PERFORMANCES AND CORRELATIONS OF PARENTAL LINES AND THEIR HYBRIDS FOR ECONOMICAL TRAITS OF CANOLA, *Brassica napus* L.

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ABSTRACT

This investigation was conducted to study the mean performances of 10 parental lines of canola and the F_1 hybrids among them. Complete diallel crosses mating design were used to obtain 45 F_1 hybrids and 45 F_1 reciprocal hybrids at three environments. These environments were: El-Gamalia 2002/2003 (E_1), El-Serw 2002/2003 (E_2) and El-Serw 2003/2004 (E_3). Different vegetative, yield traits and oil percent were studied. In the same time, the nature of association among pairs of studied traits were evaluated.

The presence of significant variation which noticed among all studied traits makes it possible and necessary to compare the means of all genotypes to evaluate their performances. However, the inbred line $_{325}$ (P_8) appeared to be the earliest, while the parental line Serw $_6$ (P_2) was the tallest and the best parent for N.se./sil. The line Serw $_{37}$ (P_3) had the larger N.pr.br./pl., while the line Serw $_4$ (P_1) started its flowering branches lower than the earlier parents and it was the largest for S.Y./pl.. The Line $_{164}$ (P_{10}) was the best parent for 1000 S.W.gms. and oil %. The Line Serw $_{98}$ (P_5) was the best for N.sil./pl.. The combined data showed that the hybrid between P_4 x P_7 produced more oil % of 46.7 %, while the hybrid between P_1 x P_8 produced the less oil of 36.7 %. The results also showed that the N.se./sil. and N.sil./pl. were very effective in the determination of S.Y./pl. in grams, while 1000 S.W.gms. trait was less effective. Therefore, selection for more N.sil./pl. and more N.se./sil. could increase the yield.

The most desirable r_{ph} and r_g correlations were obtained among S.Y./pl. gms. and N.sil./pl. which showed positive and significant values. Therefore, these two traits showed a desirable association. This finding indicated that selection program to increase one trait would, indeed, increase the other. It could be also mention that all negative r_{ph} and r_g correlations were insignificant. This result indicated that this type of association was not important, where the values of r_{ph} and r_g correlations were small. The negative r_{ph} and r_g correlations between 1000 S.W. gms. and N.sil./pl. indicated that when plant had more siliqua, the size and weight of seeds become less.

INTRODUCTION

Kandil *et al.* (1996) evaluated four parents of canola and six F_1 hybrids among them. They found significant differences among the parents and their F_1 hybrids for N.pr.br./pl., N.sil./pl., S.Y./pl. and S.Y./m² traits, while 1000 S.W. and Oil % were not significantly different. In this respect, Hammad (1998) evaluated 15 F_1 hybrids and their parents. He noticed the presence of significant variations between these genotypes for N.pr.br./pl., P.H.cms., N.se./pl., 1000 S.W. and S.Y./pl.gms traits. Similarly, Riaz *et al.* (2001) found that the magnitudes of variances among inbred lines and their F_1 hybrids over two locations were significant for P.H. and D.M. traits. In the same way, Teilep (2003) and Kassab (2004) studied different F_1 hybrids of canloa and

their parents. They illustrated the presence of variation among these genotypes for D.50%f., D.M., P.H., N.pr.br./pl., N.se./sil., N.sil./pl., 1000 S.W. and Oil % traits.

Many authors studied the mean performances of F_1 hybrids of canola among them, Kandil *et al.* (1996), Halaka (2000), Teilep (2003) and Kassab (2004). They obtained high performances F_1 hybrids of canola, although the exhibited mean values differed among them.

Concerning phenotypic and genotypic correlations in canola, Hamed (1993) found that pod yield trait was significantly correlated with oil % (0.98) and protein yield (0.97) traits. He also added that seed yield trait showed highly significant correlations with pod yield (0.94), biological yield (.80) and strew yield (0.72). In this respect, Ozer *et al.* (1999) obtained positive values of correlations among S.Y./pl. gms and each of: D.50% f., P.H., N.pr.br./pl., N.sil./pl., N.se./sil., 1000S.W. and oil % traits. On the other hand, El-Baz and El-Shakhess (2001) reported negative and significant phenotypic and genotypic correlations among P.H. and oil % which were: -0.33 and -0.37, respectively.

In another study by Sharief and Keshta (2002) and Maria et al. (2003), they found positive and significant correlation coefficients between P.H. and each of: oil %, S.Y./fe. and S.Y./pl.. On the other hand, Marinkovic et al. (2003) observed negative and significant correlation coefficients between S.Y./pl. and each of: P.H. (-0.303), H.f.br. (-0.27) and N.pr.br./pl. (-0.06). In addition, Teilep (2003) and Kassab (2004) reported that positive and significant correlation between 1000 S.W. and days to first flower, while it was negatively and significantly correlated with N.pr.br./pl..

MATERIALS AND METHODS

The genetic materials used in the present investigation included 10 inbred lines of canola, $Brassica\ napus$, L.. The seeds of all inbred lines were obtained from Oil Crops Research section, El-Serw Agricultural Research station, Agric. Res. Center, Egypt. These inbred lines were: Serw4, Serw6, Serw37, Serw64, Serw98, Serw101, Serw103, Line325, Line163 and Line164. In the growing season of 2002, all inbred lines were crossed to obtain 45 F_1 and their 45 F_{1r} hybrids through a complete diallel crosses mating design including reciprocal hybrids. In addition, the parental lines were selfed to obtain additional amounts of seeds for further investigation.

Ali (100) genotypes were evaluated at the three environments: El-Gamalia 2002/2003 (E $_1$), El-Serw 2002/2003 (E $_2$) and El-Serw 2003/2004 (E $_3$). The experimental design was a randomized complete blocks design with three replications. Each block consisted of 100 plots. The plot was a single row 4.2 m. long and 0.6 m. wide. The agricultural practices were carried out as recommended for canola plantation.

The data were recorded on the following traits:

Days to 50 % flowering (D.50%F.); plant height in centimeters (P.H.cms.); number of primary branches per plant (N.pr.br. /pl.); height of the first branch (H.f.br.cms.); weight of 1000 seeds in grams(1000 S.W.gms);

number of seeds per siliqua(N.Se./sil.); number of siliqua per plant(N.sil./pl.); seed yield per plant in grams (S.Y./pl.gms) and oil percent (oil %).

The analyses of variance for each environment and the combined analysis over environments were made according to Steel and Torrie (1960) and Cockerham and Cox (1963), respectively.

