Mean performance of the 26 F₁ reciprocal crosses (F_{1r}) for all studied traits are presented in Table 3. No specific reciprocal hybrid showed superiority over other crosses for all studied traits. The best combination for earliness was (P₃ × P₂) with mean of 02,7. The crosses (P₂ × P₁), (P₃ × P₂) and (P₂ × P₁) were the highest combinations for plant height, fruit diameter and fruit weight with mean of 280, 6,17 and 0,97, respectively. Moreover, the cross (P₁ × P₃) was the best for no. of fruit per plant, fruit length and total yield per plant with the mean of 239, 4,07 and 1030,7, respectively. Therefore, these promising crosses among F₁ hybrids and F₁ reciprocal combinations

could be used for further breeding studies to improve the economic traits in okra.

Table 7: Mean performance of F₁ hybrids for all studied traits

Hybrids		No. of	PH	No. of	FD	FL	FW	TY/P
		DF	(cm)	F/P	(cm)	(cm)	(gm)	(gm)
1	P ₁ xP _r	70.7	170	137	0.33	3.43	4.70	720.9
2	P ₁ xP _r	70.7	190	181	0.03	3.03	4.00	724.4
3	P ₁ xP _z	09.7	230	104	4.93	3.83	3.07	370.7
4	P ₁ xP _o	00.7	190	190	0.77	2.77	3.83	728.1
5	P ₁ xP ₁	09.7	220	211	0.40	4.07	0.00	1052.3
6	P ₁ xP _v	71.0	230	180	0.73	4.40	7.00	1080.2
7	P ₁ xP _Δ	71.0	190	114	0.97	4.43	0.40	710.9
8	P ₁ xP ₃	09.3	200	84	7.13	3.73	7.20	022.3
9	P ₁ xP _r	72.0	191	92	0.00	3.03	4.47	410.9
10	P _r xP _z	07.7	218	133	4.97	3.27	3.87	010.7
11	P _r xP _o	08.7	188	190	0.07	3.40	3.83	727.9
12	P _r xP ₁	08.3	238	179	4.70	4.03	4.00	717.0
13	P _r xP _v	08.0	190	149	0.07	4.47	4.03	750.3
14	P _r xP _Δ	07.7	207	178	4.77	3.73	3.70	704.8
15	P _r xP ₃	07.3	217	170	4.83	3.70	3.70	712.8
16	P _r xP _z	08.7	200	177	0.73	3.40	4.73	771.9
17	P _r xP _o	07.0	187	103	0.47	4.10	3.97	707.7
18	P _r xP ₁	07.3	190	178	4.97	7.00	0.40	907.2
19	P _r xP _v	08.7	230	100	0.13	3.03	3.47	020.0
20	P _r xP _Δ	09.7	210	110	0.13	3.37	3.80	417.7
21	P _r xP ₃	07.0	170	104	0.17	3.07	3.80	394.0
22	P _r xP _o	08.7	190	179	0.17	3.07	4.20	710.2
23	P _z xP ₁	08.0	211	148	4.87	0.70	0.07	749.9
24	P _z xP _v	73.3	220	101	0.00	3.90	0.20	784.0
25	P _z xP _Δ	07.7	113	107	0.07	3.47	4.20	700.0
26	P _z xP ₃	00.0	203	109	0.20	3.10	4.23	771.8
27	P _z xP _o	03.0	212	112	4.80	3.70	3.87	432.0
28	P _z xP _v	00.3	180	93	0.83	3.03	0.03	014.0
29	P _z xP _Δ	04.7	121	180	0.27	3.73	4.73	803.0
30	P _z xP ₃	04.3	210	179	0.70	3.20	0.77	1031.9
31	P _z xP _o	73.3	202	102	0.10	3.33	4.17	733.3
32	P _z xP ₁	73.7	223	171	0.03	3.37	4.07	704.9
33	P _z xP _v	73.0	270	180	4.43	3.23	3.33	099.7
34	P _v xP _Δ	72.7	232	142	0.73	3.83	4.93	702.9
35	P _v xP ₃	07.0	240	171	7.00	3.10	0.27	900.7
36	P _Δ xP ₃	08.3	232	140	0.23	3.07	4.87	781.3
LSD	0.00	1.94	13.7	9.0	0.37	0.72	0.83	131.72
LSD	0.01	2.07	18.1	12.0	0.49	0.90	1.10	173.84

