

Stability Parameters for Comparing Bread Wheat Genotypes under Combined Heat and Drought Stress

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A COMPARISON of ten local and twelve introduced wheat accessions was performed on twelve different induced environments (2 years \times 2 sowing dates \times 3 water stresses) to analyze genotype \times environment interactions (G \times E) and estimate stability indices of yield and its components. Mainly the total variations of studied traits were due to the main effects of environmental factors and their interaction, whereas the significant environmental variations were ranged from 10.62% (harvest index) to 43.95% (spike kernels weight). The genotypes differed significantly for all studied traits, moreover these differences ranged from 6.82 % to 50.42% of total variation in 1000-kernel weight and no. of kernels/spike, respectively. G \times E interactions were highly significant and their contributions to the total SS accounted for 40.17, 30.78, 18.17, 22.02, 25.46, 18.61 and 88.46% for heading date, no. of spikes/m², no. of kernels/spike, spike kernel weight, 1000-kernel weight, grain yield/m² and harvest index, respectively. Three genotypes (NGB10893, Sids1 and Giza168) were high yielding and stable for most of the studied traits. Thus, these three genotypes could be promoted to the next extensive breeding programs.

Keywords: Performance, Genotype \times environment interaction, Stability parameters, Wheat.

Wheat (*Triticum aestivum* L.) is a very important cereal crop in Egypt as a source of human food. Growth rate of a human population in Egypt is still relatively high, thus the demand of wheat is being progressively increased. Overcoming the gap between cereal production and consumption depends mainly on horizontal extension of cultivated area of cereals and raising the yield per unit area is encountered by unfavorable conditions such as drought, heat and high salinity of soil. The first step is to identify, the superior tolerant genotypes to be used in the breeding program. However, stable wheat cultivars that are tolerant to different environmental stresses are the ultimate goal of the national wheat research program.

Stable genotypes have the same reactions across the environments. Most favorable stability occurs with high yield or performance (Björnsson, 2002). Increasing genetic gains in yield is possible in part from narrowing the adaptation of cultivars, thus maximizing yield in particular areas by exploiting genotype \times environment interaction (G \times E). G \times E is of major importance,

because it provides information about the effect of different environments on cultivar performance and has a key role for assessment of performance stability of the breeding materials (Moldovan *et al.*, 2000 and Bose *et al.*, 2014).

The regression coefficient (b_i) and the average departure from regression line (S^2d_i) are two mathematical indices for the assessment of stability (Eberhart & Russell, 1966). A genotype with high b_i and S^2d_i reacts readily to changes in the environment and possesses considerable variability, whereas cultivars with a $b_i < 1.0$ and S^2d_i near to 0.00 react weakly to changes in growing conditions and are considered to be stable in yield (Shindin & Lokteva, 2000).

The ability of a crop cultivar to perform reasonably well in variable abiotic stresses is an important trait for both the stability of production under drought conditions (Nachit & Ouassou, 1988). Climate and weather conditions greatly influence the performance of new wheat cultivars both for yield and quality (Wajid *et al.*, 2004; Sharma *et al.*, 2006 and Abdullah *et al.*, 2007). Selection for yield stability over stress sites and years was able to improve the stress resistance (Ortiz-Ferrara *et al.*, 1991). Regression analysis as well as grain yield *per se* could be useful for identifying high yielding thermo tolerance genotypes (Abd-Elghani *et al.*, 1994). In addition, there is a strong evidence that breeding for drought resistance should include both yield and stability improvement (Clark & Townley-Smith, 1984). High temperature is one of the most important abiotic environmental factors during grain filling and may influence both the quantity and quality of the yield (Rehman *et al.*, 2007, Anwar *et al.*, 2011 and Hamidou *et al.*, 2013). All genotypes significantly produced higher grain yields under normal date of sowing compare with late date of sowing, (Sial *et al.*, 2005).

The objectives of the present study are to: (1) Evaluate the magnitude of G×E interactions, (2) Assess the stability parameters of grain yield and its components, (3) Show the degree of linear relationships either between these stability parameters or between their average of the studied traits of the 22 local and introduced wheat accessions and (4) Identify most stable genotypes under abiotic stresses (heat and drought).