Estimation of phenotypic and genotypic correlations between all pairs of studied traits required, a covariance analyses between all pairs of traits at each environment and from the combined data over the three environments according to the procedures as outlined by Singh and Chaudhary (1985).

RESULTS AND DISCUSSION

There are very few studies on canola have done in Egypt. However the Ministry of Agriculture has introduced and selected several cultivars. Now, it is the time to start a serious studies on this crop to evaluate the possibility of cultivating it in newly reclaimed lands. Indeed, the investigation will threw a light about the nature of variation among these lines, the mean performances and correlations among studied traits.

The genetic materials used in this investigation included 10 inbred lines which were involved in a complete diallel crosses mating design to produce 45 $F_{\rm 1}$ hybrids and 45 $F_{\rm 1r}$ reciprocal hybrids. All genotypes which included the 10 parental lines and the 90 $F_{\rm 1}$ hybrids were evaluated at three environments: El-Gamalia 2002 / 2003, El-Serw 2002 / 2003 and El-Serw 2003 / 2004. The vegetative, yield, yield component traits and oil percent were studied. Therefore, form the economical point of view, the evaluation of the parental lines and the $F_{\rm 1}$ hybrids among them is of paramount importance specially under several environments.

The mean performances of parental lines:

The means of all parental lines for all studied traits at three different environments and from the combined data over the three environments were calculated and the results are presented in Table 1.

In general, all parental lines showed high performances for all studied traits at environment E_1 (El-Gamalia 2002/2003), while the lowest performances for most traits were obtained at environment E_2 (El-Serw 2002/2003). This finding was expected because the soil at E_1 is more fertile and irrigated by river Nile water , while the soil at E_2 is poor in fertility, has a higher level of salinity and irrigated by drainage water.

The means of the 10 parents for D.50%F. ranged from 76 days to 110 days for parent P_8 at E_2 and parent P_6 at E_1 , respectively. The combined data over the three environments for this trait ranged from 83 days to 107 days for parent P_8 and parent P_6 , respectively. These results indicated that the parent P_8 was earlier in flowering data than the parent P_6 and all other parents.

The mean performances of parents for all studied traits at the three environments and from the combined data over the threenvironments.

	EVILLE	D.Suzak.	P.H. cms.	N. pri. br./pl.	H. f. bra.	1000 S.W.	N.se./sil.	N.sil./pl.	Se.Y./pl.	L
1	17	16		× 0.	7.1	3.3	29.4	557	512	L
	E.3	68	16	9.8	56	3.6	250	37.4	33.1	L
1		2	121		5.9	3.5	25.6	486	6##	
-	Comb.	NS N	121	10, 5	1.9	3.5	26.7	472	43.7	
	F.1	801	212	66	45.2	3.2	31.3	191	46.2	
	77	100	021	68	32.1	3.1	26.8	306	\$ 92	
,	E.		6/1	10.2	39.7	3.2	26.5	430	37.1	
1	Collin.	103	89	9.7	39.0	3.1	28.2	90+	36.6	38.0
	E	IIX.	124	671	10.5	4.0	26.4	357	396	
1	F.2	5.	55	7.11	15.8	2.2	244	323	17.8	
1	E3	66	120	14.2	9.61	2.6	24.9	77.0	38.5	
1	Comb.	101	123	12.5	15.3	2.9	25.3	698	286	
	E,	9.6	268	8.5	22.7	1.1	312	146	5 01	
	E2	96	6/6	2.5	28.5	3.0	25.5	303	46.3	
	E,	06	147	NG.	316	3.0	100	16.7	21.0	
-	Comb.	16	151	C.X.		11.6	1.137	489	18.4	
-		86	Name and Address of the Owner, where the Owner, which is the Owner, where the Owner, which is the Owner, where the Owner, which is the O	O C	Co. T.	AND DESCRIPTION OF PERSONS ASSURED AND DESCRIPTION OF PERSONS ASSURED	AND DESCRIPTION OF PERSONS ASSESSMENT OF PER	412	16.8	-
		00		0.0	8.7	3.5	22.5	647	50.7	
	-	000	130	8'/	2.6	3.6	23.9	354	30.5	
1		-	661	9.9	9,8	3.6	22.3	\$15	12.3	
1	Comb.	7.4	1+()	+%	00.00	3.5	23.0	305	41.2	
	E1	011	164	78	150	3.4	27.5	189	17.6	
	5	104	66	59	511	2.8	29.7	300	15.5	
1	13	107	129	7.2	184	2.5	30.2	475	37.7	
1	Comb.	10,	131	7.0	150	2.9	29.1	421	698	
	77	103	210	11.3	5.11	3.6	X.8.C	166	18.1	
	2 2	5.		10.6	55.3	3.3	25.1	35.1	20.2	
1	1.3	96	139	13.8	43.8	3.2	26.6	170	9119	
-	Comb.	Х6	[60]	611	+++	3.4	26.8	433	39.9	
	E.	68	153	9.2	8.3	3.8	29.1	430	18.7	-
	E2	76	8.5	9.2	6.6	3.5	25.9	107	1 × ×	
1	[2]	2 2	9=	8.01	7.0	3.7	28.1	:82	\$ OF	
1	Comb.	6.5	2011	9.7	74	3.7	27.7	373	19.2	
1	E	86	17.	8.1	12.5	2.6	28.8	13.7	36.1	
	E2	06	101	7.3	15.6	2.4	163	147	1.00	
	E ₃	91	132	7.8	212	3.0	38.8	304	0.0	
Ü	Comb.	93	135	1 1 1	16.6	36	200	200	6.67	
	E,	101	192	8.0	0.0	1	007	333	26.6	1
	E,	93	89		7.0	17.	27.5	\$28	46.7	
	E	93	126	100	9.7	7	25.8	210	22.5	
0.0	1	100		2.3	10.5	0.1	25.6	126		
		1111	113	6.3			2004	323	7.1.	

The results also cleared that P.H. cms. ranged from 212 cms. to 85 cms. for P_2 at E_1 and P_8 at E_2 , respectively. The combined data over the three environments for this trait ranged from 168 cms. to 118 cms. for P_2 and P_8 , respectively. These results indicated that the P_2 was taller than P_8 and all other parents.

In the same time, the results indicated that N.pr.bra/pl. ranged from 14.2 branches to 5.9 branches for P_3 at E_3 and P_6 at E_2 , respectively. The combined data over the three environments for this trait ranged from 12.5 branches to 7.0 branches for P_3 and P_6 , respectively. These results indicated that P_3 was the highest for N.pr.bra./pl. than P_6 and all other parents.

