# INHERITANCE OF RESISTANCE TO COTTON LEAFWORM (Spodoptera littoralis) IN SOYBEAN (Clycine max)

Habeeb, M.B.<sup>1</sup>; M.I.A. Sherif<sup>2</sup>; A.A. El-Hosary<sup>3</sup>; M.A. Naseeb<sup>1</sup> and A.M. Hablas<sup>2</sup>

- 1) Agriculture Research Centre, Cairo.
- 2) Department of Genetics, Faculty of Agriculture, Minuaiya Univ.
- 3) Department of Agronomy, Faculty of Agriculture, Moshtohor.

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مركز البحوث الزراعية \_ بالقاهــــــرة قسم الوراغة \_ كلية الزراعة \_ جامعة المنوفيــة قسم المحاصيل \_ كلية الزراعة \_ مشـــتهـــر قسم الوراغة \_ كلية الزراعة \_ جامعة المنوفيــة مركز البحوث الزراعية \_ القاهــــــرة

محمود بسطویسی حبیب ب محمد ابراهیم عبد الباری شریف علی عبد المقصود الحصری عبد المنعم محمد حبلب محمد عبد الله نصیب

# ملخص البحث

أستخدم في دراسة المقاومة لدودة ورق القطن الصنفين عاليا المقاومة للال Crawford والصنف التجاري الحساس Crawford من خلال كلاقة معايير للصغة وهذه المعايير هي عدد الشعيرات والمساحة المستهلكة من الورقة ونسبة الفقد في الأوراق وقصد أستخدم الهجينيسن الورقة ونسبة الفقد في الأوراق وقصد أستخدم الهجينيسن الانعزالية الناتجة عنها في تقدير التقديرات الوراثية المختلفة والمبنت النتائج المستحمل عليها أن قوة الهجين كانت سالبة وعالية المعنوية بالنسبة لكل من المساحة المستهلكة من الورقة ونسبة الفقد في الأوراق وبينما كانت موجسة وعالية المعنوية بالنسبة لعدد الشعيرات وذلك في كلا الهجينين وقد كان التدهور الناتج عن التربية الداخلية معنوي بالنسبة لعدد الشعيرات والمساحة المستهلكة في كلا الهجينين وأيضا لنسبة الفقد في الأوراق في الهجيسن النسبة لدليل السيادة وقد كان التأثير المستهلكة في كلا الهجينين وأيضا لنسبة الفقد في الأوراق في الهجيسن والنسبة لطبيعة فعل الهين كان التأثير المضيف والسيادي نو أهمية في وراشة وبالنسبة لطبيعة فعل الهين كان التأثير المضيف والسيادي نو أهمية في وراشة وبالنسبة لطبيعة فعل الهين كان التأثير المضيف والسيادي نو أهمية في وراشة

الثلاثة معايير للصغة ، بينما ظهرت بالاضافة لذلك معنوية تأثير الهيـــــن التغوقي بالنسبة لعدد الشعيرات والمساحة المستهلكة في الورقة ، بين مغامل الاختلاف الورائي فيما متوسطه بالنسبة للمساحة المستهلكة من الورقة ونســبة الفقد في الأوراق بينما أظهر قيما منخفضة بالنسبة لعدد الشعيرات ، وكانــت قيم المكافئ الورائي بالمفهوم العام تتراوح بين قيما عالية وقيما متوسطة فــــ الثلاث صفات ، بينما كانت القيم بالمعنى الضيق عالية في حالة عــــد الشعيرات ونسبة الفقد في الأرراق في الهجين الثاني ولكنها كانت منخفضة في المالات الأخرى ، وبالنسبة لنسبة الفقد في الأوراق في الهجين الثانـــي بينما كانت القيم متوسطة في حالة المساحة المستهلكة من الورقة في كــــلا الهجينين وفي نسبة الفقد في الأوراق في الهجين الأول ، أما في حالة عـدد الشعيرات فقد كانت القيم منخفضة في كلا الهجينين ،

#### ABSTRACT

Two highly resitant varieties D19-10455 and Celest; and the susceptible commercial variety Crawford were used to study resistance of soybean to leafworm through three criteria; i.e. hairiness; area consumed and defoliation. The two crosses D19-10455 x Crawford (I) and Celest x Crawford (II) and their segregating generations were used to estimate the different genetic parameters. Results obtained showed highly significant negative heterotic effects for leaf area consumed and defoliation in the two crosses, whereas highly significant positive heterotic effects were detected for hairiness and leaf area consumed in both crosses and in the second cross for defoliation. Values of potence ratio indicated partial deminance for resistance over susceptability and for high number of hairiness over low number. Both additive and dominance types of gene action were important in the inheritance of the three traits; although the epistatic gene effects had important attributes of inheritance with regard to hairiness and leaf area consumed in the two crosses. Genetic coefficient of variation expressed moderate values for leaf area consumed and defoliation, while low values were expressed for hairiness. High to moderate values for broad sense heritability were detected for the three traits. High values of narrow sense heritability were only obtained for hairiness and defoliation in the second cross. The expected genetic advance was higher for defoliation in the second cross, but it was moderate for leaf area consumed in both crosses and defoliation in the first cross. However, low estimates were obtained for hairiness in both crosses.