Table 2: Mean performance of F₁ reciprocal hybrids (F_{1r}) for all studied traits

Hybrids	No. of DF	PH (cm)	No. of F/P	FD (cm)	FL (cm)	FW (gm)	TY/P (gm)	
1	P _r xP ₁	08,7	223	101	0,90	3,80	3,30	098,0
2	P _r xP ₁	09,7	190	124	0,97	3,03	0,73	087,0
3	P _ε xP ₁	11,3	230	170	0,83	3,37	0,97	1013,7
4	P ₀ xP ₁	04,3	210	148	0,73	3,83	0,87	093,3
5	P _γ xP ₁	04,3	194	49	0,43	3,87	0,87	097,0
6	P _ν xP ₁	08,3	200	148	0,83	3,17	0,27	080,3
7	P _λ xP ₁	09,0	280	190	0,47	3,17	0,37	087,3
8	P _ι xP ₁	06,7	240	109	0,97	2,97	0,77	094,3
9	P _r xP _r	07,7	190	139	0,90	3,97	3,70	014,8
10	P _ε xP _r	09,3	220	171	0,13	0,13	0,13	060,0
11	P ₀ xP _r	07,0	230	140	0,93	0,37	0,10	094,0
12	P _γ xP _r	09,0	230	109	0,40	0,43	3,83	099,4
13	P _ν xP _r	10,0	212	100	0,37	3,77	0,83	090,0
14	P _λ xP _r	08,3	212	109	0,87	3,73	3,33	029,0
15	P _ι xP _r	03,7	210	102	0,10	3,80	3,90	098,0
16	P _ε xP _r	09,3	210	191	0,03	2,97	3,90	040,0
17	P ₀ xP _r	00,7	180	141	0,80	3,77	0,70	048,9
18	P _γ xP _r	09,0	230	107	0,23	0,27	0,77	048,3
19	P _ν xP _r	06,7	170	79	0,07	3,13	0,00	037,0
20	P _λ xP _r	10,3	228	171	0,97	0,07	0,17	083,4
21	P _ι xP _r	00,0	240	120	0,17	3,30	0,77	080,0
22	P ₀ xP _ε	04,7	220	130	0,40	3,70	0,80	094,9
23	P _γ xP _ε	04,0	201	239	0,73	0,07	0,33	1030,7
24	P _ν xP _ε	11,3	270	172	0,97	3,87	0,90	1010,0
25	P _λ xP _ε	07,0	270	172	0,03	3,93	3,73	000,3
26	P _ι xP _ε	07,7	241	100	0,17	3,10	3,97	090,7
27	P _γ xP ₀	09,7	217	100	0,90	3,00	3,83	090,0
28	P _ν xP ₀	09,3	197	100	0,07	3,30	0,13	069,4
29	P _λ xP ₀	13,7	230	193	0,37	0,03	0,90	041,0
30	P _ι xP ₀	07,3	230	173	0,43	2,90	0,43	092,7
31	P _ν xP _γ	06,3	239	131	0,20	3,17	0,80	028,7
32	P _λ xP _γ	06,3	280	179	0,47	3,87	0,93	032,9
33	P _ι xP _γ	12,0	228	103	0,00	3,37	0,20	032,7
34	P _λ xP _ν	00,7	212	178	0,30	3,43	0,07	067,7
35	P _ι xP _ν	12,0	180	74	0,70	2,77	0,33	019,8
36	P _ε xP _λ	11,7	272	79	0,30	3,03	3,90	007,9
LSD	0,00	1,94	13,7	9,0	0,37	0,72	0,83	131,72
	0,01	2,07	18,1	12,0	0,49	0,90	1,10	173,84