The means of the 10 parents for H.f.bra. ranged from 5.6 cms. to 54.1 cms. for P_1 at E_2 and P_7 at E_1 , respectively. The combined data over the three environments for this trait ranged from 6.4 cms to 44.4 cms for P_1 and P_7 , respectively. These results indicated that P_1 started to have flowering

branches near the surface from of soil than P7 and all other parents.

The means of the parental lines for 1000 s.w.gms. ranged from 4.2 gms to 2.2 gms. For parent P_{10} at E_2 and parent P_3 at E_2 , respectively. The combined data over the three environments for this trait ranged from 3.8 gms. to 2.6 gms. for parents P_{10} and P_9 ,respectively. Thus, these findings indicated that P_{10} was higher in 1000 s.w.gms. than parent P_9 and all other parents. Similarly, N.se./sil. ranged from 31.5 seeds to 22.5 seeds for P_4 at E_1 and P_5 at E_1 and E_3 , respectively. The combined data over the three environments for this trait ranged from 29.1 seeds to 23.0 seeds for P_6 and P_5 , respectively. These results indicated that P_6 had more N.se./sil. than P_5 and all other parents.

The results indicated that N.sil./pl. ranged from 647 siliqua to 210 siliqua for P_5 at E_1 and P_{10} at E_2 , respectively. The means from the combined data over the three environments for this trait ranged from 505 siliqua to 333 siliqua for P_5 and P_9 , respectively. Thus, these results indicated that P_5 was the highest for N.sil./pl. than P_9 and all other parents.

The means of the 10 parents for S.Y./pl. gms. ranged from 53.2 gms to 17.8 gms for P_1 at E_1 and P_3 and P_9 at E_2 , respectively. The combined data over the three environments for this trait ranged from 43.7 gms. to 26.6 gms for P_1 and P_9 , respectively. These results indicated that P_1 was the highest for S.Y./pl. gms. than P_9 and all other parents.

The means of the 10 parental lines for oil % ranged from 43.1 % to 36.4 % for P_4 , P_6 and P_{10} at E_1 and P_9 at E_3 , respectively. The combined data over the three environments for this trait ranged from 41.5 % to 37.2 % for P_{10} and P_9 , respectively. These results indicated that P_{10} was the highest parent for oil % than P_9 and all other parents.

In general, the means of the 10 parental lines for all studied traits did not show the superiority of certain parent for all traits. However, the line $_{325}$ (P_8) appeared to be the earliest, while the line $Serw_6$ (P_2) was the tallest and the best parent for N.se./sil. The line $Serw_{37}$ (P_3) had the larger N.pr.br./pl. While the line $Serw_4$ (P_1) started its flowering branches lower than the earlier parents and the largest S.Y./pl.. While, the Line₁₆₄ (P_{10}) was the best parent for 1000 S.W.gms. and oil %. The Line $Serw_{98}$ (P_5) was the best for N.sil./pl..

The F₁ hybrids and their reciprocals:

According to the complete diallel crosses mating design of the 10 parental lines, the resulted 45 F_1 hybrids and their 45 F_{1r} reciprocal hybrids at the three environments and from the combined data over environments were evaluated for all studied traits and their mean performances are presented in Table 2. The means of all F_1 and F_{1r} hybrids were arranged so that the first parent was the female parent, while the second parent was the male parent.

In general, the means of all F_1 hybrids varied not only from one F_1 hybrids to another but also, for the same F_1 hybrid at the different environments. These results indicated that environments exerted important effects on the same genotypes. The results also showed the presence of significant differences among all the F_1 hybrids and their F_{1r} reciprocals hybrids which were true at the three environments and certainly from the combined data. These differences between the hybrids and their reciprocals ranged from insignificant up to highly significant.

For number of days to 50 % flowering (D.50%f.) all F_1 hybrids and their reciprocals appeared to have larger means indicating that they flowered later than their parents. These results indicated that F_1 hybrids had a longer period of vegetative growth which would be noticed when the plant height trait would be studied. It is a common fact that plants which have longer period of vegetative growth have also more yield. The earliest plants were also shorter and had their in H.f.br. near the surface of the soil.

The results also cleared that the earliest F_1 hybrid was the F_1 hybrid between $P_8 \times P_4$ which flowered after 80 days from planting at E_2 , while the latest F_1 hybrid in flowering was the cross between $P_6 \times P_7$ which flowered after 124 days from planting at E_1 . The means obtained from the combined data showed similar trend where the earlier F_1 hybrid over the three environments was also the same hybrid between $P_8 \times P_4$ which gave 50 % flowering rate after 85 days from planting and the latest F_1 hybrid was also the F_1 hybrid between $P_6 \times P_7$ which gave 50 % flowering rate after 115 days from planting. Therefore, the range of 50 % flowering rate of the F_1 hybrids significantly varied when compared with the range of the parents for the same trait which showed a range of 83 days to 107 days.

The results also, indicated that the reciprocal hybrids were close to each other with few differences ranged from not significant up to highly significant. It was also noticed that all F_1 hybrids and their reciprocals were later at E_1 than the other two environments. These results indicated that the rich soil which irrigated by river Nile water gave more and stronger vegetative growth causing the hybrids to flower later than the same F_1 hybrids at the other two environments which were poorer in their fertility.

For plant height in centimeters (P.H.cms.) all F_1 hybrids and their reciprocals appeared to have larger means indicating that they were taller than their parents. These findings indicated that F_1 hybrids had a taller plants which in turn caused an increase in N.pr.br./pl..

The obtained means showed that the tallest F_1 hybrid was the F_1 hybrid between $P_3 \times P_4$ which had a height of 232 cms. at E_1 , while the shortest F_1 hybrids was the hybrid between $P_6 \times P_9$ which was 95 cms. in height at E_2 . The means obtained from the combined data over the three

environments showed that the tallest F_1 hybrid was the F_1 hybrid $P_4 \times P_{10}$ which had 198 cms. in length and the shortest F_1 hybrid was the F_1 hybrid $P_8 \times P_1$ which showed 128 cms. in length. Therefore, P.H. cms. significantly varied and had a wider range than their parents which showed a range of 118 cms. to 168 cms. for the same trait.

The results also, indicated that the reciprocal hybrids were not close to each other with differences ranged from not significant up to highly significant. It was an important observation to noticed that all F_1 hybrids and their reciprocals were taller at E_1 than at the other two environments.