#### INTRODUCTION

Soybean is susceptible to several diseases and insects. One of the most harmful insects is cotton leafworm, Spodoptera littoralis (Boisd). This pest was found in all soybean growing areas in Egypt. Cotton leafworm becomes widespread and serious. So, it is imperative to breed varieties having a high degree of resistance to cotton leafworm. Currently, varieties such as Crawford group IV appear to be susceptible to leaf-feeding insects such as cotton leafworm. Therefore, insect control will be difficult and costly. However, several studies with cotton leafworm have demonstrated consistently higher level of resistance in Celest and D79-10455 under field and laboratory conditions. The objective of the present investigation is to study the inheritance of resistance to cotton leafworm insect in soybean.

#### MATERIALS AND METHODS

The present investigation was carried out at the Sakha Agriculture Research Station, during the three successive seasons, 1984, 1985 and 1986. Three varieties of soybean were used in this worm. Two of them were highly resistant to leafworm insect, i.e. D19-10455 and Celest and the other one was susceptible i.e. Crawford (Commercial variety).

In 1984 season, the three varieties were sown in four planting dates to avoid differences in flowering time and to secure enough hybrid seeds. During this season, two crosses were made as follows: D19-10455 x Crawford (I) and Celest x Crawford (II).

In the 1985 season, the hybrid seeds and the three parents were sown.  $F_1$  plants were self-pollinated and backcrossed to both parents to obtain  $F_2$ 's and the backcrosses seeds, also were crosses repeated between parents to obtain the  $F_1$  seeds.

In the 1986 season an experiment for each of the two crosses was carried out. Each trial contained six populations, i.e., the two parents, their  $F_1$ ,  $F_2$  and the two backcrosses to each parent. The experimental design was randomized complete block design with three replications. Each plot consisted of one ridge 300 cm long and 60 cm. wide. Hills were spaced 20 cm. apart with one seed per hill. Each replicate consisted of 15 plots per hill. Each replicate consisted of 15 plots for  $F_2$  plants, five plots for each of BC<sub>1</sub> and BC 2 and two plots for any nonsegregating population.

Data were taken only on guarded plants. The study of the inheritance of resistance to cotton leafworm was evaluated using the following three criteria:

- Hairiness: Number of hairs on the lower surface of the leaf were counted, and the leaflet from first trifoliolate of the upper third of plant was taken as a standard (Kamel, 1963). The binocular field (0.5 cm²) was used to determine the average number of hairs for each plant.
- 2. The area of leaf tissue consumed: One fourth instar stage larvae were placed in a glass container, 1000 ml. capacity and were allowed to feed on fresh leaflets, excised randomly from the upper third of each plant, including their petioles. The area of leaf tissue consumed after 24 hours was measured (Thobbi, 1962 and Meisner et al., 1983).
- 3. Leaf feeding damage or foliage loss (defoliation): Visual ratings of percent defoliation were recorded, as the average of three times (every seven days) beginning two weeks after flowering, on each plant in the field experiment without insect control under the natural field infection. A standard area diagram for estimating the percentage of defoliation was reported by Smith and Brim, 1979 as shown in Figure (1).

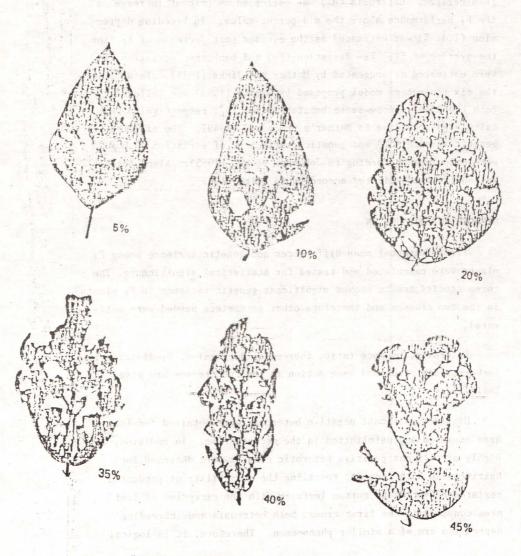


Fig. (1): Standard area diagram estimating the percentage of defoliation.

