GENETIC VARIABILITY AND HERITABILITY IN SOME GENOTYPES OF MAIZE

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ABSTRACT

This investigation was carried out at Sakha and Gemmeiza Research stations during the summer season of 2003 to evaluate the behaviour of twelve maize genotypes received from Field Crops Research Institute, Agricultural Research Center, to determine the phenotypic, genotypic, environmental variations, heritability in broad sense, genotypic and phenotypic coefficients of variation. Combined data over the two locations indicated that average performance for grain yield ranged from 164.8 gm (T.W.C. 320) to 258.2 gm (S.C. 10). At the two locations and combined data, plants of Dahab were earlier than the other genotypes in days to mid silk and days tasseling ,Pioneer 3062 gave the highest number of rows/ear. S.C. 10 gave the highest number of kernels/row, whereas, Bashayer gave the highest weight of 100 kernel (gm) .Estimates of phenotypic (ph), genotypic (g) and environmental variance reached highest estimates concerning grain yield/plant (gm) and number of kernels/row. Estimates of genotypic and phenotypic coefficient of variation revealed high values in grain yield/plant (gm), number of rows/ear, number of kernels/row and 100 kernel weight (gm). Grain yield/plant (gm) and days to tasseling showed high broad sense heritability estimates (over 80%).

INTRODUCTION

Maize (zea mays L) is very important to western hemisphere, where it became the dominant food staple in the early civilization of that part of the world. Maize was rapidly accepted as a food staple in many parts of the world because of its large size, good yield, ease of cultivation, versatility and storage character. It is well established that the majority of the people depend mainly on cereal grains as their staple food. In Egypt, maize has been considered an important food in the people's diet, including, charcoal coasted maize, bread, snacks as well as maize doughnut and sweet taxed goods.

The measurements of phenotypic , genotypic and environmental components of variance in maize yield and other characters, have been a matter of great importance . Estimates of the genetic parameters in the

population have impacts in methods of practing section.

The objective of the present investigation was to study the variability and covariability of some introduction of maize with respect to yielding ability and important agronomic characters. The result of this study would be helpful to the plant breeder planning programs for developing high yielding varieties of maize adapted to the Egyptian conditions.

MATERIALS AND METHODS

This research was carried out at Sakha and Gemmeiza Research Stations to study the relative advantage of gene make-up effect as well as genetic behaviour of some maize varieties on the qualitative and quantitative criteria for maize.

The present study include twelve maize varieties, S.C. 10, Watania 1, pioneer 3062, S.C.Bachayer, Dahab, Taba, T.W.C. 310, T.W. C.320, T.W.C. 321, T.W.C. 322, Nefertety and pioneer 3057. A randomized complete block design in four replications was used, each plot was 3x7 m and consisted of ten row, 70 cm apart. Intra-hill spacing was 25 cm. The agricultural practices were done as recommended. At harvest five plants selected randomly from the three guard rows in each plot to collect data on the following characters.

- 1- Number of rows / ear
- 2- Number of kernels / row
 - 3- 100 kernel weight (gm)
 - 4- Grain yield / plant (gm)
 - 5- Days to mid silk
 - 6- Days to tasseling

Statistical analysis

The collected data were subjected to two types of statistical analysis:

- Comparison between means, were done according to Duncan multiple range test (1955).
- 2- Statistical manipulation of the data:
 - Analysis of variance for each location and combined analysis over the two locations for each variable was made according to the technique outlined by Steel et al. (1997)
 - Covariance for each pair of variables were performed for a randomized complete block design according to Snedecor and Cochran (1980).

The expectation of mean squares are given in Table 1 according to Singh and Chuadhary (1985), as a Fixed model.

n, r and a are number of environments, replications and genotypes, respectively.