Combining ability analysis

Mean squares of general, specific combining ability and reciprocal effects for all studied traits are given in Table 3. The results exhibited that mean squares of general combining ability (GCA), specific combining ability (SCA) and reciprocal effects were highly significant for all studied traits. These results indicate that both GCA and SCA were important in the

inheritance of these traits. However, the magnitudes of GCA were larger than those of SCA for all studied traits pointed out the predominance of the additive gene action. In addition, significant reciprocal effect mean squares were observed for all studied traits, indicating that these traits were controlled by extra-nuclear factors as well as nuclear factors. These results are in agreement with those reported by Prakash *et al* (2002), Rewale *et al* (2003), El-Sherbeny *et al.*, (2005), El-Gendy and El-Sherbeny (2005), Sinthil *et al* (2006) and Singh *et al* (2006).

Table 4: The analysis of variance and mean squares for combining ability analysis

SV	DF	No. of DF	PH (cm)	No. of F/P	FD (cm)	FL (cm)	FW (gm)	TY/P (gm)
GCA	8	9,68**	2.48,2**	1337,4**	0,939**	1,07,0**	1,300**	42848,2**
SCA	27	6,01**	479,7**	766,4**	0,009**	0,309**	0,338**	28940,0**
Reciprocal	36	7,12**	1116,0**	150,0**	0,091**	0,230**	0,449**	36006,6**
Error	142	0,48	24,0	11,4	0,017**	0,066	0,089	2216,6

GCA effects (g_i)

Estimates of general combining ability effects (g_i) of each parent for all studied traits are presented in Table 5. (P₀) was the best general combiner for all studied traits except fruit length, fruit weight and plant length. While (P₁) was good general combiner for fruit diameter, fruit weight and total yield per plant. (P₂) was good general combiner for fruit length. (P₃) was good general combiner for fruit diameter. (P₄) was good general combiner for plant length, number of fruit per plant and total yield. (P₅) was good general combiner for fruit length, plant length and number of fruit per plant. (P₆) was the best general combiner for all studied traits except fruit length, number of fruit per plant and earliness. (P₇) was the best general combiner for plant length and number of fruit per plant. (P₈) was the best general combiner for fruit diameter, plant length and earliness. Generally, the results showed that (P₀) and (P₈) were the best general combiners for earliness, while (P₁), (P₄), (P₆) and (P₇) were found to be good general combiners for total yield per plant.

It could be suggested that these parental genotypes possess favorable genes to improve hybrids for earliness and yield components.

Table 5: Estimates of general combining ability effects (g_i) of each parental lines for all studied traits

Genotypes	No. of DF	PH (cm)	No. of F/P	FD (cm)	FL (cm)	FW (gm)	TY/P (gm)
P ₀	0,434*	-0,12	-2,20**	0,312**	-0,076	0,227**	34,4**
P ₁	-0,138	-7,99**	0,04	-0,373**	0,200**	-0,601**	-88,9**
P ₂	0,029	-16,26**	-8,82**	0,193**	0,019	-0,030	-46,6**
P ₃	-0,114	7,74**	13,68**	-0,090**	0,078	-0,020	63,1**
P ₄	-1,780**	-19,14**	7,90**	0,081*	-0,100	0,020	38,2**
P ₅	0,220	10,76**	6,97**	-0,414**	0,014**	-0,113	14,2
P ₆	1,000**	4,33**	-7,89**	0,260**	-0,191**	0,027**	43,2**
P ₇	0,886**	3,74**	6,04**	-0,042	-0,010	-0,111	16,0
P ₈	-0,042**	11,74**	-16,67**	0,077*	-0,474**	0,006	-73,9**
SE(g _i)	0,170	1,24	0,80	0,033	0,060	0,070	11,9

SCA effects (Sij)

Estimated specific combining ability effects (S_{ij}) of each cross combination for all studied traits are found in Table 6. The results revealed that the cross combination ($P_1 \times P_7$), ($P_1 \times P_8$), ($P_7 \times P_8$), ($P_8 \times P_7$), ($P_8 \times P_1$), ($P_7 \times P_1$), showed desirable negative significant SCA effects for earliness. Moreover, seven, seven, five and twelve out of thirty six crosses exhibited positive SCA effects for fruit diameter (cm), fruit length (cm), fruit weight (gm) and plant height (cm), respectively. Concerning to total yield per plant, fifteen and nine out of the thirty six hybrids were the best yielding crosses for number of fruit per plant, and total yield/plant, respectively.

