

# DIALLEL CROSS ANALYSIS FOR SOME QUANTITATIVE TRAITS IN YELLOW MAIZE UNDER STRESS AND NORMAL IRRIGATION TREATMENTS

## I. BIOCHEMICAL GENETIC MARKERS FOR HETEROISIS AND COMBINING ABILITY

A.A. Abdel -Sattar<sup>1</sup> and M.F. Ahmed<sup>2</sup>

1- Genetics and Cytology Department, National Research Center, Cairo, Egypt.

2- Agronomy Dept., Faculty of Agric., Ain Shams Univ., Cairo, Egypt.

### ABSTRACT

*5X5 half diallel cross along with their parental maize inbred lines were evaluated under normal and drought environments. The obtained data revealed highly significant differences between environments for all studied traits. Partitioning of genetic variations of crosses into parents and their hybrids showed that variance due to general combining ability (GCA) was significant for all studied traits under both environments and for their combined analysis except for ear length under normal environments and the combined analysis and 100 kernels weight under drought. Variance due to specific combining ability (SCA) was significant for all studied traits under both environments and their combined analysis except for number of rows/ear under both environments and their combined data and 100-kernels weight for the combined analysis. The ratio of GCA/SCA revealed that the largest part of the total genetic variability was mainly due to non-additive gene action for most of the studied traits under normal and drought environments. Mean squares due to the interaction between each of GCA and SCA with environments were highly significant for all studied traits except for ear length, ear diameter, number of kernels per row and 100-kernels weight for GCA x environment and number of rows per ear for SCA x environment indicating that, both additive and non-additive gene effects were influenced by environments. The best general combiners for grain yield per plant and two or more of its attributes were the inbred lines P1 and P4 under normal conditions and P3 under drought conditions. Six out of ten crosses under normal conditions and three crosses under drought conditions showed positive and significant SCA effects for grain yield per plant and two or more of its attributes. In this regard, the best F<sub>1</sub> cross was P1 x P4 under both normal and drought conditions. The electrophoretic patterns (SDS-PAGE) for water soluble proteins of the five maize inbred lines and their ten F<sub>1</sub> hybrids showed that, the electrophoretic bands could be a useful tool for the identification and characterization of the used five inbred lines of yellow maize. Using soluble leaves protein electrophoresis could be effective in the identification of the highly heterotic hybrids and those having high specific combining ability effects under drought conditions as biochemical genetic markers associated with hybrid vigor and specific combining ability in yellow maize hybrids.*

**Key words:** *Diallel cross, Yellow maize, Water stress, Heterosis, Combining ability, Susceptibility index, Electrophoretic patterns.*

## INTRODUCTION

Drought, like many other environmental stresses, has adverse effects on crop yield. Low water availability is one of the major causes for crop yield reductions affecting the majority of the farmed regions around the world. As water resources for agronomic uses become more limiting, the development of drought-tolerant lines becomes increasingly more important (Bruce *et al* 2002). It has been observed that grain yield is significantly reduced when plants are grown under water shortage (Hall *et al* 1981, Quattar *et al* 1987, Sobrado 1990, Attia *et al* 1994, El-Ganayni *et al* 2000 and Yadav *et al* 2003). Maize breeders were deeply involved during the last years, in attempts to improve high yielding cultivars under drought stress environments (Edmeades *et al* 1992, Saneoka *et al* 1996, Abd El-Saboer 1997 and Younis and Aref 2001).

Information on the relative importance of general combining ability (GCA) and specific combining ability (SCA) are essential for breeding maize to tolerate drought stress conditions. Generally, GCA is associated with additive genes while SCA is attributed primarily to non-additive (dominance and epistasis) genes. It is very essential that, the breeder should evaluate the potentialities of the available germplasm for new recombinations and eventually combining ability which have proved to be of considerable use in breeding crop plants. In this regard, several studies have been reported in maize (Desai and Singh 2001, Abd El-Aty and Katta 2002, Shafey *et al* 2003 and El-Morshidy *et al* 2003). Several investigators tried to identify and characterise maize inbreds via proteins electrophoresis (Ohms 1985 and Abd El-Tawab *et al* 1989, 2001).

Therefore, this investigation aims at studying (1) the magnitude of heterosis and combining ability under normal and water stress conditions in ten maize hybrids and their parents, and (2) the efficiency of protein electrophoresis in the identification and characterization of the five maize inbred lines and their hybrids for heterosis and combining ability under drought conditions using SDS-PAGE for soluble protein patterns.

## MATERIALS AND METHODS

The genetic material used in this investigation included new five yellow maize (*Zea mays* L.) lines (AY1, AY2, AY3, AY4 and AY5), representing a wide range of diversity for several agronomic characters and drought resistance. These lines were developed by the first author through the breeding program at Dept. of Genetic and Cytology, National Research Center. These lines were derived from single cross (S.C.)150 ,S.C.151,

S.C.158, three way cross (T.W.C.)352 and double cross (D.C.) Dahab x Sabeiny.

In 2003 season, all possible cross combinations excluding reciprocals were made among the five inbred lines giving a total of 10 crosses. In 2004 season, the five parents and their ten crosses were evaluated under two moisture regimes at the Experimental Farm of the National Research Center, at Shalakan, Kalubia Governorate, Egypt. The first moisture regime was the normal (full) irrigation as it is carried out for this crop in the area (experiment I) and the second was made by withholding irrigation after 2<sup>nd</sup> irrigation for three successive irrigations (experiment II).

Each experiment was conducted in a randomized complete block design with three replications. The experimental plot included three rows of five meters long and 70 cm wide. Planting was in hills spaced at 25 cm apart. The common agricultural practices of growing maize were applied properly as recommended in the district. At harvest plant height, ear height, ear length, ear diameter, number of rows / ear, number of kernels / row, 100-grain weight and grain yield per plant were recorded on a sample of 10 guarded plants in the middle row of each plot. The drought susceptibility index (s) was used to characterize the relative stress resistance of all genotypes. The susceptibility indices were calculated independently using original data for yield and some yield components using a generalized formula (*Fischer and Maurer 1978*) as follows:

$$S = (1 - Y_d / Y_p) / D, \text{ Where}$$

S = An index of drought susceptibility

Y<sub>d</sub> = Performance of a genotype under drought stress.

Y<sub>p</sub> = Performance of the same genotype under normal irrigation.

D = Drought intensity = 1 - (mean Y<sub>d</sub> of all genotypes / mean Y<sub>p</sub> of all genotypes).

The data obtained for drought susceptibility index values for each trait were analyzed on an individual plant basis. An ordinary analysis of variance for each water regime, and the combined analysis over both regimes were performed according to Snedecor and Cochran (1981). General and specific combining ability variances and effects were obtained by employing Griffing's (1956) diallel cross analysis method 2 model I.