The genotypic and phenotypic variance (σ^2 g and σ^2 ph) were calculated from the combined analysis as follows:

$$\sigma^2 g = (Ma - Mas)/rn$$

where $\sigma^2 g = Genotypic variance$

Ma = Mean square of varieties

Mas = Mean square of the locations \mathbf{x} varieties interaction

r = Number of replications

n = Number of environments Phenotypic variance, σ^2 ph, was computed according to the following

formula:

$$\sigma^{2} ph = \sigma^{2}g + (\sigma^{2}gs / n) + (\sigma^{2}e / rn)$$
where

$$\sigma^{2}gs = (Mas - Me) / r$$

$$\sigma^{2}e = Me (error mean square)$$

Genotypic and phenotypic covariances were calculated in the same way.

- Genotypic and phenotypic coefficients of variance were calculated according to Burton (1951) as follows:

G.C.V. =
$$\frac{\sqrt{\sigma^2} g}{\overline{x}} x100$$
P.C.V. =
$$\frac{\sqrt{\sigma^2 ph}}{\overline{x}} x100$$

- Broad sense heritability (H%) was calculated as described by Hanson (1963) using the following formula:

(H%) = (genotypic variance / phenotypic variance) x 100

RESULTS AND DISCUSSION

Analysis of variance for each location and the combined analysis over the two locations for yield and other characters were done and presented in Table 2. Combined analysis indicated that Location mean squares were highly significant for the studied traits except number of rows / ear. Replications x locations interaction mean squares were significant for combined analysis in all studied traits except number of kernels / row and number of rows / ear. Genotypes mean squares were highly significant in all traits for each invironment and its combined also. Significant mean squares of the genotypes and locations interaction were founed in all traits indicating that genotypes carried genes with different additive and additive x additive effects which seemed to be inconstant from environment to invironment.

Mean performance of genotypes for all studied traits indicated in diversity as shown in Table 3. Examining this table, we noticed that a genotypes showed fluctuation in yield and its attributes over locations, therefore the genotypes were different in its productivity, this can be attributed to the minor affects of both environment and genetical factors.

Combined analysis over two locations Gemmeiza and Sakha regarding grain yield / plant (gm) and the other traits were reached to the significant level. The highest number of rows / ear was recorded by pioneer 3062 and Dahab. The highest number of kernels / row was recorded by S.C. 10 and T.W.C. 310. The highest 100 kernel weight (gm) was recorded by S.C.Bashayer, S.C. 10, T.W.C. 310, T.W.C. 320, T.W.C. 321 and T.W.C. 322. The highest grain yield/plant (gm) was recorded by S.C.10. The better number of days to mid silk and tasseling date were recorded by Dahab Table 3.

Estimates of phenotypic, genotypic and environmental variance for yield and some characters in maize genotypes obtained from the combined analysis of variance are presented in Table 4. Results showed that the highest values of phenotypic and genotypic variances were 497.67 and 437.01 respectively for grain yield / plant(gm), 4.44 and 2.86 for number of kernels / row.

Table 1: Combined analysis of variance and covariance of a randomized complete block design for Gemmeiza and Sakha locations.

Expectation of Ms Mean cross product $\sigma^2 + r\sigma^2 gs + rn\sigma^2 g$ Mpa $\sigma^2 + r\sigma^2 gs$ Mpa	ZAOOOE	
ts(locations) n-1 product norms $n(r-1)$ ma $\sigma^2 + r\sigma^2 g_S + rn\sigma^2 g$ mpa $notypes$ $(n-1)(r-1)$ mas $\sigma^2 + r\sigma^2 g_S$ mpa	Mean	Expectation of M.C.P.
ts(locations) n-1	cross	
ns $n(r-1)$ Ma $\sigma^2 e + r\sigma^2 g s + rn\sigma^2 g$ Mpa $(n-1)(r-1)$ Mas $\sigma^2 e + r\sigma^2 g s$ Mpas	product	
ins $n(r-1)$ Ma $\sigma^2 e + r\sigma^2 g s + rn\sigma^2 g$ Mpa inotypes $(n-1)(r-1)$ Mas $\sigma^2 e + r\sigma^2 g s$ Mpas		
a-1 Ma $\sigma^2 + r\sigma^2 gs + rn\sigma^2 g$ Mpa (n-1)(r-1) Mas $\sigma^2 e + r\sigma^2 gs$ Mpas		
$(n-1)(r-1)$ Mas $\sigma^2 e + r\sigma^2 gs$ Mpas	Мра	COVe+rCOVas+rnCOVa
1	Mpas	COVe+rCOVgs
n(n-1)(r-1) Me 0.2e Mp		,

Table 2: Mean squares for grain yield, its components and other agronomic traits at the two locations and combined

analysis.