The genetic variance within  $F_2$  population was firstly evaluated. If that variance is significant, various genetical parameters were them derived. Heterosis (H%) was expressed as percent increase of the  $F_1$  performance above the mid-parent value. In breeding depression (I.d. %) was estimated as the average part decrease of  $F_2$  from the average of  $F_1$ .  $F_2$ - deviation ( $E_1$ ) and backcross deviation ( $E_2$ ) were estimated as suggested by Mather and Jinks (1971). In addition, the six parameters model proposed by Gamble (1962) was followed. Both broad and narrow-sense heritabilities ( $h_b^2$ , respectively) were calculated according to Mather's procedure (1949). The expected genetic advance ( $\Delta G$ ) and genetic coefficient of variation (G.C.V.%) were calculated according to Johanson et al. (1955). Also, potenc ration (P) was estimated according to Smith (1952).

#### RESULTS AND DISCUSSION

First, parental mean differences and genetic variance among  $F_2$  plants were calculated and tested for statistical significance. The three studied traits showed significant genetic variance in  $F_2$  plants in the two crosses and therefore other parameters needed were estimated.

Heterosis, potence ratio, inbreeding depression,  $F_2$ -deviation, backcross deviation and gene action in the two crosses are given in Table (1).

Highly significant negative heterosis were obtained for leaf area consumed and defoliation in the second cross. In addition, highly significant positive heterotic effects were detected for hairiness in both crosses, revealing the possibility of producing resistance hybrids to cotton leafworm with the exception of leaf area consumed in the first cross, both heterosis and inbreeding depression are of a similar phenomenon. Therefore, it is logical

Table (1): Heterosis, inbreeding depression, potence ratio, F2 deviation, backcross deviation and gene action for resistance to cotton leafworm in the two crosses D79-10455 X Grawford (I) and Celest X Crawford (II).

	9	Heterosis	Inbreeding	Potence	7.	٠ ١	Gene action six parameters (Gamble procedure)	rameters (	sample proc	equire)
Character	Cross	% ************************************	depression ratio deviation $E_1$	ratio	deviation El	deviation E <sub>2</sub>		aa	ad	48 PP
Hairiness	I Tail	3.50**	4.54**	0.18**	- 0.32**	0.62**	10.72**- 1.64** 2.9	0** 2.52	** 0.44	- 3.76 **
	(882 () E <sub>19</sub>	** 6.04	5.73	0.20	0.20 - 0.39	- 0.27	0,20 0,39 0,27 12,67 2,89 1,79 1,02 1,00 0,40	** 1.02	1.00	0.40
Leaf area	rdoT-	10.11 -1.31	**59.4	0.07	2.02**	- 6.58**	53.11** 5.40**-21.9	2** -21.24*	* - 4.28**	34.39**
consumed	nors v	-32.64	-47.03	** 89.0-	6.19	- 0.99	- 0.99 39.92 ** 21.65 -39.90 -26.74 ** 2.28 29.71 **	** -26.74	2.28 29.71	29.71
Defoliation	d bands	- 2.34	- 3.59	- 0.29	0.79	- 0.18	34.30 ** 2.:0 ** 4.32 * - 3.52	2* - 3.52	- 0.34 3.57	1.87
	i de	-21.90	-19.97	- 0.55	1.41	- 0.77	25.29 ** 10.71 ** - 9.	71 - 4.10	- 0.54	2,57

\* Significan at 5% level.

\*\* Significant

to anticipate that heterosis in the  $F_1$  will be followed by appreciable reduction in the  $F_2$  performance. Leaf area consumed in the first cross, did not showed any heterosis, however, a highly significant value for inbreeding depression was detected. The difference most parameters may lead to the observed absence of heterotic effects (Marani, 1968).

It could be concluded that heterosis and inbreeding depression effects affecting the resistance of cotton leafworm, refer to the existense of some non-additive genetic types of gene action. Lambert and Rilen, 1984 showed that here is a strong trend for larvae of different insects reared on soybean F<sub>1</sub> plants of the crosses to be smaller than the mid-parent values. Also, they mentioned that mortality tended to be greater for larvae reared on resistant geneotypes, and larval weight for those reared on F<sub>1</sub> plants tended to be similar to those reared on the most resistant parents.

Nature of gene action was also studied according to the relationships illustrated by Gamble (1962). For estimating various parameters of gene effects, the variety with larger mean in each trait was usually considered as P<sub>1</sub>. In all traits, the mean effect of parameters "m" was highly significant.

The additive genetic estimates were highly significantly negative for hairiness and positive for other traits (Table 1). These results indicate the potentiality of improving the performance of these traits by using pedigree seleciton program. Similar results were obtained by Abou-Tour (1986) for leaf area consumed, who detected additive effects through generation mean analyses. The estimates of dominance effects were significantly positive for hairiness and significantly negative for the other traits, it can be concluded that the dominance gene effects are effective in the inheritance of all traits. Similar conclusion was obtained by Abou-Tour (1986), and Kornegay and Temple (1986) for leaf-feeding ratings. On the other hand, Lambert and Kilen

(1984) found that there was a slight dominance effect for resistance.