In the protein electrophoretical study, two leaves of each of the five parents and their 10 F's were used for SDS-protein analysis. Sodium dodecylsulphate polyacrylamide gel electrophoresis (SDS-PAGE) was

performed on water soluble protein fractions (albumin and globulin) according to the method of Laemmli (1970) as modified by Studier (1973). The SDS-protein gel was scanned and analyzed using Gel Doc 2000 BioRad System.

## RESULTS AND DISCUSSION

### Analysis of variance

Mean square estimates for all studied traits under normal, water stress and for combined data across them are shown in Table (1). Highly significant differences between environments were found for all studied traits. Mean squares due to genotypes, parents and crosses were also significant for all recorded traits under normal and water stress environments and their for combined data except ear length, ear diameter, number of rows per ear and 100 kernels weight for parents and number of rows per ear and number of kernels per row for crosses under normal environments. Parents versus crosses mean squares were significant for all traits under all environments except ear length, ear height and number of rows per ear under drought environments which were insignificant.

The mean squares due to interaction of genotypes with the two environments were highly significant for all studied traits except ear diameter and number of rows per ear which were insignificant, indicating different responses of these genotypes from normal to stress conditions. The partitioning of crosses mean squares into those due to parents and those due to F's showed that variance due to general combining ability (GCA) was significant for all studied traits under both environments and for their combined data except for ear length under normal environments and for the combined and 100 kernels weight under drought environments. Variance due to specific combining ability (SCA) was also significant for all studied traits under both environments and for their combined data except number of rows/ear under both environments and for their combined data and 100-kernels weight in the combined analysis. The ratio of GCA / SCA was less than unity for all studied traits under both environments and for their combined except ear height and number of rows per ear for both environments and their combined data, ear length and number of kernels per row under drought environments and plant height and number of kernels per row in combined analysis which were more than unity. These findings indicates that the largest part of the total genetic variability is due to non-additive gene action for the majority of the studied traits. These results are in agreement with those reported by El-Hosary *et al* 1990, Mostafa *et al* 1996, El-Morshidy *et al* 2003 and Shafey *et al* 2003.

**Table 1. Mean squares estimates for all studied traits in 5 x 5 yellow maize diallel crosses under normal and drought environments (Env.) as well as for combined data.**

S.O.V.	Env.	Df	Plant height cm	Ear height cm	Ear length cm	Ear diameter cm	No. of Rows / ear	No. of Kernels / row	100 kernels weight g	Grain yield/ plant g
Env.	Combined	1	100935.51**	27144.10**	203.10**	7.57**	51.38**	751.11**	1095.51**	111021.34**
Reps./ Env.	Normal	2	72.96	67.20	0.58	0.02	0.96	5.36	11.09	41.36
	Drought	2	68.62	110.07	1.08	0.03	0.16	12.69	2.02	18.02
	Combined	4	65.51	88.63	0.83	0.02	0.56	9.02	6.56	22.68
Genotypes (G.)	Normal	14	3099.84**	998.94**	2.88**	0.25**	3.59**	50.33**	55.02**	2019.60**
	Drought	14	2717.64**	420.63**	3.59**	0.63**	3.40**	45.15**	28.63**	616.14**
	Combined	14	2241.26**	863.50**	4.01**	0.73**	5.12**	73.49**	72.09**	2149.76**
Parents (P)	Normal	4	2655.83**	1154.17**	1.69	0.21	3.77	50.43**	12.10	1147.90**
	Drought	4	4579.17**	829.17**	6.48**	0.54**	6.43**	48.90**	16.90**	336.50**
	Combined	4	4607.08**	1881.25**	5.53**	0.50**	7.00**	85.03**	25.47**	757.87**
Crosses (C.)	Normal	9	1204.54**	278.24**	2.91**	0.25**	2.95	28.46	30.83**	955.78**
	Drought	9	1080.40**	272.93**	2.53**	0.50**	2.08**	34.89**	16.48**	361.47**
	Combined	9	1310.45**	218.30**	2.90**	0.66**	3.60**	36.70**	34.01**	1079.91**
P. vs. C.	Normal	1	21933.62**	6864.40**	7.40**	0.48**	8.71**	246.68**	444.44**	15080.28**
	Drought	1	10006.68**	115.60	1.63	2.21**	3.21	122.5**	184.9**	4026.71**
	Combined	1	1155.20**	2599.20**	7.99**	2.37**	11.25**	358.42**	601.34**	17346.05**
G. x Env.	Combined	14	3576.23**	556.00**	2.46**	0.15	1.88	21.99**	11.56**	485.94**
P. x Env.	Combined	4	2627.92**	102.00	2.63	0.26	3.20	14.30	3.53	726.53**
C. x Env.	Combined	9	974.49**	332.88**	2.54**	0.09	1.42	26.66**	13.50**	237.34**
P. vs. C. x Env.	Combined	1	30785.89**	4380.81**	1.05	0.31	0.67	10.76	28.01**	1760.95**
GCA	Normal	4	2910.36**	1354.09**	0.49	0.24**	8.68**	45.13**	32.92**	1369.39**
	Drought	4	2090.13**	481.46**	3.67**	0.30**	9.06**	80.81**	8.84	327.77**
	Combined	4	3987.60**	1630.69**	1.95	0.48**	13.45**	110.53**	35.87**	904.33**
SCA	Normal	10	3175.64**	856.89**	3.84**	0.26**	1.56	52.41**	63.86**	2279.64**
	Drought	10	2968.64**	396.30**	3.56**	0.77**	1.14	31.21**	36.54**	731.48**
	Combined	10	1542.72**	556.62**	4.84**	0.84**	1.79	58.67**	86.58	2647.93**
GCA x Env.	Combined	4	1812.88**	204.86**	2.21	0.06	4.29**	14.61	5.89	792.83**
SCA x Env.	Combined	10	4601.56**	696.56**	2.56**	0.19**	0.91	24.95**	13.83**	363.18**
Error	Normal	28	50.81	57.27	1.11	0.08	1.60	12.93	4.78	86.80
	Drought	28	43.98	43.42	1.07	0.09	1.07	8.21	4.12	61.23
	Combined	56	47.40	50.35	1.09	0.09	1.33	10.57	4.44	74.01
GCA/SCA	Normal		0.92	1.58	0.13	0.92	5.56	0.86	0.52	0.60
	Drought		8.70	1.22	1.03	0.39	7.95	2.56	0.24	0.45
	Combined		2.58	2.93	0.40	0.57	7.51	1.88	0.41	0.31

\*,\*\* indicate significant at 0.05 and 0.01 probability levels, respectively.

Highly significant mean squares due to general combining ability X environment and specific combining ability X environment were detected for all studied traits except ear length, ear diameter, number of kernels per row and 100 kernels weight for GCA X environment and number of rows per ear for SCA X environment, indicating that both additive and non-additive gene effects were influenced by environments.