The results indicated that E_1 yielded plants with more vegetative growth than the same F_1 hybrid at the other two environments which were

poor in soil and irrigation water with high degree of salinity.

The number of primary branches per plant (N.pr.br./pl.) for the F_1 hybrids and their reciprocals tended to have larger means than their parents. The reported means in Table 2 showed that the highest F_1 hybrid for this trait was the F_1 hybrid between P_1 x P_6 at E_1 and P_2 x P_7 at E_3 which had 14.8 branches per plant, while the lowest F_1 hybrid for this trait was the cross between P_5 x P_4 which had 6.5 branches at E_2 . The means obtained from the combined data over the three environments showed that the highest F_1 hybrids were the F_1 hybrids between P_2 x P_7 ; P_3 x P_6 ; P_6 x P_8 and P_{10} x P_7 which showed 13.5 branches per plant, while the lowest F_1 hybrid was the F_1 hybrid between P_5 x P_4 which gave 8.3 branches per plant. Therefore, N.pr.br./pl significantly varied in the F_1 hybrids than the range of the parents for the same trait which showed a range of for 7.0 to 12.5 branches per plant.

The results also indicated that the F_1 hybrids and their reciprocals were not close to each other with differences ranged from not significant up to highly significant. It was very important observation to notice that all F_1 hybrids and their reciprocals were higher at E_1 than the other two environments.

The means of height of first branch in centimeters (H.f.br.cms.) for all F_1 hybrids and their reciprocals appeared to have smaller means indicating that they started to produce their flowering branches near the soil surface. The reported means showed that the F_1 hybrid P_3 x P_6 at E_3 started to have flowering branches at 2.5 cms. from soil surface, while the F_1 hybrid P_6 x P_2 at E_1 started to produce the first flowering branches at 38.9 cms. from soil surface. The obtained means from the combined data over the three environments showed that the F_1 hybrid P_4 x P_8 started to have its flowering branch at 3.4 cms. from soil surface, while the F_1 hybrid between P_6 x P_2 started at 28.5 cms from soil surface. Therefore, H.f.br.cms. significantly varied in the F_1 hybrids and had a wider range than the range of the parents which showed a range of 6.4 cms. to 44.4 cms. from the soil surface. The results also, indicated that the F_1 hybrids were not close to their reciprocals with differences ranged from not significant up to highly significant.

In general, it was apparent that variations within parental lines were present but in different magnitudes for each trait. Accordingly, the two traits D.50 % f. and N.pr.br./pl. showed less variations than the two traits P.H.cms.

and H.f.br.cms. which showed larger amounts of variation.

The mean performances of F₁ hybrids (F₁) and their F₁ reciprocal hybrids (F_{1,1}) for all studied traits at the three environments and from the combined data over the three environments.

		D.50	%F.			P.Hcms.	cms.			N.pr.br./pl	n:/pl.			H. f. bra.cms.	TI.CINS.	
	E	E,	E	Comb.	E,	E,	F.3	Comb.	E,	E;	Ε,	Comb.	E,	E,	E3	Comb.
p ₁ xp ₂	112	95	97		219	138	180	179	14.4	10.8	11.9	12.3	22.3	21.5	25.9	23.2
P.xp.	103	94	96		215	152	195	187	12.9	11.3	14.1	12.8	17.3	18.3	17.3	17.6
DixDs	112	98	66		166	109	131	135	11.4	8.4	10.3	10.0	13.2	13.0	19.3	15.2
Pixpi	104	94	93		167	111	142	140	10.9	8.8	10.2	10.0	11.9	7.5	6.6	8.6
DIXDA	114	66	102	105	214	120	153	162	10.7	9.5	9.8	10.0	12.7	7.8	10.1	10.2
P.XD.	102	94	97		206	119	166	164	12.2	8.5	11.7	10.8	11.0	5.6	8.9	8.5
DixDe	116	103	106		183	125	161	156	12.4	6.6	10.4	10.9	13.5	14.1	15.9	14.5
P.x.D.	101	91	92		180	109	142	144	10.0	7.6	10.8	9.5	9.7	15.0	15.6	13.4
D, VD.	117	103	107		187	127	169	161	14.8	11.1	12.3	12.7	3.8	4.5	7.1	5.1
P.XD.	101	98	101		198	126	178	168	10.6	8.3	11.2	10.0	15.8	17.1	18.5	17.1
D.XD-	119	107	110		191	116	150	152	9.2	9.0	9.1	9.1	23.5	18.0	23.0	21.5
P-xp.	109	97	102		213	107	143	154	10.1	9.5	10.4	10.0	4.8	9.2	6.5	6.8
DIADS	102	92	94		159	113	144	139	13.1	8.5	10.8	10.8	10.3	13.0	17.1	13.5
Pavni	96	86	93		177	26	111	128	9.9	7.6	9.6	06	9.5	5.7	9.1	8.1
DiXDo	101	92	93		179	135	174	163	14.0	7.3	10.8	10.7	11.5	8.0	10.4	10.0
PoXpi	98	89	91		175	120	167	154	11.6	7.8	10.2	6.6	23.7	18.9	20.8	21.1
DiXDia	104	63	96		201	103	133	148	9.6	8.8	9.4	9.3	12.9	10.3	11.2	11.5
Piaxbi	101	66	103		187	125	161	158	9.7	8.2	10.9	9.6	19.3	9.7	18.9	16.0
DiXDi	113	100	102		219	158	169	192	12.6	11.5	14.1	12.7	21.8	12.8	20.9	18.5
Paxp,	111	101	103		226	120	168	171	12.4	7.5	10.0	10.0	23.1	23.9	27.6	24.9
DAND	109	66	100		227	146	195	190	14.1	7.5	13.5	11.7	20.2	18.7	19.0	19.3
P.ND,	104	89	93		219	144	189	184	-14.3	11.4	14.6	13.4	17.3	13.5	16.5	15.7
Pykus	116	100	105		204	118	152	158	13.2	9.4	12.3	11.6	12.3	5.5	9.7	9.2
Dexin,	107	98	100		218	123	160	167	11.3	7.7	10.7	9.9	10.8	12.2	11.8	11.6
P, v.D.	118	104	107		196	107	148	151	11.0	0.6	11.2	10.4	15.9	25.6	13.9	18.5
DoxD:	111	92	94		214	107	152	157	9.5	9.5	8.3	9.1	38.9	21.3	25.4	28.5
P,xD;	120	106	108		224	137	185	182	2	11.2		13.5	34.0	8.0	6.6	15.8
D-XD;	120	105	111		229	127	181	179	14.2	10.6	12.8	12.5	30.7	19.7	22.7	24.4
Paxps	101	92	92		197	118	188	168		9.2	12.8	11.5	17.5	4.5	29.0	17.0
DAXDS	101	91	98		216	147	182	182	10.6	7.8	9.0	9.1	5.4	4.0	8.1	5.9