Hairiness in the first cross exhibited significant estimates for the three types of digenic epistasis although the significant estimates were experessed for "aa" and "ad" only in the second cross. Also, the significant values of various types of epistatic effect were accompanied by highly significant estimates for  $E_1$  and  $E_2$  (Table 1). Therefore it can be said that the three types of gene action are important contributors to this trait.

For leaf area consumed, the significant estimates were obtained for the three types of epistatic effects in the first cross, but of different magnitudes and directions. It is interesting to note that if the estimated values for dominance and various epistatic parameters were of nearly equal magnitudes and opposite directions, no appreciable heterotic effect can be detected (Marani, 1968 and El-Hosary, 1983). In the II cross, significant values were detected for additive x additive and dominance x dominance types of epistasis. It is noteworthy that both types of epistasis were accompanied by significant estimates of E1. Also, the heterotic effect previously obtained may be due to both dominance and epistatic gene action.

For defoliation, insignificant estimates were detected for three types of epistasis. It is noteworthy that the three epistatic types were accompanied by insignificant estimates for E<sub>2</sub> and E<sub>1</sub> heterotic effect in the first cross; and E in the second cross.

In both crosses, genetic coefficient of variation was of moderate values for leaf area consumed and defoliation (Table 2), while the hairiness had low values of G.C.V.%. Therefore, it is impossible to estimate the magnitude of heritable variation, when figured out with help of other heritability and the genetic advance (Swarup and Changale, 1962).

Table (2): Heritability percentage in broad and narrow sense, genetic coefficient of variation for resistance to cotton leafworm in two crosses, D<sub>79-10455</sub> X Crawford (I) and Celest X Crawford (II).

Cross	h' broad %	h' narrow	GS	GS %	G.C.V.
mt iso Co	egentinale va	3- 2- 2- 2- 2- 2- 2- 2- 2- 2- 2- 2- 2- 2-	35,74 1101	of a	
I	91.55	31.82	0.69	6.44	9.36
II	90.62	79.78	2.19	17.28	10.03
-,	ins end. 1784	es (syrecolo	100 200	ay ros	
I	92.94	28.47	6.16	11.60	19.06
II	70.98	18.43	4.59	11.50	25.50
1170 102	artiels of th				
1		-5 500 04.		high all his	
I	58.54	32.89	4.13	12.83	14.50
II	87.22	74.42	11.97	61.76	37.63
	I II II II	I 91.55 II 90.62  I 92.94 II 70.98	I 91.55 31.82 II 90.62 79.78  I 92.94 28.47 II 70.98 18.43	I 91.55 31.82 0.69 II 90.62 79.78 2.19  I 92.94 28.47 6.16 II 70.98 18.43 4.59	I 91.55 31.82 0.69 6.44 II 90.62 79.78 2.19 17.28  I 92.94 28.47 6.16 11.60 II 70.98 18.43 4.59 11.50

High heritability values in broad sense were detected in both crosses for all traits (Table 2). Poze (1974) found high heritability estimates in soybean for resistance to Epilachna varrestis. For hairiness and defoliation high values for narrow sense heritability were obtained in the second cross, which are nearly equal with their corresponding broad sense ones. For the other cases high heritability in the broad sense was accompanied by low value for narrow ones. This result revealed that the non-additive genetic variance has a great role in the existence of variability in these cases. This finding is in agreement with results previously obtained by means of gene action studies where estimates of dominance and various types of epistatic effect were mostly predominant. Abou-Tour (1986) found that heritability in narrow sense ranged from 34.47 to 63.58 in 1983 and from 40.36 to 56.35 in 1984 test for leaf feeding ratings.

The predicted genetic advance was rather higher for defoliation in the second cross; and moderate values for leaf area consumed in both crosses and for defoliation in the first cross. However, hairiness in both crosses gave low estimates of the predicted genetic advance (Table 2).

Johanson et al. (1955) working with soybean reported that heritability estimates along with genetic gain are usually more usefull in predicting the resultant effect of selection that heritability values alone. On the other hand Dixit et al. (1970) pointed out that high heritability is not always associated with high genetic advance, but to make selection effective, high heritability should be associated with high genetic gain.

High to moderate genetic advance were found to associate with high to moderate heritability estimates in narrow sense in all traits. Therefore, selection for these traits should be effective and satisfactory for successful breeding purposes. It is noteworthy that defoliation in the second cross showed the highest predicted genetic advance, Abou-Tour (1986) working on cotton, reported that genetic advance in two crosses was 13.33% to 22.30% in 1983 and 14.90%-22.51 in 1984.

Relatively low genetic advance was associated with moderate heritability value in hairiness in first cross. It could be concluded that selection for this trait would be less effective than former five cases.

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