### Mean performance, heterosis over better parent and susceptibility index

Mean performance for all studied traits under normal and drought conditions as well as combined data are presented in Table (2). Mean values for these traits exhibited the parental diversity and the genotypes differential response from normal to drought conditions. Regarding to grain yield per plant; inbred line-1 ranked the first as the best under normal conditions

**Table 2. Mean performance for all studied traits in 5 x 5 yellow maize diallel cross under normal and drought conditions as well as combined data.**

Genotypes	Env.	Plant height cm	Ear height cm	Ear length cm	Ear diameter cm	No. of rows/ ear	No. of kernels/ row	100 kernels weight g	Grain yield/ plant g
inbred	Normal	183.33	95.00	16.33	4.67	13.00	31.00	28.00	160.00
line P1	Drought	168.33	85.00	12.83	3.50	9.67	22.67	20.67	71.67
Ay1	Combined	175.83	90.00	14.58	4.08	11.33	26.83	24.33	115.83
P1 x P2	Normal	221.67	120.00	15.50	4.23	14.33	34.67	32.00	154.33
	Drought	148.00	80.00	13.33	3.60	12.00	25.33	29.00	90.00
	Combined	180.83	100.00	14.42	3.92	13.17	30.00	30.50	122.17
P1 x P3	Normal	265.00	141.67	17.83	4.33	15.00	36.00	36.67	150.00
	Drought	126.67	80.00	14.83	4.13	13.00	32.33	31.33	88.67
	Combined	195.83	110.83	16.33	4.23	14.00	34.17	34.00	119.33
P1 x P4	Normal	206.67	118.33	18.50	4.93	13.33	39.33	41.33	211.67
	Drought	128.33	82.00	12.50	4.57	11.67	23.33	31.33	108.33
	Combined	167.50	100.17	15.50	4.75	12.50	31.33	36.33	160.00
P1 x P5	Normal	223.33	121.00	17.00	4.73	13.67	27.67	40.00	155.00
	Drought	136.67	86.00	13.33	4.67	11.33	25.33	28.33	66.00
	Combined	180.00	103.50	15.17	4.70	12.50	26.50	34.17	118.50
inbred	Normal	158.33	85.00	17.50	4.33	12.33	34.33	26.33	121.00
line P2	Drought	210.00	80.00	15.00	4.00	13.00	30.00	23.00	80.00
Ay2	Combined	184.17	82.58	16.25	4.17	12.67	32.17	24.67	100.50
P2 x P3	Normal	250.00	141.67	17.50	4.53	14.67	32.33	34.67	170.00
	Drought	143.33	77.33	13.17	3.87	13.67	30.00	26.33	93.33
	Combined	196.67	109.50	15.33	4.20	14.17	31.17	30.50	131.67
P2 x P4	Normal	271.67	139.33	17.33	4.43	13.67	36.00	32.00	161.00
	Drought	153.33	80.67	14.83	3.43	11.33	32.67	27.67	95.67
	Combined	212.50	110.00	16.08	3.73	12.50	34.33	29.83	128.33
P2 x P5	Normal	243.33	131.00	17.00	4.20	12.00	33.00	35.33	164.33
	Drought	163.33	91.67	14.00	3.53	12.67	26.00	24.33	93.33
	Combined	203.33	111.33	15.50	3.87	12.33	29.50	29.83	128.83
inbred	Normal	240.00	133.33	15.83	4.13	14.67	30.67	28.00	123.33
line P3	Drought	230.00	118.33	14.00	3.83	13.00	25.67	21.67	85.00
Ay3	Combined	235.00	125.83	14.92	3.98	13.83	28.17	24.83	104.17
P3 x P4	Normal	251.67	142.67	15.33	4.77	14.67	36.00	34.33	172.67
	Drought	148.33	88.00	13.67	3.93	12.00	31.00	25.33	94.67
	Combined	200.00	115.33	14.50	4.35	13.33	33.50	29.83	133.67
P3 x P5	Normal	245.00	129.67	17.60	4.57	13.67	33.33	32.33	165.00
	Drought	152.00	90.33	14.20	3.93	12.33	28.33	26.67	89.00
	Combined	198.50	110.00	15.90	4.25	13.00	30.83	29.50	127.00
inbred	Normal	200.00	116.67	15.67	4.10	12.00	24.67	31.33	136.00
line P4	Drought	156.67	90.00	14.50	3.00	10.67	24.67	26.00	62.67
Ay4	Combined	178.33	103.33	15.08	3.85	11.33	24.67	28.67	99.33
P4 x P5	Normal	236.67	126.67	17.33	4.33	12.33	33.33	36.00	181.67
	Drought	191.67	110.00	15.50	4.03	11.00	24.67	26.67	101.67
	Combined	214.17	118.33	16.42	4.18	11.67	29.00	31.33	141.67
inbred	Normal	191.67	95.00	15.83	4.00	12.00	25.33	30.33	108.33
line P5	Drought	135.00	76.67	11.33	3.17	11.33	19.00	25.67	60.67
Ay5	Combined	163.33	85.83	13.58	3.58	11.67	22.17	28.00	84.50
LSD	Normal	11.92	12.66	1.76	0.48	2.11	6.01	3.66	15.58
0.05	Drought	11.09	11.02	1.73	0.52	1.73	4.79	3.39	13.09
	Combined	7.96	8.21	1.21	0.35	1.33	3.76	2.44	9.95
LSD	Normal	17.34	18.42	2.56	0.70	3.07	8.74	5.33	22.67
0.01	Drought	16.14	16.03	2.52	0.76	2.52	6.97	4.93	19.04
	Combined	10.93	11.27	1.66	0.48	1.83	5.16	3.35	13.66

while inbred line-3 was the best under drought conditions. The hybrid P1 x P4 was the best in grain yield under normal and drought environments. The hybrids P2 x P3, P2 x P5, P3 x P5, P4 x P5 and P1 x P4 under normal conditions and the hybrids P1 x P4 and P4 x P5 under drought conditions, respectively exceeded their better parents for yielding ability (Table 3). It is also clear from Table (3) that the best hybrids under normal conditions were P2 x P4 for plant height, P1 x P4 for ear length, number of kernels per row