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+	E	£3	E.3	-	1700	133	180	172	26	8.9	10.3	9.6	19.4	10.8	16.0	
P ₂ xp ₉	108	95	95		507	144	181	178	8	633	9.5	9.6	4.6	5.0	9.7	6.5
D ₉ XP ₂	111	94	96		717	141	0	0 00	10.2	7 4	63	8.9	16.3	9.8	11.3	12.1
PaxDan	113	96	97		211	126	801	000	100	0 2	100	10.8		18.0	21.8	19.5
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P ₃ xp ₆	001	200	010	+	192	116	155	155	12.6	10.9	12.9	17.7	13.5	20.00	2.0	2.0
03	16	000	104	+	201	133	175	170	14.4	11.6	14.5	13.5	10.8	3.9	5.5	0.0
P ₃ xp ₆	109	200	5	+	000	120	185	174	12.6	7.9	11.7	10.7	10.0	10.6	12.2	11.0
03	109	96	200	-	007	200	187	178	118	8.3	12.6	10.9	13.5	23.2	16.0	17.6
P ₃ xp ₇	115	102	106	+	210	000	124	200	130	77	8.6	10.2	8.8	5.9	9.3	8.0
p ₇ xp ₃	112	101	102	105	211	771	150	151	66	87	8.6	9.5	14.7	7.6	9.2	10.5
Paxpa	97	68	91	-	1/3	771	000	2 4	110	a a	103	10.0	15.2	7.9	12.9	12.0
2	91	82	84	_	186	111	158	761	77.0	0 0	128	110	14.6	15.4	18.7	16.2
D'AD	103	92	94	_	155	102	138	132	0 0	1 0	2 0	α	12.0	107	10.4	11.0
2 0	100	86	89	-	203	148	187	179	10.2	7.7	7.07	20.0	7 4	17.5	8	141
payb3	000	00	96	1	207	108	151	155	13.3	10.3	13.7	4.2	2	1.0	2 -	0.4
010	201	000	000	+	108	133	181	171	12.0		10.5	10.4	7.6		0.	10.1
p3	103	200	100	+	200	118	157	154	10.3	8.2	9.7	9.4	19.1	15.2	19.2	0.71
Ps	106	16	18	+	000	100	700	171	9.7		8.7	8.3	16.9	14.9	18.1	16.6
p4	104	95	16	+	213	200	147	143	10.5		8.6	9.7	23.6	18.8	23.7	22.0
P ₄ xp ₆	110	96	100	+	7/1	0 0	1423	157	13.4	7.9	11.9	11.1	9.7	4.9	7.4	7.3
D4	113	104	101	+	617	90	200	101	111		143	12.0	9.8	16.1	11.3	12.4
P,xD7	112	86	104	-	225	154	183	1 2	128	D A	110	113	10.1	11.0	12.8	11.3
d	109	95	98		224	127	1/4	0	2.00		10 8		4.0	3.2	3.1	3.4
200	07	87	93		214	128	179	1/4	4.0	0.0	200	107	u u	48	116	7.3
108		80	85	+	207	117	166	164	14.1		0.2	1 0	20.0		13.1	12
, D4		8 8	000	-	187	101	137	142	12.3	6.6	10.5	0.0	1.7	7.1.7	0 0	12
p4xp ₉		36	100	+	213	129	172	171	11.1	6.6	9.4	10.1	14.5	13.4	12.3	4.0
cp4		33	701	+	700	188	199	198	11.8	8.6	10.1	10.2		27.8		25.8
D10		96	200	101	177	200	144	158	102	8.4	10.1	9.6	15.5	5.8	9.3	10.2
2	100	98	102	_	7117	111	-	200			-		THE REAL PROPERTY AND ADDRESS OF THE PERSONS ASSESSMENT ASSESSMENT ASS	PERSONAL RESIDENCE AND ADDRESS OF	CONTRACTOR OF THE PERSON NAMED IN	

Table 2 continued:

10.3 11.5 16.0 9.9 10.7 13.9 5.0 13.7 23.3 11.4 14.0 14.2 6.5 6.5 2.8 2.8 15.4 17.8 17.8 16.1 18.4 8.6 13.0 4.7 8.3 8.3 18.9 19.4 8.0 6.5 9.4 15.0 10.8 4.6 10.1 14.6 9.7 14.4 4.0 13.6 17.4 3.9 18.4 13.9 13.6 10.7 18.8 17.0 5.1 10.4 8.6 9.7 11.4 10.9 11.3 13.0 12.4 12.5 10.4 9.0 9.7 9.4 10.1 12.2 10.8 9.2 9.6 10.6 12.3 9.6 8.3 12.5 12.1 12.8 12.2 11.0 12.5 11.5 10.5 8.6 7.7 10.1 10.1 7.7 7.7 7.6 9.5 11.6 10.1 10.6 10.4 11.1 9.2 10.7 14.5 12.3 13.0 12.2 9.1 14.0 14.1 13.8 14.6 9.9 1135 105 135 135 135 105 110 110 1110 213 215 183 183 220 220 220 220 190 190 190 190 217 195 219 219 220 220 178 178 181 181 163 111 117 117 120 120 96 99 101 P₅xp₁₀ P₆XP₁₀ Psxps P₁₀XP₅ P7XP10 p₉xp₆ Proxpe P₁₀XP₇ p₁₀Xp₉ p₈xp₅ p₇xp₆ Pexpa pexpe Pexpa P₇xp₈ P₈xp₇ P₉Xp₇