Table 6: Estimates of specific combining ability effects (S_{ij}) of each cross for all studied traits

Crosses	No. of DF	PH (cm)	No. of F/P	FD (cm)	FL (cm)	FW (gm)	TY/P (gm)
$P_1 \times P_7$	1,00*	-9,89**	-2,90	-0,164*	-0,230	-0,279	-88,8
$P_1 \times P_8$	1,33**	-10,41**	10,41**	-0,097	-0,332*	-0,429**	2,2
$P_7 \times P_8$	1,81**	8,09**	-22,09**	-0,176*	-0,070	-0,039	-70,8*
$P_8 \times P_7$	-2,02**	2,47	10,20**	-0,030	-0,242	-0,000**	-12,4
$P_1 \times P_7$	-2,02**	-20,43**	-22,88**	0,177*	0,106	0,216	-69,1*
$P_1 \times P_8$	-0,14	21,49**	20,98**	-0,130	0,377*	0,276	187,2**
$P_7 \times P_8$	0,31	19,09**	-0,40	0,100	0,218	0,174	0,3
$P_8 \times P_7$	-0,26	-0,91*	-7,73**	0,320**	0,177	0,097**	6,3
$P_7 \times P_1$	1,07**	0,26	-24,38**	0,039	-0,192	0,216	-67,1*
$P_8 \times P_1$	-0,12	2,26	-10,38**	0,177*	-0,301	0,123	-49,1
$P_7 \times P_8$	1,38**	18,97**	8,41**	0,201*	0,070	0,040	30,7
$P_8 \times P_7$	0,21	9,23**	13,24**	-0,004	0,296	0,128	72,2*
$P_7 \times P_1$	-0,24	-9,84**	8,70**	-0,011	0,380**	0,204	80,4**
$P_8 \times P_1$	-1,62**	-3,74	8,27**	-0,109*	-0,270	-0,224	-26,1
$P_7 \times P_8$	-2,19**	-7,24*	3,98	-0,078	0,201	-0,108	2,7
$P_8 \times P_7$	0,71	1,73	20,98**	0,143	-0,087**	-0,177	76,0**
$P_7 \times P_8$	-0,29	2,11	-0,23	0,017	0,347*	-0,210	-29,3
$P_8 \times P_7$	-0,40	-3,79	16,20**	-0,021	0,927**	0,728**	194,7**
$P_7 \times P_1$	-1,74**	-2,36	-16,90**	-0,190*	-0,418**	-0,012**	-183,8**
$P_8 \times P_1$	0,71	17,23**	-0,37**	0,008	0,039	0,126	14,7
$P_7 \times P_1$	-1,87**	-4,77	-10,76**	0,000	0,210	0,209	-7,8
$P_8 \times P_1$	0,19	2,11	-20,23**	-0,040	-0,013	-0,003	-99,1**
$P_1 \times P_7$	-2,48**	-9,29**	24,70**	-0,033	0,868**	0,330	100,1**
$P_1 \times P_8$	3,07**	18,74**	7,00**	0,227**	0,322**	0,040**	128,1**
$P_7 \times P_8$	-1,81**	-34,27**	-9,37**	-0,104	-0,037	-0,010*	-114,7**
$P_8 \times P_7$	-1,38**	10,73**	9,34**	-0,140	-0,177	-0,384*	-21,0
$P_7 \times P_1$	-0,48	0,09	-32,07**	-0,109*	-0,482**	-0,070**	-214,1**
$P_8 \times P_1$	-0,26	-10,98**	-26,76**	0,017	0,039	0,278	-100,1**
$P_7 \times P_8$	1,69**	-20,89**	23,91**	-0,074	0,330*	0,399*	176,9**
$P_8 \times P_7$	-0,21	10,71**	31,73**	0,077	-0,044	0,076**	247,0**
$P_7 \times P_1$	0,24	8,78**	-0,73**	-0,037	-0,746**	-0,434*	-91,9**
$P_8 \times P_1$	0,02	10,21**	3,34	0,370**	-0,006**	0,221	47,8
$P_7 \times P_8$	4,40**	4,71	3,00	-0,288**	-0,413**	-0,729**	-89,7**
$P_8 \times P_7$	-1,10**	-2,86	8,20**	-0,042	0,170	-0,170	9,6
$P_7 \times P_1$	0,17	-22,86**	-1,09	0,172*	-0,120	-0,236	-24,6
$P_8 \times P_1$	1,29**	14,73**	-28,07**	-0,109	0,110	-0,010	-113,0**
SE (Sij)	0,43	3,00	2,07	0,081	0,107	0,182	28,8