**Table 3. Percentage of heterosis over better parent for all studied traits under normal and Drought conditions.**

Genotype	Env.	Plant height cm	Ear Height cm	Ear Length cm	Ear Diameter cm	No. of rows / ear	No. of kernels/ row	100 kernels weight g	Grain yield/ plant g
P1 x P2	Normal	20.91**	26.32**	-11.43*	-9.29	10.26	0.97	14.29*	-3.54
	Drought	-32.80**	-5.88	-14.89	-13.60	-5.26	-8.43*	29.85**	12.50
P1 x P3	Normal	10.42**	6.25	9.18	-7.14	2.27	16.13	30.95**	-6.25
	Drought	-44.93**	-32.39**	5.95	7.83	0.00	25.97**	44.62**	4.31
P1 x P4	Normal	3.33	1.43	13.27*	5.71	2.56	26.88**	31.92**	32.29**
	Drought	-23.76**	-8.89	-13.79*	30.48**	9.38	-5.41	20.51**	51.16**
P1 x P5	Normal	16.52**	27.37**	4.08	1.43**	5.13	-10.75	31.87**	-3.13
	Drought	-18.81**	1.18	3.90	33.33**	0.00	11.77	10.39	-7.91
P2 x P3	Normal	4.17	6.25	0.00	4.62	0.00	-5.83	23.81**	37.84**
	Drought	-37.68**	-34.65**	-15.96*	-7.20	5.13	8.43	17.91	9.80
P2 x P4	Normal	35.83**	19.43**	-0.95	-6.92	10.81	4.85	2.13	18.38**
	Drought	-26.40**	-10.37	-5.32	-17.60*	-10.53*	18.07	6.41	19.58**
P2 x P5	Normal	26.96**	37.90**	-2.86	-3.08	-2.70	-3.88	16.48**	35.81**
	Drought	-21.60**	14.58*	-10.64	-15.20	0.00	-6.02	-5.20	16.66*
P3 x P4	Normal	4.86*	7.00	-3.16	15.32*	0.00	17.39	9.57	26.96**
	Drought	-35.51**	-25.63**	-5.75	2.61	-7.70	20.78*	-2.56	11.37
P3 x P5	Normal	2.98	-2.75	11.16*	10.48	-6.82	8.70	6.60	33.78**
	Drought	-33.91**	-23.66**	1.43	2.61	-5.13	10.39	3.90	4.71
P4 x P5	Normal	18.33**	8.57	9.47*	5.69	2.78	31.58**	14.90**	33.58**
	Drought	22.34**	22.22**	6.90	27.37**	-2.94	0.00	2.56	62.23**

\*,\*\* indicate significant at 0.05 and 0.01 probability levels, respectively.

and 100 kernels weight, P3 x P4 for ear diameter and ear height and P1 x P3 for number of rows per ear. The best hybrids under drought conditions were P4 x P5 for plant height, ear length and ear height, P1 x P5 for ear diameter, P2 x P3 for number of rows per ear, P2 x P4 for number of kernels per row and P1 x P3 for 100 kernels weight.

Stress susceptibility index "S" was used to estimate the relative stress injury because it accounted for variation in yield potential and stress intensity. Low stress susceptibility ( $S < 1$ ) is synonymous with higher stress tolerance (Fischer and Maurer 1978). The results indicated that S values (Table 4) varied between parents and their F1 crosses for all studied traits. P3, P2, P2 x P4, P1 x P3 and P1 x P2 were the best drought tolerant genotypes for grain yield per plant and one or more of its attributes.

**Table 4. Stress susceptibility index (S) of the fifteen genotypes for grain yield and its components.**

Genotypes	No. of rows / ear	No. of kernels / row	100 kernels weight g	Grain yield/ Plant g
P1	2.28	1.51	1.25	1.22
P1 x P2	1.45	1.52	0.45	0.92
P1 x P3	1.19	0.57	0.70	0.91
P1 x P4	1.11	2.29	1.15	1.08
P1 x P5	1.52	0.48	1.39	1.27
P2	-0.48	0.71	0.60	0.75
P2 x P3	0.61	0.41	1.15	1.00
P2 x P4	1.52	0.52	0.65	0.90
P2 x P5	-0.50	1.19	1.49	0.96
P3	1.01	0.92	1.08	0.69
P3 x P4	1.62	0.78	1.25	1.00
P3 x P5	0.87	0.84	0.84	1.02
P4	0.99	0.00	0.81	1.20
P4 x P5	0.96	1.46	1.24	0.98
P5	0.50	1.41	0.73	0.98

## Combining ability estimates

Estimates of the general combining ability effects ( $g_i$ ) of each parental genotype for all studied traits are presented in Table (5). Inbred lines P1, P2 and P5 exhibited negative and highly significant ( $g_i$ ) effects for plant and ear height, respectively under the normal environment. While P1, P4 and P5, respectively for plant height and P1 and P2 for ear height showed negative and highly significant ( $g_i$ ) effects under drought conditions, indicating that, those inbred lines possess favorable genes for developing improved hybrids with short plants. However, the inbred lines P1 and P4 under normal conditions and P3 under drought conditions exhibited positive and highly significant ( $g_i$ ) effects for grain yield per plant. The inbred line P1 showed positive and highly significant ( $g_i$ ) effects for ear diameter under normal and drought conditions. The inbred line P3 under normal conditions and P2 and P3 under drought conditions exhibited positive and highly significant ( $g_i$ ) effects for number of rows per ear. The inbred lines P2 and P3 under drought conditions and P2 under normal conditions showed positive and highly significant ( $g_i$ ) effects for number of kernels per row. The inbred lines P1 and P4 under normal conditions and P4 under drought conditions exhibited positive and highly significant ( $g_i$ ) effects for 100 kernels weight. This is of practical interest in breeding programs for developing high grain yield per plant and one or more of the remaining traits under normal and drought conditions.

Table 5. Estimates of general combining ability effects for the parental genotypes evaluated under normal and stress environments (env.) as well as for their combined (comb.) data..

Parental Genotype	Env.	Plant height cm	Ear height cm	Ear length cm	Ear diameter cm	No. of rows / ear	No. of kernels/ row	100 kernels weight g	Grain yield/ plant g
P1	Normal	-10.29**	-6.26**	0.09	0.17**	0.26	0.66	0.93*	8.18**
	Drought	-12.16**	-4.06**	-0.45*	0.16**	-0.59**	-1.25*	0.53	-2.28
	Combined	-11.22	-5.16	-0.18	0.16	-0.17	-0.30	0.73	2.95
P2	Normal	-7.43**	-4.69**	0.21	-0.10	-0.17	1.37*	-1.83**	-6.01**
	Drought	9.51**	-5.25**	0.36	-0.06	0.60**	1.94**	-0.61	2.87
	Combined	1.04	-4.97	0.29	-0.08	0.21	1.66	-1.22	-1.57
P3	Normal	19.48**	12.51**	-0.13	0.02	0.97	0.56	-0.78	-4.2*
	Drought	10.98**	6.56**	0.15	0.09	0.79**	1.80**	-0.66	3.34*
	Combined	15.23	9.53	0.01	0.05	0.88	1.18	-0.72	-0.43
P4	Normal	1.62	3.65**	-0.14	-0.01	-0.36	-0.15	0.98*	9.32**
	Drought	-2.64*	2.04	0.38	-0.13*	-0.59**	0.09	0.77*	1.91
	Combined	-0.51	2.84	0.12	-0.07	-0.48	-0.03	0.88	5.62
P5	Normal	-3.38*	-5.21**	-0.03	-0.08	-0.70**	-2.44**	0.70	-7.30**
	Drought	-5.69**	0.71	-0.45*	-0.05	-0.21	-2.58**	-0.04	-5.85**
	Combined	-4.53	-2.25	-0.24	-0.07	-0.45	-2.51	0.33	-6.57
LSD	( $g_i-g_j$ )								
0.05	Normal	4.51	4.79	0.67	0.18	0.80	2.28	1.38	5.90
	Drought	4.20	4.17	0.66	0.20	0.65	1.81	1.28	4.95
	Combined	3.01	3.10	0.46	0.13	0.50	1.42	0.92	3.76
0.01	Normal	6.07	6.45	0.90	0.25	1.08	3.06	1.86	7.95
	Drought	5.65	5.61	0.88	0.27	0.88	2.44	1.73	6.67
	Combined	4.00	4.12	0.63	0.17	0.67	1.89	1.22	5.00