S.O.V.	D.F	D.F		No. of rows/ear	ear	No.	No. of kernels/row	/row	101	100 kernel weight	ight
	Sin.	Com.	L1	L2	Comb.	L1	L2	Comb.	7	L2	Comb.
Locations		-			2.67			74.73			37.75*
Reps x L.	3	9	0.97	0.97	0.97	3.28	0.28	1.78	2.41	12.36	7.39
Genotypes	11	11	5.54	7.9	10.23	30.97	17.11	35.48	13.88	15.22	19.8
LxG		11			3.21			12.6			9.31
Error	33	99	1.09	0.43	0.76	6.16	2.23	4.19	4.03	4.38	4.2
C.V%			7.45	4.77	6.28	5.5	3.19	4.46	5.9	6.4	6.15
S.O.V.	D.F	D.F	Gra	Grain yield /plant [gm]	t [gm]	۵	Days to mid silk	silk	Da	Days to tasseling	ling
	Sin.	Com.					[days]			[days]	1
			L1	L2	Comb.	L1	L2	Comb.	17	L2	Comb.
Locations		-			858.61			100.04			216.0
Reps x L.	3	9	12.01**	75.3**	43.7**	1.13	3.3**	2.22*	1.58	7.41**	4.49**
Genotypes	11	11	3559.01	1680.96	4368.17	7.66	18.14	21.39**	6.82	17.58	20.94
LxG		11			871.89			4.41			3.48
Error	33	99	109.28	87.69	98.49	0.83	0.83	0.83	0.97	0.85	0.91
C.V%			4.94	4.31	4.63	1.42	1.47	1.45	1.48	1.45	1.47

Table 3: Mean performance for yield, yield components and some agronomic traits at Gemmeiza, Sakha nd combined data

Southone	N	No. of rows/ear	ır	Ž	No. of kernels/row	row	100 Ke	100 Kerrier Weight [girl]	[6]
Genorypes			- 1	-	1.3	Comb		L2	Comb.
		77	Comp.		41		0,00	1000	OA ACAL
4.4	44 E obo	14 Ohc	14 25cd	48.6 a	49.5 a	49.05 a	63.12 a	37.8 ab	34.4090
S.C10	14.5 anc	14.000		14000	47 K oh	46 21de	32.5 c	33.8 a	33.15 b
Watania{1}	15.5 a	13.5 c	14.5 C	44.3d-C	41.0 an		24 6	2200	32 7 h
0000	AFF	4602	15753	41.57bc	42.5 d	47.04 g	31.30	20.00	24.1
Pioneer 3062	10.0 d	460.0	44004	48.6.2	47.0 ab	47.8 b	37.75 b	34.35 a	36.05 a
S.C.Bashayer	13.0cd	13.040	1.000	20.01		40 04	22 72 c	32 55ab	33.14 b
Dahah	15.0 ab	16.0 a	15.5 ab	40.82 c	44.0 cd	44.41 9	20.00	2000	000
Jalian	7 1 2 0 7	4250	125 00	45 4 ab	47.75ab	46.58cd	32.25 c	28.55 C	30.4 C
Taba	13.5 D-a	20.01	20.0		47 7Eab	40 0k	34 87hc	32.77ab	33.83ab
0000	12000	13.5 C	13.25 e	48.25 a	47.73ab	40.00	24.0	-	
.W.C.310	20.00	000	40 05 6	42 22hc	49.53	45.91 d-f	33.0c	35.42 a	34.21ab
T.W.C.320	12.5 d	12.00	167.71	44.3400		320 27	24 07ah	22 43	34 14ab
100 0 181 T	12 Kh.d	13 0cd	13.25e	44.6a-c	45.5bc	45.051	24.01 an	20.10	
.W.C. 321	2000		3 30 07	A7 75 0	47 0 ah	47.38bc	34.87ab	33.37 a	34.13ab
T W C 322	12.5 d	12.0 d	167.21	47.10	20.14		07 0760	22 22ah	22 25 h
Information.	14 5 30	12.0 d	13.25 e	45.1 a-c	46.0 bc	45.55 er	37.37 DC	32.3240	20.00
Neierlery	25.		44 7560	42 87hc	48 0 ah	45.44 ef	31.87 c	29.42bc	30.05
Dionogra057	15.5 a	14.0 bc	14./ 200	47.0100	40.0				