Specific combining ability effects (S_{ij}) of each reciprocal cross combination (F_{rj}) for all studied traits are found in Table 5. The results showed that no reciprocal cross was the best for all studied traits. However, nine and five out of thirty six reciprocal hybrids exhibited significant SCA effects for earliness and plant height, respectively. For yield and its component, sixteen, four, three, six and nine out of thirty six reciprocal crosses revealed desirable SCA for number of fruit per plant, fruit diameter, fruit length, fruit weight and total yield per plant, respectively.

Table 5: Estimates of specific combining ability effects (S_{ij}) of each reciprocal cross (F_{rj}) for all studied traits

Crosses	No. of DF	PH (cm)	No. of F/P	FD (cm)	FL (cm)	FW (gm)	TY/P (gm)
$P_{r1} \times P_1$	1,0*	-24,00**	-7,00**	0,217*	-0,183	0,60**	63,7
$P_{r2} \times P_1$	0,0	0,0	28,00**	-0,217*	-0,200	-0,377	68,9*
$P_{r3} \times P_1$	-0,83	-2,00	-33,00**	-0,400**	0,233	-1,200**	-321,0**
$P_{r4} \times P_1$	0,77	-10,00**	21,00**	-0,033	-0,083**	-0,017*	2,4
$P_{r5} \times P_1$	2,67**	13,00**	81,00**	-0,017	0,300	0,077	40,4**
$P_{r6} \times P_1$	1,33**	-12,00**	16,00**	-0,000	0,617**	0,377	100,0**
$P_{r7} \times P_1$	1,00*	-40,00**	-38,00**	0,200**	0,633**	0,017*	-100,0**
$P_{r8} \times P_1$	1,33**	-17,00**	-37,00**	0,083	0,333	0,777**	-110,0**
$P_{r9} \times P_1$	2,17**	-2,00	-23,00**	0,300**	-0,217	0,383	-01,9
$P_{r10} \times P_1$	-1,33**	-1,00	-14,00**	-0,083	-0,433**	-0,133	-74,9*
$P_{r11} \times P_1$	0,83	-21,17**	20,00**	0,317**	-0,433**	-0,133	76,4*
$P_{r12} \times P_1$	-0,33	4,00	10,00**	0,100	-0,700**	0,083	03,0
$P_{r13} \times P_1$	-1,00*	-8,00*	-0,00	-0,100	0,300	-0,100	-24,9
$P_{r14} \times P_1$	-0,83	-3,00	4,00	-0,100	0,000	0,133	37,9
$P_{r15} \times P_1$	1,83**	3,00	34,00**	-0,133	-0,100	-0,100	10,4**
$P_{r16} \times P_1$	-0,33	-0,00	-12,00**	0,000	0,217	0,377	13,0
$P_{r17} \times P_1$	0,77	3,00	6,00*	-0,177	0,177	-0,317	-21,1
$P_{r18} \times P_1$	-0,83	-22,00**	0,00*	-0,133	0,877**	0,317	79,0*
$P_{r19} \times P_1$	1,00*	27,00**	30,00**	-0,477**	-0,000	-1,017**	41,8
$P_{r20} \times P_1$	-0,33	-6,00	-30,00**	-0,417**	-0,300	-0,633**	-233,4**
$P_{r21} \times P_1$	1,00*	-37,00**	-8,00**	-0,000**	0,133	-0,933**	-142,7**
$P_{r22} \times P_1$	2,00**	-17,00**	19,00**	-0,117	-0,077	-0,300	42,6
$P_{r23} \times P_1$	2,00**	-20,00**	-40,00**	0,077	0,077**	0,377	-142,9**
$P_{r24} \times P_1$	1,00*	-22,00**	-10,00**	-0,233**	0,017	-0,300	-100,2**
$P_{r25} \times P_1$	0,33	-81,00**	-3,00	0,017	-0,233	0,233	20,1
$P_{r26} \times P_1$	-1,33**	6,00	4,00	0,017	0,000	0,133	38,1
$P_{r27} \times P_1$	-3,33**	-2,00	-19,00**	-0,000	0,100	0,017	-71,3*