\*, \*\* indicate significant at 0.05 and 0.01 probability levels, respectively.



Specific combining ability effects of the ten crosses for all studied traits under normal and drought conditions are presented in Table (6). Seven out of ten crosses under drought conditions and one hybrid (P1 x P4) under normal conditions for plant height and six crosses under drought conditions and two hybrids under normal conditions for ear height exhibited negative and significant specific combining ability effects, indicating that these crosses possess favorable genes for producing improved hybrids with short plant and low ear placement. For grain yield per plant, six out of ten crosses under normal conditions and three crosses under drought conditions showed positive and significant SCA effects. Regarding to ear length and ear diameter two out of ten crosses under normal and drought conditions, respectively exhibited positive and significant SCA effects. For number of kernels per row two out of ten hybrids under drought and one hybrid (P1 x P4) under normal conditions exhibited positive and significant SCA effects. For 100- kernels weight five out of ten crosses under normal and three crosses under drought conditions exhibited positive and significant SCA effects.

When a cross ranks highest for SCA effect such as the cross (1 x 4) for grain yield per plant and three or more of its attributes and at the same time ranks the best for its mean performance in the same trait, such cross would be considered as a good breeding material to improve this specific trait. Similar results were reported by Desai and Singh (2001), Abd El-Aty and Katta, (2002), El-Morshidy *et al* (2003) and El-Shouny *et al* (2003).

### **SDS-Protein electrophoresis**

Electrophoretic assays have been widely used as a rapid and accurate test to identify and characterize different crop cultivars. By the use of appropriate and refined techniques, it is now possible to actually fingerprint each cultivar to assess its identity and its agronomic properties. SDS-PAGE analysis of water soluble proteins was successfully used for the identification and prediction of heterosis and combining ability in maize and different field crops (Abd El-Tawab *et al* 1989 and Esmail *et al* 1999).

The electrophoretic patterns (SDS-PAGE) for water soluble proteins (albumin and globulin) of the five maize inbred lines and their ten hybrids are illustrated in Figure (1) and Table (7). From the SDS-PAGE analysis, 43 bands were detected with different molecular weights (MW) and relative mobilities (Rm). Numbers inside the table represent the intensity percentages (%) of each band out of the total column.

**Table 6. Estimates of specific combining ability effects for ten yellow maize crosses evaluated under normal and drought environments as well as for their combined data.**

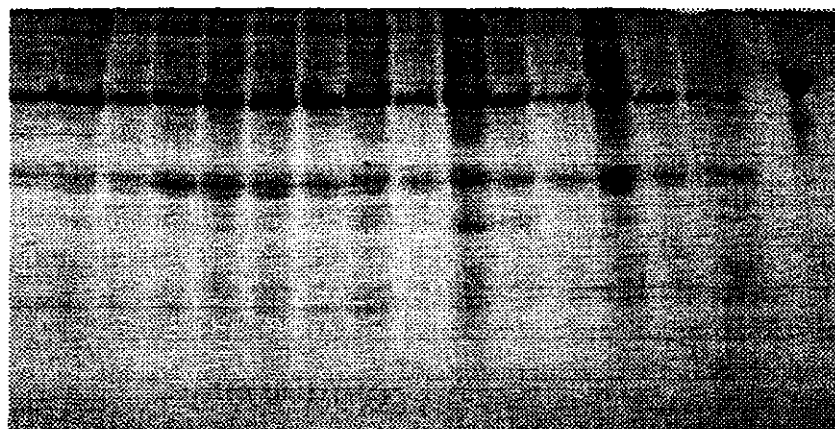
Genotypes	Env.	Plant height cm	Ear height cm	Ear length cm	Ear diameter cm	No. of rows/ ear	No. of kernels/ row	100 kernel weight g	Grain yield/ plant g
P1 x P2	Normal	13.49**	8.48*	-1.61**	-0.23	0.83	0.13	-0.35	-3.46
	Drought	-16.25**	1.57	-0.38	-0.31*	0.08	-2.10	2.81**	4.03
	Combined	-1.38	5.02*	-1.00**	-0.27**	0.45	-0.98	1.23	0.29
P1 x P3	Normal	29.92**	12.95**	1.06*	-0.25	0.35	2.27	3.27**	-9.60*
	Drought	-31.00**	-10.24**	1.33**	0.07	0.88	5.05**	5.19**	2.22
	Combined	-0.57	1.36	1.20**	-0.09	0.62	3.66**	4.23**	-3.70
P1 x P4	Normal	-10.56**	-1.52	1.74**	0.38**	0.02	6.32**	6.18**	38.55**
	Drought	-15.78**	-3.71	-1.24*	0.73**	0.94	-2.24	3.76**	23.32**
	Combined	-13.17**	-2.62	0.25	0.56**	0.48	2.04	4.97**	30.93**
P1 x P5	Normal	11.11**	10.00**	0.13	0.24	0.68	-3.06	5.13**	-1.51
	Drought	-4.40	1.62	0.43	0.75**	0.22	2.43	1.57	-11.25**
	Combined	3.36	5.81*	0.28	0.50**	0.45	-0.32	3.35**	-6.38*
P2 x P3	Normal	12.06**	11.38**	0.61	0.22	0.44	-2.11	4.03**	24.59**
	Drought	-36.06**	-11.71**	-1.15*	0.02	0.37	-0.48	1.33	1.75
	Combined	-12.00**	-0.17	-0.27	0.12	0.41	-1.30	2.68**	13.17**
P2 x P4	Normal	51.59**	17.91**	0.46	-0.25	0.78	2.27	-0.40	2.06
	Drought	-12.44**	-3.86	0.29	-0.19	-0.59	3.91**	1.24	5.51
	Combined	19.57**	7.02**	0.37	-0.22*	0.10	3.09**	0.42	3.79
P2 x P5	Normal	28.25**	18.43**	0.01	-0.02	-0.56	1.56	3.22**	22.02**
	Drought	0.60	8.48**	0.28	-0.16	0.37	-0.10	-1.29	10.94**
	Combined	14.43**	13.45**	0.15	-0.09	-0.10	0.73	0.97	16.48**
P3 x P4	Normal	4.68	4.05	-1.20*	0.37**	0.64	3.08	0.89	11.92**
	Drought	-18.92**	-8.33**	0.67	0.16	-0.11	2.38	-1.05	4.03
	Combined	-7.12**	-2.14	-0.94**	0.26**	0.26	2.73**	-0.08	7.98**
P3 x P5	Normal	3.02	-0.10	0.96	0.23	-0.03	2.70	-0.83	20.87**
	Drought	-12.21**	-4.67	0.69	0.08	-0.16	2.38	1.10	6.13
	Combined	-4.60	-2.38	0.83*	0.16	-0.10	2.54*	0.14	13.50**
P4 x P5	Normal	12.54**	5.76	0.71	0.03	-0.03	3.41	1.08	24.02**
	Drought	41.08**	19.52**	1.76**	0.41	-0.11	0.43	-0.33	20.22**
	Combined	26.81**	12.64**	1.23**	0.22*	-0.07	1.92	0.37	22.12**
<b>LSD (Sij - Sjk)</b>									
0.05	Normal	11.05	11.73	1.63	0.45	1.96	5.57	3.39	14.44
	Drought	10.27	10.21	1.61	0.48	1.60	4.44	3.14	12.13
	Combined	7.36	7.59	1.12	0.32	1.23	3.48	2.25	9.20
0.01	Normal	14.87	15.79	2.19	0.60	2.64	7.50	4.56	19.44
	Drought	13.83	13.75	2.16	0.65	2.15	5.98	4.23	16.33
	Combined	9.79	10.09	1.48	0.43	1.64	4.62	3.00	12.23
<b>LSD (Sij - Sjk)</b>									
0.05	Normal	10.08	10.70	1.49	0.41	1.79	5.08	3.09	13.18
	Drought	9.39	9.33	1.47	0.44	1.46	4.05	2.87	11.07
	Combined	6.72	6.93	1.02	0.29	1.13	3.17	2.06	8.40
0.01	Normal	13.58	14.41	2.00	0.55	2.41	6.85	4.17	17.74
	Drought	12.64	12.56	1.97	0.59	1.97	5.46	3.86	14.90
	Combined	8.94	9.21	1.36	0.39	1.50	4.22	2.74	11.17