		I dialont from		Davs to mi	Days to mid silk [days]		Days to te	Days to tasselling luays	ayo
Genotypes	Grain yie	Grain yield/plant [giii]		Day of the		-		13	Comb
16	-	12	Comb.	7	L2	Comp.			
	1 20	264 22	258 23	63.0 d-f	63.2ob	63.13bc	66.5b-d	64.25b	63.38DC
S.C.10	255.3d	201.24	22.00	62 5 c.f	60.08	61.75de	p-q0'99	61.5 c	63.75de
Natania[1]	240.1a	234.300	431.40	9 30 00	60 75do	6150	65.25cd	61.75c	63.5 e
Pioneer 3062	227.6a	235.6 b	231.6 D	167.70	00.100	70000	CE E od	64 5 h	65 0 bc
C C hachaver	229.03	227.3 b-d	228.1 b	62.75 ef	62.5 DC	07.03Cd	03.3	2000	60.40 6
O.C.Dasilayer	227.00	2126 00	21066	62.5 f	57.75 f	60.13 f	65.0 d	59.25 d	07.131
Dahab	201.04	20.000	70007	6703	66 75 3	66.88a	69.75 a	68.0 a	68.88 a
Taba	205.7 a	180.7 g	193.4 u	01.0 a	7 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	CO Ead	SE FOR	63.75 b	64.63cd
T 181 C 240	21413	209 2 ef	211.6 c	63.25c-1	p-0 c/.19	07.30	20.00		100
.W.C.310	414.1 a	1001	46400	64 75 hc	62 0 b-d	63.38 bc	67.5 b	63.75 b	65.63bc
T.W.C.320	132.3 a	137.761	104.0 0	201.10	60 60	62 5 hc	67.5 h	64.0 b	65.75 b
T W C 321	210.1 a	219.6 c-e	214.9 C	04.5 D-d	04.3 00	20.00	00 7550	CA 25 h	65 5 hc
0000	240 2 2	242 4 d-f	211.2 c	64.5 b-d	62.25 bd	63.38 DC	00.700	04.40	20.00
.W.C.322	410.3 a	2000		CEER	62 75 h	64.13 e	66.75bc	64.25 b	65.5 pc
Nefertety	200.1 a	209.6 er	204.9 C	00.00	2000	62 63 04	66 75hc	63.5 b	65.13bc
Dioneer 3057	205.9 a	209.4 ef	207.6 c	64.25 D-e	01.00	04.00	200		

With regard to environmental effect, results in Table 4 showed that grain yield / plant (gm), 100 kernel weight and number of kernels / row were greatly affected by environmental conditions with variance estimates of 98.45, 4.26 and 4.19 respectively. The lowest values of environmental variance were recorded by number of rows / ear (0.76) and days to mid silk (0.79). These results are similar to those reported by Samia et al (2002). In general estimating genotypic and invironmental variance is of great importance because the variance indicates the performance of various genotypes in given invironmental circumstances. This help the breeder to produce genotypes that represent combinations of genes for a particular invironment.