$P_{r28} \times P_1$	-2,00**	-6,00	-28,00**	0,133	0,117	0,200	-127,7**
$P_{r29} \times P_1$	-4,00**	-04,00**	-6,00**	-0,000	-0,100	-0,083	-44,0
$P_{r30} \times P_1$	-1,00**	-10,00**	8,00**	0,133	0,100	0,677**	104,6**
$P_{r31} \times P_1$	3,00**	6,33	10,00**	-0,000	0,083	-0,317	2,4
$P_{r32} \times P_1$	3,67**	-28,00**	-4,00	-0,217*	-0,200	-0,433**	-89,0**
$P_{r33} \times P_1$	0,0	21,00**	38,00**	-0,283**	-0,077	-0,433**	83,0*
$P_{r34} \times P_1$	3,00**	10,00**	-13,00**	0,217*	0,200	0,183	-31,9
$P_{r35} \times P_1$	-3,00**	30,00**	48,00**	0,100	0,217	0,477*	290,4**
$P_{r36} \times P_1$	-1,67**	-10,00**	30,00**	-0,033	-0,233	0,433*	187,7**
SE (S_{ij})	0,49	3,47	2,39	0,093	0,182	0,211	33,3

It could be noticed that the excellent cross combinations were obtained from crossing (good x good), (good x poor) and (poor x poor) general combiners. Therefore, it is not necessary that parents having estimates of high GCA effects would also give high estimates of SCA effects in their respective cross combinations. These results suggest the important role of non additive gene action in the inheritance of the studied traits.

Nature of gene action

Based on the analysis of combining ability, the different genetic parameters were estimated and the obtained results are presented in Table 1. The results indicated that the magnitudes of the non additive genetic variance (VD) were larger than those of additive ones (VA) for all studied traits except for fruit diameter and fruit weight. In this direction, Dhankhar and Dhankhar (2001), Prakash et al (2002) and Solankey and Singh (2010) stated that non additive genetic variance was higher than the additive one for days to flowering, plant height, number of branches, number of pods per plant and pod yield per plant. Considerable values of reciprocal effects variance were observed in all studied traits, exhibiting the important role of cytoplasmic factors in the expression of these traits. Furthermore, the broad sense heritability estimates (H_b^2 %) were more than 50% and larger than their corresponding narrow sense heritability (H_n^2 %) for all studied traits. However, estimates of narrow sense heritability were 13.9%, 32.4, 40.0, 47.1, 56.8 for earliness, fruit length, fruit weight, plant height and fruit diameter, respectively. With respect to yield components, the estimates of narrow sense heritability ranged from 11.3 % to 17.3% for total yield per plant and No. of fruit per plant, respectively. These results verified the predominance of non additive gene action in the inheritance of these traits. These results are in agreement with those obtained by Prakash et al (2002) and Salameh and Kasrawi (2007).