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38.0 39.8 43.2 36 40.1 39.3 38.2 43.3 45.1 43.2 43.1 38.3 38.1 43.1 39.5 42.8 40.0 37.8 38.5 44.8 43.0 38.1 41.7 40.4 35.9 42 48.0 48.1 38.3 46.2 43.1 40.4 39.3 41.2 45.2 45.0 48.1 37.2 39.3 48.3 48.2 44.1 42.1 42 35. 52.9 46.0 42.4 39.7 39.2 63.8 48.9 85.6 36.2 27.3 56.4 46.8 30.0 42.0 47.7 48.3 46.7 32.8 42.5 63.4 49.7 36.9 26.7 65.1 47.0 29.2 29.2 42.4 40.9 35.2 17.8 58.3 19.8 31.6 53.7 25.4 25.3 16.6 54.6 34.5 32. 57.4 64.2 38.1 81.4 81.4 81.4 81.4 74.4 776.3 39.5 76.3 60.3 67.3 59.1 40.7 52.6 57.5 54.9 49.4 58.7 420 269 269 269 326 326 383 383 383 467 467 567 214 E, 586 499 291 565 426 204 280 565 141 554 387 190 470 950 487 424 424 902 785 734 476 907 565 644 878 392 611 633 887 463 674 487 E 3.9 3.6 3.8 3.5 3.2 3.6 Table 2 continued: E. 3.8 2.9 3.7 3.7 3.7 3.8 3.8 3.8 3.8 2 2 3 4 2 2 2 3 4 8 9 8 4 8 9 3.6 DIXPID PIXPE Parpi P₁XP₇ P-xp1 pixps pixp. Poxp. PigXpi P₃xp₂ p2Xp4 P₄xp; pexp2 P₆xp₂ Psxp1

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44.8 39.0 42.9 38.5 42.0 43.8 42.9 45.1 44.8 46.3 46.5 39.7 16.7 44.7 40 38.2 37.8 41.8 41.5 42.8 44.9 37.1 41.2 6 38. 44 44 43.2 45.9 44.1 39.3 40.3 43.2 41.2 45.1 48.1 47.1 43.2 42.2 44.8 45.4 47.1 47.3 48.0 46. 48.1 44.2 41.3 58.9 57.5 61.4 55.2 50.3 54.5 35.8 63.4 32.5 50.8 42.5 61.0 46.9 49.6 39.4 42.4 38 55.0 48.5 54.2 37.4 59.5 57.3 61.5 33.8 55.1 40.8 66.3 39.8 52.1 E 49 39 42 48.0 47.7 445.5 39.0 46.3 46.3 33.9 40.6 40.6 38.5 57.4 57.0 64.6 65.1 63.5 66.0 66.0 76.2 776.2 57.3 57.3 50.3 71. 53. 45. 67 56. 48 345 507 501 706 706 161 715 598 676 556 459 334 334 342 342 383 386 760 760 760 550 121 747 527 527 387 339 653 374 665 557 590 456 344 574 574 553 558 489 567 320 5 1056 386 797 719 719 648 647 792 686 686 685 681 1083 496 995 724 596 29.4 31.4 31.4 26.1 27.2 27.7 27.7 27.6 27.6 25.4 27.6 27.6 26.8 26.8 25.1 28.3 27.2 30.8 34.3 34.3 33.6 25.8 25.9 25.9 00 28. 28. 3.0 3.0 4.1 3.6 2.8 3.5 6 3.6 3.1 3.6 3 able 2 continued: E2 3.1 3.1 4.0 4.4 4.4 4.4 4.4 2.9 3.4 3.7 3.6 2.2 3.3 7 2.7 2.7 2.9 2.9 2.9 2.5 2.9 3.8 P10XP3 p₆xp₃ p₇xp₃ P₃xp₈ P3XP10 p₄xp₃ p₅xp₃ p₆xp₄

45.2 40.5 42.1 42.6 38.0 43.8 39.8 44 39 40.6 40.0 43.2 44.2 41.3 41.2 42.4 44.5 43.1 40.1 38.1 42.1 40.1 42. 40.0 8 6 42.0 42.2 37.9 41.2 40.4 42.3 40.2 46.1 41.1 42.3 42.2 41.4 40 36. 39 36 38 38 45.1 42.0 42.4 47.3 43.1 47.0 42.2 44.1 45.2 44.7 42.0 46.3 44.1 45 48 42.1 48 45 38 44 43 43. 59.2 59.2 54.5 54.5 51.0 54.5 30.7 61.5 62.2 2 0 32.47 40 43.0 46 48. 35 45 50 60 43.2 67.1 52.4 55.9 39.4 50.2 58.5 42.3 54.1 29.9 34.5 61.0 39.2 62.0 0.09 46.5 45.7 48 20.9 36.8 31.5 36.4 28.3 30.6 54.5 41.7 44.3 28.3 41.6 32.6 45.4 31.3 37.4 40.8 51.5 52.6 35.8 48.6 25.7 50.7 49.7 41.4 0 72.1 50.4 72.0 78. 63 59. 63. 58 46. 61 54 57 615 320 320 527 383 383 559 640 510 606 500 500 709 361 605 253 519 655 456 409 686 786 588 471 070 590 249 427 376 431 793 488 344 581 N.siVpl. 635 165 E₂ 437 2217 289 289 455 360 391 320 320 597 521 521 186 442 328 563 546 629 560 388 452 1042 916 547 708 966 734 444 704 326 525 609 582 E₂ 25.4 31.0 25.7 25.7 19.9 32.5 27.7 29.5 24.4 29.6 28.2 23.5 24.4 26.9 226.9 225.5 227.7 225.5 225.2 226.4 227.7 226.4 22 19.3 S.W.gms. 2.7 2.9 2.9 2.9 3.1 3.1 4.2 00 0 2.6 00 1 8 3.1 3 2 continued: 000 4.2 2 2 3 3 3 4 5 5 5 5 5 5 5 3.4 3.3 23.23.99.89.49.90.23.00.2 3.1 3.6 2 9 3.4 8 0 3.6 3.3 3.6 2.7 2.8 3.6 3.1 H a a a Table P7xp10 P₆xP₁₀ P₆xp₁₀ P₁₀xP₅ P₁₀XP₆ P₇xp₈ P₁₀XP₇ P₆xp₇ P₆xp₆ P₇xp₉ P₉xp₇ P_sxp_e p_exp₅ P_exp₇ p₇xp₅ PsxpB p_uxp_s P₆xp₉ P₆xp₆ 9dx6d P₈Xp₇

Weight of 1000 seeds in grams (1000S.W.gms.) varied from one hybrid to another. The hybrid $P_7 \times P_1$ gave the lightest weight of 1.9 gms. at E_1 . On the other hand, the hybrid $P_5 \times P_{10}$ gave the heaviest weight of 4.6 gms. at E_3 . It appeared that fertile soil yielded lighter seed weight than poor soils with high level of salinity. These results were clearly apparent for all F_1 hybrids and their reciprocals where heavier seed weight was obtained at poor soils. The combined data showed that the hybrid $P_5 \times P_{10}$ had the heaviest seeds with the weight of 4.1 gms.. The same hybrid was also the heaviest hybrid for weight of seed at the three environments. The two F_1 hybrids $P_7 \times P_{10}$ had the lightest seed weight of 2.5 gms..