\*,\*\* indicate significant at 0.05 and 0.01 probability levels, respectively.

**Table 7. Densitometer analysis of water soluble proteins (SDS-PAGE) showing number of bands (B.no.), Relative mobility (Rm), molecular weight (Mw) and intensity as a percentage of total concentration for 5 x 5 maize diallel crosses.**

B. No.	R.m	M.W. K.Da	Inbred Parents					Hybrids														
			1	2	3	4	5	1x2	1x3	1x4	1x5	2x3	2x4	2x5	3x4	3x5	4x5					
1	0.011	224.77				4.22	2.78													3.82		
2	0.022	211.02		4.65	4.61							4.01		3.89	4.29					1.88	2.25	
3	0.051	167.41			2.75	2.78	3.98						1.78	3.98	2.96				2.92			
4	0.056	162.67		2.78					2.94	2.93									2.93	1.98	1.46	
5	0.061	158.86									2.27							3.04				
6	0.070	151.99	3.05	5.53	3.90	4.31	5.15	3.48	4.00	4.43	3.65	2.78	5.27						3.23	2.83	1.50	
7	0.101	135.52	2.84	2.3	2.57	3.12	2.59	2.58	1.74	1.86	1.95	2.75	1.85						2.58			
8	0.124	122.76	3.25	3.76	4.00	3.30	2.79	2.34	2.62	3.28	2.92	1.97	2.54	2.62	3.72	2.53	2.02					
9	0.152	111.44		1.80	3.76								2.57						2.36	2.13		
10	0.158	111.42	2.90			2.08	3.79					2.55										
11	0.164	106.92		3.58				2.81	2.39	1.99	1.88	3.48	3.06					3.20	1.43			
12	0.177	100.81				1.98					1.40											
13	0.187	91.47	3.10																		2.70	
14	0.191	94.05			2.28																	
15	0.205	88.58				2.79	3.14	2.04		2.98	1.19	2.78	2.54	2.39	1.85							
16	0.243	81.96	10.7	12.5	11.7	12.2	8.86	12.8	15.8	14.3	10.3	10.1	13.4	7.99	8.65	7.30	4.04					
17	0.276	73.34		2.70	1.72	1.01		1.97			2.91								1.89	0.92		
18	0.282	72.45	1.80				1.36		1.87	1.38		1.79	1.23						1.56	1.45		
19	0.296	67.22		2.44	1.38		1.63	1.78			1.53		1.65					1.77				
20	0.306	66.02				1.87			1.31	1.17		2.07		1.65				1.23				
21	0.312	65.16	1.52			1.04	1.25	0.76		0.95												
22	0.323	62.68		1.09								1.20		0.97	1.22						1.67	
23	0.333	61.65		1.35			1.14	1.06		1.19			1.13									
24	0.348	59.67			1.25			0.74	0.81	0.92				0.51	1.13							
25	0.435	47.53	2.47								1.63	1.66	1.59								4.72	
26	0.450	46.03		2.98	3.01						3.14	2.48	4.05	1.95	3.46	4.56	3.87				1.05	
27	0.457	44.68	1.05																			
28	0.466	44.17				4.48	7.08	6.66	4.16		1.33		0.80	1.22	2.13	0.72						
29	0.476	43.24		1.66	1.12					1.07		0.91										
30	0.488	42.36				1.28	2.32	1.83	0.91			0.55								0.77		
31	0.525	40.05		0.45	0.62	0.32			0.65	1.16		0.70	0.56	1.08	1.34						1.42	
32	0.544	38.93	0.99				0.73	0.35		0.70											1.36	
33	0.563	37.84		0.63	0.53	0.49			0.75	1.00									1.21			
34	0.575	30.85					0.99	0.59			2.40								0.69		1.17	
35	0.636	34.54					0.78	0.85														
36	0.695	52.07					0.87	1.04											0.74		1.26	
37	0.702	31.69																			0.94	
38	0.720	31.13									0.65	0.79	0.49	0.80	0.76						2.10	
39	0.744	29.93					2.95	0.80		0.85		0.79										
40	0.753	29.68	1.47				0.59	1.09											1.35	1.26	1.51	2.70
41	0.774	28.27									1.81	1.61	1.74	1.09								
42	0.783	28.29		1.91	2.20	0.99																
43	0.792	27.57					1.34	1.54														