Estimates of phenotypic coefficient of variation P.C.V., genotypic coefficient of variation (G.C.V.), and Heritability (H%) of twelve genotypes of maize over both locations of Gemmeiza and Sakha are shown in Table 5. The results indicated that grain yield / plant (gm) ranked first with P.C.V. value of 10.4 followed by number of rows / ear (8.15). Meanwhile, tasseling date (days) had the lowest value of P.C.V. being 2.49. Data in Table 5 revealed that grain yield / plant (gm) gave the highest coefficient of genotypic variation being 9.75 followed by number of rows / ear being 6.75, but the lowest G.C.V. were for days to tasseling (2.27). and days to mid silk(2.32)

For heritability broad sense estimates, data in Table 5 indicated that grain yield / plant (gm) recorded highest heritability value 87.8% followed by days to tasseling (83.4) and days to mid silk (79.27). The above mentioned results are in line with those obtained by Barakat (1998). On the other hand selection for a given character among the studying genotypes would seem very effective, but these high estimates of heritability for some characters may be due to both of the high genetic variability which occurred between these genotypes and to the confounded effects of the environmental conditions. It seems reasonable to think of heritability as a measure of degree to which the phenotype reflects the genotype and in so doing accept a nonrestrictive definition, Rasmusson and Glass, (1967).

It could be concluded that grain yield / plant (gm), number of kernels / row and days to tasseling are important characters in maize breeding programs. These traits had high heritabilities as well as high genetic variabilities. Thus breeding for these characters would be effective in improving maize crop. Also this result attributed directly to the presence of more addition genetic variance, (Abul-Saad et al 1998-1999 and Barakat 1998).

Table 4: Estimation of phenotypic, genotypic and environmental variance for yield and some characters for twelve genotypes of maize over the two locations.

		Variance	
Characters	Phenotypic	Genotypic	Environmental
No. of rows/ear	1.28	0.88	0.76
No. of kernels/row	4.44	2.86	4.19
100 kernel weight [gm]	2.62	1.66	4.26
Grain yield/plant [gm]	497.67	437.01	98.45
Days to mid silk [days]	2.7	2.14	0.79
\Days to traceling [days]	2.62	2.18	0.91

Table 5: Estimates of means, phenotypic coefficient of variation [P.C.V.], genotypic coefficient of variation [G.C.V.] and heritability [H%] of yield and some characters for twelve genotypes of maize over the two locations.

Characters	Means	P.C.V.	G.C.V.	H%
No. of rows/ear	13.88	8.15	6.75	68.59
No. of kernels/row	45.95	4.58	3.68	64.48
100 kernel weight [gm]	33.35	4.85	3.86	63.37
Grain yield/plant [gm]	214.49	10.4	9.75	87.8
Days to mid silk [days]	62.96	2.61	2.32	79.27
Days to tasseling [days]	65.06	2.49	2.27	83.4

REFERENCES

- Aboul-Saad, Sh.F.; M.M.A.Ragheb and A.A.Abd El. Aziz (1998-1999). Genetic variance and correletion studies in a yellow, maize population. Bull. Fac. Agric., Cairo Univ., 45:811-826.
- Barakat, A.A. (1998). Genetic on some maize (zea mays L.) populations. Ph.D. Thesis, Fac. Agric, Al-Azhar Univ., Egypt.
- Burton, G.W. (1951). Quantitative inheritance in pear / millet (pennistum glaucum). Agron.J.43:409-417
- Duncan, D.B.(1955). Multiple range and multiple F tests. Biometriecs, 11,1-42.
- Hanson, W.D.(1963). Heritability. Statistical Gen. and PL. Breed. Symposium, North Carolina publication, 982:125-140.
- Johnson, V.A.; K.J.Biever; A. Haunold and J.W. Sehmidt (1966). Inheritance of plant height, yield of grain and other plant and seed characteristics in a cross of hard red winter wheat, *Triticum*
- aestivum L. Crop Sci., 6: 336-338.
- Rasmusson, D.C. and R.L.Glass. (1967). Estimates of genetic and environmental variability in in barley. Crop Sci. 7:185-188.
- Snedecor, G.W. and W.G. Cochran (1980). Statistical methods. Seventh Edition. Iowa State Univ. Press, Ames, Iowa, U.S.A.
- Samia, G.A.; M.S. Sohier and S.M. Salama (2002). Estimating prediction Equations of yield and its characters in maize using some macro climatic and micro environmental factors, J.Agric. Sci. Mansoura Univ., 27(7): 4355-4370,2002.
- Singh, R.K. and B.D.chaudhary, (1985): Biometrical methods in quantitative genetic analysis. Kalyani Publishers, Ludhiana, New Delhi.
- Steel, R.G.D.; T P.J.H. Torri and Dickey, A.D.(1997): Principles and procedures of statistics. MC. Graw hill Book company Inc., New york. 280.