Table 1: Estimates of genetic parameters and heritability in broad (H_b^2 %) and narrow (H_n^2 %) sense for all studied traits.

Genetic Components	No. of DF	PH (cm)	No. of F/P	FD (cm)	FL (cm)	FW (gm)	TY/P (gm)
VA	0.024	224.1	81.08	0.126	0.102	0.140	1986.174
VD	2.764	227.8	377.47	0.21	0.146	0.120	1336.192
Vr	3.321	046.0	719.00	0.037	0.084	0.180	17169.992
VE	0.482	24.0	11.44	0.017	0.066	0.089	2216.223
H_b^2 %	87.2	94.9	97.07	89.4	78.9	70.3	87.4
H_n^2 %	13.9	47.1	17.34	56.8	32.4	40.0	11.3

In conclusion, it could be noticed that most studied traits were mainly controlled by non additive effects and cytoplasmic factors. Thus, the genetic material used in this study could be used for hybridization for producing promising crosses to improve economic traits in okra.

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القدرة على التآلف وطبيعة فعل الجين في الباميا

سهير السيد عبده الجندي^١، حازم عبد الرحمن عبيد الله علي^٢، إيهاب عوض الله إبراهيم^١،
محمد حمام زين العابدين الدقيشي^٣
^١قسم بحوث الخضار- معهد بحوث البساتين - مركز البحوث الزراعية- مصر.
^٢قسم البساتين- كلية الزراعة - جامعة سوهاج- مصر.
^٣قسم البساتين- كلية الزراعة - جامعة اسيوط- مصر.

تم إجراء هذا البحث بمحطة البرامون الزراعية بالمنصورة - معهد بحوث البساتين- مركز البحوث الزراعية. أجري هذا البحث لدراسة القدرة على التآلف وطبيعة الفعل الجيني لبعض الصفات الاقتصادية في الباميا وذلك باستخدام نظام التهجين الدائري الكامل بين تسع سلالات وراثية محلية من الباميا.

ويمكن تلخيص أهم النتائج فيما يلي:

أوضحت نتائج تحليل التباين وجود فروق معنوية بين التراكيب الوراثية لكل الصفات المدروسة. كانت تقديرات القدرة العامة والخاصة علي التآلف معنوية جدا لكل الصفات تحت الدراسة مما يؤكد أهمية التباين الوراثي المضيف وغير المضيف في وراثية الصفات تحت الدراسة. -أوضحت النتائج أن الآباء $P_0, P_1, P_2, P_3, P_4, P_5, P_6, P_7, P_8$ ذات قدرة عامة عالية علي التآلف لصفة التبرير بينما كانت الآباء $P_1, P_2, P_3, P_4, P_5, P_6, P_7, P_8$ ذات قدرة عامة علي التآلف لصفة المحصول الكلي. كانت الهجن ($P_0 \times P_1$), ($P_0 \times P_2$), ($P_0 \times P_3$), ($P_0 \times P_4$), ($P_0 \times P_5$) and ($P_0 \times P_6$) ذات قدرة خاصة عالية علي التآلف لصفة التزهير المبكر ومعظم صفات المحصول. كانت قيمة التباين الوراثي غير المضيف أكبر من التباين الوراثي المضيف لصفة التبرير، و صفات المحصول ومكوناته. كانت أعلى قيم لدرجة التوريث في المدى الواسع أكبر من 75% وكانت أعلى من قيم درجة التوريث في المدى الضيق لكل الصفات قيد الدراسة حيث كانت قيمة درجة التوريث في المدى الضيق (9,13%) لصفة التبرير بينما كانت (3,11%) لصفة المحصول الكلي. طبقا لنتائج التحليل الوراثي للصفات تحت الدراسة يمكن استخدام هذه الهجن المباشرة في الحصول علي أعلى محصول من الباميا.

قام بتحكيم البحث

أ.د / ممدوح محمد عبد المقصود
أ.د / جلال احمد رزق الشربيني

أ.د / ممدوح محمد عبد المقصود
أ.د / جلال احمد رزق الشربيني