Numbers inside the table represent proper intensity percentages of each band



SDS-SP 1→15

Figure 1. SDS Electrophoretic patterns of water soluble protein in 15 genotypes of yellow maize

Two bands are commonly present in all five parents and their ten hybrids of MW 122.76 and 81.96 KDa. These bands were considered as marker bands for these genotypes. The five inbred lines were discriminated from each other by some unique bands, where inbred line P1 showed three bands of MW 91.47, 47.53 and 44.68 KDa. The inbred line P2 characterized by three bands of MW 162.67, 106.92 and 62.68 KDa. The inbred line P3 showed two bands of MW 94.05 and 59.67 KDa. Two bands of MW 100.81 and 66.02 KDa. characterized the inbred line P4. Five bands of MW 30.85, 34.54, 32.07 29.93 and 27.57 KDa. characterized the inbred line P5. From these results it is concluded that the analysis of soluble protein electrophoretic bands could be a useful tool for the identification and characterization of the five inbred lines of yellow maize. Variations among different field crops in SDS-protein banding patterns have been reported by many authors (Abd El-Tawab *et al* 1978, 1989, Afiah *et al* 1999 and Esmail *et al* 1999). Regarding the hybrids, six out of ten crosses (P1 x P2, P1 x P4, P2 x P3, P2 x P4, P3 x P4 and P4 x P5) exhibited number of bands which exceeded their respective parents (Table7) and were distinguished by having more hybrid bands. In the same time, most of these hybrids showed substantial hybrid vigor with regard to grain yield per plant under drought conditions (Table3) and positive significant specific combining ability effects under drought conditions (Table 6). Four hybrids (P1 x P3, P1 x P5, P2 x P5 and P3 x P5) exhibited a number of bands which did not exceed the number of bands of their parents. These crosses showed insignificant and negative heterosis and specific combining ability under drought conditions (Tables 3 and 6), except the hybrid P2 x P5 which had positive and significant heterosis and specific combining ability.

These results indicated to some extent the effectiveness of using soluble seed protein electrophoresis in the identification of the highly heterotic hybrids and high specific combining ability under drought conditions as biochemical genetic markers associated with hybrid vigor and specific combining ability in yellow maize hybrids.

## REFERENCES

- Abd El-Aty, M.S. and Y.S. Katta (2002). Estimation of heterosis and combining ability for yield and other agronomic traits in Maize hybrids (*Zea mays* L.). J. Agric. Sci. Mansoura Univ. 27(8): 5137-5146.
- Abd El-Saboor, G.A. (1997). Genetic structure of *Zea mays* L. population adopted to dry conditions. M.Sc. Thesis, Fac. of Agric., Assuit University.
- Abdel-Tawab, F.M., A.M. Omar and M.A. Rashed (1978). Heterosis and combining ability in maize (*Zea mays*, L.). II. Chemical constituents and serological analysis. Iraqi J. Agric. Sci. 12:97-108.

- Abdel-Tawab, F.M., Eman M. Fahmy, M.A. Rashed and M.H. Abou Deif (1989). Protein and isozyme polymorphism as related to heterosis and combining ability in Maize. *Egypt. J. Genet. Cytol.*18: 203-217.
- Abdel-Tawab, F.M., M.A. Rashed, F.M. El-Domyati, T.Z. Salam, S.A. Azer and A.F. Khafaga (2001). Marker-assisted selection for salt tolerance in maize (*Zea mays*, L.). *Egypt. J. Genet. Cytol.*30: 175-188.
- Afiyah, S.A.N., H.Z. Hassan, S.A.M. Khattab, S.A. Ibrahim and A.Z.E. Abdel Salam (1999). Genetic analysis of bread wheat diallel crosses under saline and normal conditions. I. Biochemical genetic marker for heterosis and combining ability. *Desert Inst. Bull., Egypt.* 49(1): 189-218.
- Attia, M.M., H.A. Agrama and H.E. Khalifa (1994). Effect of irrigation intervals on yield of some corn varieties in calcareous soil of West Nubaria region. *J. Agric. Sci. Mansoura Univ.* 19: 3155-3162.
- Bruce, W.B., G.O. Edmeades and T.C. Barker (2002). Molecular and physiological approaches to maize improvement for drought tolerance. *J. Exper. Botany* 53(3):13-25.
- Desai, S.A. and R.D. Singh (2001). Combining ability studies for some morphophysiological and biochemical traits related to drought tolerance in maize (*Zea mays* L.). *Indian J.Gent.*61(1):34-36.
- Edmeades, G.O., J. Bolanos and H.R. Lafitte (1992). Progress in breeding for drought tolerance in maize. 47<sup>th</sup> Annual Corn and Sorghum Res. Conference; 93-111.
- El-Ganayni, A.A., A.M. Al-Naggar, H.Y. El-Sherbieny and M.Y. El-Sayed (2000). Genotypic differences among 18 maize populations in drought tolerance at different growth stages. *J. Agric. Sci. Mansoura Univ.* 25(2): 713-727.
- El-Hosary, A.A., G.A. Sary and A.A. Abdel -Sattar(1990). Studies on combining ability and heterosis in maize (*Zea mays* L.). *Growth attributes. Egypt. J. Agron.*15(1-2): 23-34.
- El-Morshidy, M.A., E.A. Hassaballa, S.F. El-Saad and M.A. Abd El-Moula (2003). Combining ability and type of gene action in maize under favorable and water stress environments. *Egypt. J. Plant. Breed.* 7(1): 55-75.
- El-Shouny, K.A., Olfat H. El-Bagoury, H.Y. El-Sherbieny and S.A. Al-Ahmed (2003). Combining ability estimates for yield and its components in yellow maize (*Zea mays*, L.) under two plant densities. *Proc. Third Pl. Breed. Con.* April 26 (Giza) *Egypt J. Plant Breed.* 7(1): 399-417.
- Esmail, A.M., A.M. El-Marakby, M.A. Rashed and M.F. Ahmed (1999). Prediction of heterosis and combining ability in some cotton crosses via serological and electrophoretic analysis. *Ann. Agric. Sci. Ain Shams Univ. Cairo* 44(2): 523-536.