التباين الوراثى والمظهرى والبيئى ودرجة التوريث لبعض التراكيب الوراثية في

احمد مؤمن عبد العزيز ' - محمود عادل عبد الخالق '

- ١- المعمل المركزى لبحوث التصميم والتحليل الاحصائى مركز البحوث الزراعية الجيزة مصر
 - ٢- الادارة المركزية لفحص واعتماد التقاوى -وزارة الزراعة واستصلاح الاراضى

اجرى هذا البحث بهدف تقدير بعض السلوكيات الوراثية لاثنى عشر تركيبا وراثيا من اصناف وهجن الذرة الشامية ، وتمت زراعة التجربة بمحطة البحوث الزراعية بسخا ومحطة البحوث الزراعية بالجميزة والتابعتين لمركز البحوث الزراعية خلال الموسم الصيفى ٢٠٠٣ واوضحت النتائج الأتى:

- ١- اظهر تحليل التباين الفردى والتجميعى اختلافات واسعة ومعنوية بين التراكيب الوراثية لجميع الصفات المدروسة عدى صفة عدد الصفوف في الكوز في التحليل التجميعي، وفي التفاعل بين الجهات والتراكيب الوراثية ظهرت معنوية عالية لجميع الصفات المختبرة.
- ۲- اظهر الصنف دهب تبكيرا في صفة خروج الحريرة والازهار المذكرة عن بقيــة التراكيــب الوراثية..
 - اعطى الصنف بيونير ٣٠٦٢ اعلى عدد للصفوف في الكوز
 - ٤- تميز الهجين فردى ١٠ باعلى عدد من الحبوب في الصف يليه الهجين الثلاثي ٣١٠.
 - وفى صفة وزن ١٠٠ حبة تفوق الصنف بشاير عن بقية التراكيب الوراثية.
 - ٦- اظهر الهجين الفردى ١٠ اعلي كمية محصول على النبات (٢٥٨,٢ جم) عن بقية الاصناف
- ٧- حققت صفة كمية المحصول بالنبات اعلى قيم للتباين المظرى والوراثى حيث كانتا٤٩٧,٦٧٤ و
 ٤٣٧,٠١ على الترتيب.وكذلك صغة عدد الحبوب في الصف حيث كان قيمة التباين المظهرى ٤,٤٤ والتباين الوراثي ٢,٨٦٣ .
- ٨- حققت صفات كمية محصول النبات ووزن ١٠٠ حبة اعلى تاثرا بالظروف البيئيـة بينمـا سجلت صفة عدد الصفوف في الكوز اقل درجة تاثر بالظروف البيئية.
- ٩- حققت صفة كمية محصول النبات اعلى قيمة لمعامل الاختلاف المظهرى يليها صفة عدد الصفوف في الكوز بينما كانت صفة عدد الايام لظهور النورة المذكرة اقل معامل اختلاف مظهري.
- ١٠ حققت صفة كمية محصول النبات اعلى قيمة لمعامل الاختلاف الوراثى يليها صفة عدد الصفوف في الكوز بينما سجلت الدراسة اقل قيمة لمعامل الاختلاف الوراثي في صفتي عدد الايام لظهور الحريرة والنورة المذكرة.
- ١١-سجلت صفة كمية محصول النبات اعلى درجة توريث يليها صفتى عدد الايام لظهور النورة المذكرة وعدد الايام لظهور الحريرة.
- مما ذكر نجد أن صفات كمية محصول النبات وعدد الصفوف في الكوز وعدد الايام لظهور النورة المذكرة وعدد الأيام لظهور الحريرة من الصفات التي يمكن ان يعتد بها في برامج تربية الذرة.