Number of seeds per siliqua (N.se./sil.) varied not only from one hybrid to another but also from one environment to another for the same hybrid. The hybrid between $P_7 \times P_6$ gave the smaller number of seeds of 19.3 seeds at E_2 . On the other hand, the hybrid between $P_2 \times P_9$ gave the larger number of seeds of 35.5 seeds at E_1 . The combined data showed that the hybrid $P_5 \times P_8$ had the larger number of seeds of 32.9 seeds, while the hybrid

P₇ x P₆ had the fewer number of seeds of 22 seeds.

Number of siliqua per plant (N.sil./pl.) varied not only from one hybrid to another but also from one environment to another for the same hybrid. The hybrid between $P_1 \times P_{10}$ gave the fewer number of siliqua of 141 siliqua at E_2 . On the other hand, the hybrid $P_{10} \times P_7$ gave the larger number of siliqua of 1086 siliqua at E_1 . The combined data showed that the hybrid $P_7 \times P_6$ had the largest number of siliqua of 1051 siliqua per plant, while the hybrid $P_6 \times P_{10}$ had the fewest number of siliqua of 253 siliqua.

Seed yield per plant in grams (S.Y/pl. gms.) varied not only from one hybrid to another but also from one environment to another for the same hybrid. The hybrid $P_1 \times P_{10}$ gave the lowest yield of 16.6 gms. for this trait at E_2 . On the other hand, the hybrid $P_1 \times P_6$ gave the highest yield of 81.4 gms. at E_1 . The combined data showed that the hybrid $P_1 \times P_6$ had the highest yield per plant of 69.7 gms., while the hybrid $P_1 \times P_{10}$ had the lowest yield of 27.3 gms. per plant.

Oil percent (Oil %) varied not only from one hybrid to another but also from one environment to another for the same hybrid. The hybrid $P_1 \times P_8$ gave the lowest percent ratio of (35.1) oil % at E_1 . On the other hand, the hybrid $P_5 \times P_2$ gave the highest percent ration of 48.3 % oil at E_1 . The combined data showed that the hybrid $P_4 \times P_7$ produced more oil % of 46.7 %, while the hybrid $P_1 \times P_8$ produced less oil of 36.7 %.

The results obtained for yield and yield component traits showed that N.se./sil. and N.sil./pl. were very effective in determination S.Y./pl. in grams, while 1000 S.W.gms. was less effective. Therefore, selection for more N.sil./pl. and more N.se./sil. would increase yield. In this connection similar results obtained by Kandil *et al.* (1996), Halaka (2000), Teilep (2003) and Kassab (2004).

Correlation coefficient measures the power of association between any two traits. When the correlation coefficient is positive and significant it indicates that the increase in one trait would accordingly increase the other. The analyses of covariance in addition to the analyses of variance of all pairs of traits would result in the estimation of phenotypic (r_{ph}) and genotypic (r_{q})

correlations between these pairs of traits. It is usually expected that the magnitude of r_{ph} correlation is larger than its corresponding estimate of r_g correlation. This is true because the environmental factors affecting the r_{ph} correlation more than the r_g correlation.

In the present investigation, the r_{ph} and r_{g} correlations were separately estimated for the parents and the F_{1} hybrids. The results of these estimates are presented in Tables 3 and 4 for parents and hybrids, respectively.

Phenotypic and genotypic correlations within the parental lines:

For parental lines, the results showed variable directions of correlations since some estimates were negative and the others were positive. Most of these estimates were insignificantly positive or negative with except the rph and ra correlation values obtained between plant height (P.H.cms.) and height of the first branch (H.f.br.cms.), 1000 seed weight in grams (1000S.W.gms.) and each of number of siliqua per plant (N.sil./pl.) and seed yield per plant in grams (S.Y./pl. gms.) and finally between number of siliqua per plant (N.sil./pl.) and seed yield per plant in grams (S.Y./pl. gms.). The values of rob and ro correlation coefficients between those pairs of traits reported earlier were positive and significant. This finding indicated the presence of high association between them. The positive and significant rep and ra correlations between P.H. cms. and H.f.br. cms. indicated that the height of first branch started higher from the surface of the soil, when the plant was tall. Although the direction of r_{ph} and r_{q} correlations were positive, it is undesirable because, in this case, plants would give less number of primary branches per plant. This result was insured by the presence of negative roh and ro correlations between P.H. cms. and N.pr.br./pl..

The most desirable r_{ph} and r_g correlations were obtained between N.sil./pl. and S.Y./pl. gms. and between 1000 S.W.gms. and S.Y./pl. gms. which showed positive and significant values. Therefore, these three traits showed a desirable association. This finding indicated that selection program to increase one trait would, indeed increase the others. It is very important to indicate that S.Y./pl. gms was positively correlated with oil %, although it was not significant. This results is economically desirable because the increase in

both S.Y./pl. gms. and in oil % would increase the final oil yield.

It should be mentioned that all negative r_{ph} and r_g correlations were insignificant indicating that this type of association was not important specially the values of r_{ph} and r_g correlations were very small. The negative r_{ph} and r_g correlations between 1000 S.W. gms and N.se./sil. indicating that when siliqua had more seeds, the size and the weight of seeds become lighter.

In general, it could be concluded that S.Y./pl. gms is a function of N.sil./pl. and 1000 S.W. gms.. The selection for heavier seed and more N.sil./pl. would, indeed, increase the total seed yield and therefore, the total oil yield. Phenotypic and genotypic correlations within hybrids:

Table 3: Phenotypic (r_{ph}) and genotypic (r_g) correlation between pairs of all studied traits for the 10 parents obtained from the combined data over three environments.

Traits	Corr.	P.H.	N.pr.br. /pl.	H.f.br.	1000 s.w.	N.seed/ sil.	N. sil./pl.	S.Y./ pl.	Oil %
D.50%f.	rph	0.445	-0.070	0.463	-0.487	0.195	-0.022	-0.288	0.158
	ra	0.533	-0.073	0.506	0677	0.280	-0.013	-0.308	0.195
P.H.	Γph		-0.023	0.86**	-0.047	0.119	-0.093	0.077	-0.102
Cms	rg		-0.026	0.91**	0.021	0.093	-0.068	0.069	-0.132
N.pri.	Γph			0.274	0.183	-0.312	0.028	0.061	-0.254
bra./pl.	rq			0.297	0.279	-0.429	-0.002	0.061	-0.261
H.f.bra.	Γph				-0.254	0.307	-0.019	-0.011	-0.266
	Γg				-0.308	0.399	-0.018	0.002	-0.255
1000 s.w.	Γph				Carrier a - 11 -	-0.395	0.314	0.663**	0.323
	rg					-0.628	0.941**	0.880**	0.380
N.seed/	Γph						-0.456	-0.158	0.153
sil.	ra						-0.587	-0.207	0.241
N. sil./pl.	Γph							0.807**	0.201
	rg							0.950**	0.252
S.Y./pl	rph								0.360
	Γg								0.348

^{*} Significant at 5 % level.