- Fischer, R.A. and R. Maurer (1978).** Drought resistance in spring wheat cultivars I-Grain yield responses. *Aust. J. Agric. Res.* 29: 897-912.
- Griffing, B. (1956).** Concept of general and specific combining ability in relation to diallel crossing systems. *Aust. J. Biol. Sci.* 9: 463-493.
- Hall, A.J., H.D. Ginzo and J.H. Lemcoff (1981).** Water stress before and during flowering in maize and its effects on yield, its components and their determinants. *Maydica* 26: 19-38.
- Laemmli, U.K. (1970).** Cleavage of structural proteins during the assembly of the head bacteriophage T4. *Nature* 227: 680-685.
- Mostafa, M.A.; A.A. Abd El-Aziz and H.Y. El-Sherbiney (1996).** Diallel analysis of grain yield and natural resistance to late wilt disease in newly developed inbred lines of maize. *Bull. Fac. Agric. Cairo Univ.* 47: 393-404.
- Ohms, J.P. (1985).** Determination of the maternal type in hybrid maize seed by isoelectric focusing of corn proteins. 2<sup>nd</sup> International Symposium "Cultivar identification and evaluation" 5-9 May 1985 Braunschweig, Germany.
- Quattar, S., R.J. Jones and R.K. Crookston (1987).** Effect of water deficits during grain filling on the pattern of maize kernel growth and development. *Crop Sci.* 27: 726-730.
- Saneoka, H., S. Ogata and W. Agata (1996).** Cultivar differences in dry matter production and leaf water relations in water stress maize (*Zea mays*, L.). *Grassland Sci.*, 41(4): 294-301.
- Shafey, S.A.; H.E. Yassien; I.E. M.A. El-Beially and O.A.M. Gad-Alla (2003).** Estimates of combining ability and heterosis effects for growth, earliness and yield in maize (*Zea mays* L.). *J. Agric. Sci. Mansoura Univ.* 28(1): 55-67.
- Snedecor, G.W. and W.G. Cochran (1981).** Statistical methods applied to experiments in agriculture and biology .7<sup>th</sup> ed. Iowa State College, Press, Ames, Iowa, U.S.A.
- Sobrado, M.A. (1990).** Drought responses of tropical corn. I. Leaf area and yield components in the field. *Maydica* 35: 221-226.
- Studier, F.W. (1973).** Analysis of bacteriophage T7 early RNAs and proteins of slab gels. *J. Mol. Biol.* 79: 237-248.
- Yadav, T.P., R.D. Singh and J.S. Bhat (2003).** Genetics studies under different levels of moisture stress in maize (*Zea mays* L.). *Indian J. Genet.* 63(2): 119-123.
- Younis, F.G. and Kh.A. Aref (2001).** Effect of favourable and drought stress conditions on variance components for yield and it's attributes of maize hybrids. *J. Agric. Sci. Mansoura Univ.* 26(2): 667-679.

# تحليل الهجن التبادلية لبعض الصفات الكمية فى الذرة الصفراء تحت الظروف

## المثلث وظروف الإجهاد المائى

### ١ . الالته الوراثية البيوكيماوية لقوة الهجين والقدرة على التالف

عبد الستار أحمد عبد الستار<sup>١</sup> و مصطفى فزاع أحمد<sup>٢</sup>

١- قسم الوراثة والسيولوجى\_المركز القومى للبحوث\_القاهرة\_مصر .

٢ - قسم المحاصيل\_كلية الزراعة\_جامعة عين شمس-القاهرة-مصر .

يهدف البحث إلى دراسة قوة الهجين والقدرة على الالتلاف فى الذرة الصفراء تحت الظروف المثلثى وظروف الإجهاد المائى ( منع الرى بعد الريه الثانيه ولمدة ثلاث ريات متتاليه ) بمحطة البحوث والتجارب الزراعية بالمركز القومى للبحوث\_شلفان\_قيوبية وكذلك التعرف على الألة الوراثية البيوكيماوية لتحمل الجفاف باستخدام خمسة سلالات مرباة داخليا وكلاء الهجن التبادلية دون العكسية (عشرة هجن)، حيث تم تقييمها فى موسم ٢٠٠٤ فى تصميم تجريبي قطاعات كاملة الصنولية من ثلاثة مكررات وتم أخذ القراءات على صفات ارتفاع النبات(سم)، ارتفاع الكوز (سم)، طول الكوز (سم)، قطر الكوز (سم)، عدد الحبوب بالصف، عدد الصفوف بالكوز، وزن ١٠٠ حبه (جرام) وإنتاجيه النبات من الحبوب (جرام). وتم تحليل النتائج بطريقة جريفنج ١٩٥٦ (الموديل الاول\_ الطريقة الثانية)، ويمكن تلخيص أهم النتائج فيما يلى:-

١- كان متوسط مجموع المربعات المرجح إلى كل من التركيب الورثيه والآباء والهجن على المستوى لكل الصفات المدروسة تحت كلا من الظروف المثلثى وظروف الجفاف ما عدا طول الكوز وقطر الكوز وعدد الصفوف بالكوز ووزن ١٠٠ حبه للآباء وعدد الصفوف بالكوز وعدد الحبوب بالصف للهجن تحت الظروف المثلثى .

٢- أظهر تباين اليبينات أروفاً معنوية كبيرة فى كل الصفات تحت الدراسة .

٣- أظهرت النسبة بين تباين كلا من القدرة العامة والخاصة على التالف أن الجزء الأكبر من التباين الوراثى الكلى يرجع إلى التباين الغير مضيف وذلك لمعظم الصفات المدروسة تحت الظروف المثلثى وظروف الإجهاد المائى .

٤- أوضح التفاعل بين كل من القدرة العامة والخاصة على التالف والبيئة أنه على المعنوية لمعظم الصفات تحت الدراسة ما عدا صفات طول الكوز وقطر الكوز وعدد الحبوب بالصف ووزن المائة حبة للتفاعل القدرة العامة على التالف والبيئة وصفة عدد الصفوف للكوز لتفاعل القدرة الخاصة على التالف والبيئة حيث كان غير معنوى مشيراً إلى تأثير كلا من الفعل الجينى المضيف والفعل الجينى الغير مضيف بالتغير فى ظروف البيئة .

٥- كانت أحسن التركيب الوراثيه الأبوية بالنسبة للقدرة العامة على التالف فى صفة محصول النبات من الحبوب وإثنين أو أكثر من الصفات المساهمه فيه هى سلالتان P1 و P4 تحت الظروف المثلثى والسلالة P3 تحت ظروف الجفاف .

٦- أظهرت ستة هجن تحت ظروف الرى العادى وثلاثة هجن تحت ظروف الجفاف لثرة خاصة على التالف عالية المعنوية وموجبة بالنسبة لصفة محصول النبات من الحبوب وإثنين أو أكثر من الصفات المساهمة وكان أحسن هذه الهجن هو الهجين P1xP4 تحت كلا من الظروف المثلثى وظروف الجفاف .

٧- أظهر تنفيد الكهرسى للبروتينات الذائبة أنه أداة فعالة للتعرف على الخمس سلالات المرباة دلتغيا من الـذرة الصفراء كما أظهر أيضاً كفاءة فى التعرف على قوة الهجين والقدرة الخاصة على الانتلاف تحت ظروف الجفاف فى هجن الـذرة الصفراء.