Table 4: Phenotypic (r_{ph}) and genotypic (r_g) correlation between pairs of all studied traits for F_1 hybrids and F_{1r} reciprocal hybrids obtained from the combined data over three environments.

Trait	Corr.	P.H.	N.pr.br.	H.f.br.	1000	N.seed/	N.	S.Y./	Oil %
		cms	/pl.		s.w.	sil.	sil./pl.	pl.	
D.50%f.	rph	0.062	0.027	0.071	0.024	-0.096	0.044	0.016	0.014
	ra	0.059	0.027	0.070	0.027	-0.101	0.045	0.016	0.011
P.H.	rph		0.166	0.067	-0.024	-0.095	0.087	0.061	0.122
Cms	ra		0.166	0.061	-0.019	-0.085	0.082	0.062	0.120
N.pri.	roh			-0.010	0.023	-0.023	0.044	0.056	0.034
bra./pl.	ra			-0.004	0.015	-0.029	0.051	0.057	0.037
H.f.bra.	rph				-0.036	-0.076	-0.031	-0.063	0.003
	ra				-0.039	-0.088	-0.029	-0.065	0.005
1000 s.w.	Γ _{ph}					0.086	-0.185	0.021	-0.094
	rg					0.078	-0.118	0.018	-0.095
N.seed/	rph						-0.171	-0.019	0.008
sil.	ra						-0.128	-0.019	0.003
N. sil./pl.	rph							0.289**	0.016
tolica Salesy (1991)	ra							0.289**	0.017
S.Y./pl	rph								-0.016
	rg								-0.017

^{**} Significant at 1 % level.

The results showed variable directions for correlations, since some estimates were negative and some others were positive. The most desirable r_{ph} and r_g correlations were obtained between S.Y./pl. gms. and N.sil./pl. which showed positive and significant values. Therefore, these two traits showed a desirable association indicating that selection to increase one trait would, indeed, increase the other. It should be mention that all negative r_{ph} and r_g correlations were insignificant indicating that this type of association was not important, the values of r_{ph} and r_g correlations were small. The negative r_{ph} and r_g correlations between 1000 S.W. gms. and N.sil./pl.

^{**} Significant at 1 % level.

^{*} Significant at 5 % level.

indicated that: when plant had more siliqua, the size and weight of seeds become less.

The results of this investigation for r_{ph} and r_g were different from the results obtained by Sharief and Keshta (2002), Marinkovic *et al.* (2003), Teilep (2003) and Kassab (2004). On the other hand, the results obtained by El-Baz and El-Shakhess (2001) and Maria *et al.* (2003) were very close to the results obtained in this investigation.

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سلوك الآباء والهجن بينها وطبيعة الارتباط للصفات الاقتصادية في الكانولا عصلى ماهر العصدل*، أشرف حسسين عبد الهادى*، معاطى معاطى قسطة**، السيد فهمى العزونى**

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** قسم بحوث المحاصيل الزيتية - معهد بحوث المحاصيل الحقلية - مركز البحوث الزراعية

تعتبر الكانولا من المحاصيل الجديدة في مصر والتي تزرع من أجل الحصول على الزيت من بنورها حيث تحتوى على نسبة عالية من الزيت (٤٠% - ٤٥%) تقريبا ، ونظرا لكونها محصولا جديدا فإن الأبحاث التي أجريت عليها بغرض التحسين الوراثي قليلة ، ولذا كانت هذه الدراسة إحدى الخطوات المهمة لمد النقص في هذا الاتجاه.

استخدم في هذا البحث ١٠ سلالات نقية من الكانولا تم الحصول عليها من قسم بحسوث المحاصيل الزيتية استخدم في هذا البحث ١٠ سلالات نقية من الكانولا تم الحصول عليها من قسم بحسوثة وهمي: سرو؟ بمحطة البحوث الزراعية بالسرو - مركز البحوث الزراعية ، وهي سبع سلالات مصرية وهمي: سرو؟ (P1) ، سرو ٢٠ (P5) ، سرو ٢٠ (P5) ، سرو ٢٠ (P5) ، سرو ٢٠ (P5) ، ستورد ٢٦ (P5) ، مستورد وهي: مستورد وهي: مستورد وهي: مستورد وهي: مستورد وهي المستورد وهي المستورد وهي المستورد وهي المتحدل عليه المستورد وهي: الجمالية (E1) ، مستورد ولا الأول والآباء في ثلاث بيئات مختلفة وهمي: الجمالية (E1) ٢٠٠٢/ ٢٠٠٢ ، السرو (E3) ٢٠٠٢ / ٢٠٠٢ ، السرو (E3) ٢٠٠٢ / ٢٠٠٢

- وجود تباينات عالية بين التراكيب الو راثية المختلفة .

كانت السلالة ٣٢٥ ((P_8) هي الأكثر تبكيرا ، بينما السلالة سرو (P_2) أفضل الأباء في ارتفاع النبات وعدد البذور/ القرن ، والسلالة سرو (P_3) الأكبر في عدد الغروع على النبات ، بينما السلالة سرو (P_1) افضل محصول / النبات بالجرام ، و السلالة ١٦٤ ((P_1)) الأفضل في وزن الألف بنزرة ونسبة الزيت. وكان الهجين بين $(P_1 \times P_2)$ أفضل الهجن في نسبة الزيت ، بينما الهجين بين $(P_1 \times P_3)$ الأقل في نسبة الزيت.

وقد أظهرت الدراسة وجود ارتباط مظهري ووراثي موجب وعالي المعنوية بين ارتفاع النبات/ السم وارتفاع أول فرع بالسنتيمتر وكذلك بين وزن ال١٠٠٠ بذرة وكل من عدد القرون/ النبات ومحصول البذور/النبات بالجرام لسلالات الأباء. كذلك وجد ارتباط وراثي ومظهري موجب وعالي المعنوية بين عدد القرون/ النبات ومحصول البذور/ النبات بالجرام للهجن.