Materials and Methods

Plant material and experimental design

Twenty two wheat genotypes from diverse origin including 10 local and 12 introduced genotypes were used in this study (Table 1). The experiments were conducted at the experimental farm of Faculty of Agriculture, Sohag University, Egypt. Two experiments were performed in each season at two different sowing dates (D) 20th November and 20th December, 2011/2012 and 2012/2013 and three water irrigation treatments. The irrigation treatments were; (1) I₁ as normal irrigation in which wheat plants were supplied by water irrigation over all
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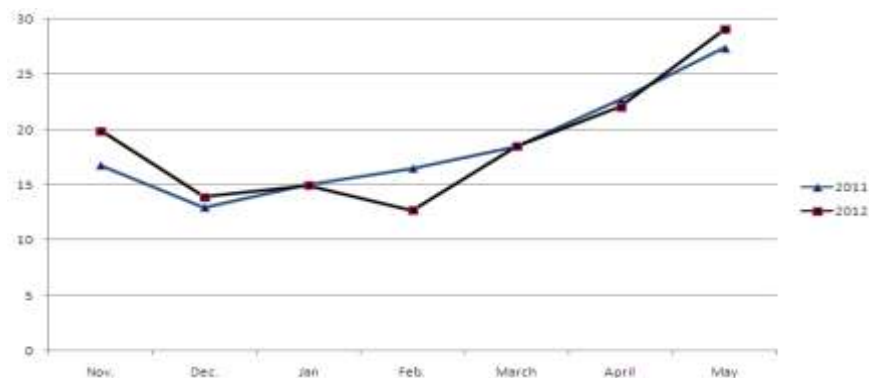
growth stages (10 irrigation frequencies), (2) I_{II} water irrigations were reduced by three irrigations from the beginning of anthesis stage till harvest stage (7 irrigation frequencies) and (3) I_{III} was 5 irrigation frequencies only started from booting stage till the milk-ripe stage.. Overall, twelve different environments were created by manipulating environmental factors which were two years × two dates × three water irrigation treatments. Both the trend of temperature as a climatic factor and the soil status (particle-size distribution, soil texture and chemical analysis) as an edaphically factor reflected the variation between the two winter growing seasons of 2011/2012 and 2012/2013 as shown Fig. 1 and Table 2.

TABLE 1. Pedigree and source of 22 wheat accessions used in the study .

Entry No.	Genotype	Pedigree	Source
1	NGB90533	---	Nord gene bank
2	Giza 164	Kvz/Buha"s"/Kai/Bb=Veery"s"	Egypt
3	Gemmeiza 3		"
4	NGB10992	---	Nord gene bank
5	NGB11185	---	"
6	NGB11418	---	"
7	NGB4769	---	"
8	NGB4823	---	"
9	NGB6404	---	"
10	NGB6406	---	"
11	NGB8188	---	"
12	NGB8218	---	"
13	NGB10893	---	"
14	Sids 1	HD2172/Pavon"s"/1158.57/Maya74"s"	Egypt
15	Sakha 69	Inia/R1.4220//7C/Yr"s"	"
16	NGB10991	---	Nord gene bank
17	Gemmiza 7	---	Egypt
18	Sakha 8	Indus 66 x Norteno"s"/PK3418-65-ISW-OS	"
19	Gemmeiza 1	Maya 74/on//1160/147/3/Bb/Gall/ Chat"s"	"
20	Sahel 1	NS 732/PIMA//Verry"s"	"
21	Giza 165	Cno/Mfd//Man"s"	"
22	Giza 168	Mill/Kauz//Kauz.	"

TABLE 2. Soil status at the experimental farm of Faculty of Agriculture in Sohag University in both seasons (2011/12 and 2012/13) .

Soil status	2011/2012	2012/2013
1- Mechanical properties of surface-soil (0-30 cm)		
Sand (%)	50.35	49.51
Silt (%)	18.72	19.40
Clay (%)	30.93	31.09
Soil texture	Sandy-clay	Sandy-clay
Organic mater (%)	2.81	2.46
Total N (%)	0.153	0.181
2- Chemical properties of surface-soil (0-30 cm)		
N PPM	30	70
P ₂ O ₅	17	47
K ₂ O	778	746
Fe	2.88	6.36
Zn	2.18	3.34
Mn	8.56	12.86
Cu	0.58	1.26
Soluble ions (meq/100g soil (1:5))		
Ca ⁺⁺	0.4	0.6
Mg ⁺⁺	1.6	3.4
Na ⁺⁺	1.73	2.6
K ⁺	0.95	0.37
HCO ₃ ⁻	0.2	0.8
CL	1.6	2.0
SO ₄	2.88	4.17
CaCO ₃ %	5.4	5.6
EC (ds/m) (1:5)	0.5	0.7
pH (1:2.5)	7.8	7.4

**Fig. 1.** The trend of temperature (°C) during growing months of wheat plants in both seasons (2011/12 and 2012/13) .

Split-plot design with three replications was used for every planting date, in which the water irrigation treatments were assigned to the main plots and the genotypes were randomly distributed to the sub plots. Data were recorded on days to 50% heading, number of spikes/m², number of kernels/spike, spike grain weight (gm), 1000-kernel weight (gm), grain yield/m² (gm) and Harvest index (gm) was calculated by using the following formula:

$$\text{Harvest index} = (\text{Grain yield})/(\text{Grain} + \text{straw yield})$$

Statistical analyses

The combined analysis, both in detail and collectively, was performed on the recorded data of grain yield and its components of the 22 genotypes over all the twelve environments according to Gomez & Gomez (1994). The stability parameters C.V. %, b_i , B_i , S^2d_i and r^2 were assessed to each of the 22 genotypes over all environments. Where C.V.% was estimated according to Francis & Kannenberg (1978), b_i and S^2d_i were estimated by using Eberhart & Russell's model (1966) and β_i as described by Perkins & Jinks (1968). Consequently, a stable genotype is a genotype has a regression coefficient of unity ($b_i = 1.0$) and a deviation from regression mean squares equals zero ($S^2d_i = 0$) and hence an ideal genotype would have both a high average performance over a wide range of environments together with stability parameters as defined by Eberhart & Russell (1966). The coefficient of determination (r^2) was proposed to use by Pinthus (1973), because it measures the proportion of a genotype's production variation that is attributable to the linear regression as an index of production stability over environments.

A correlation among stability indices (\bar{X} , C.V. %, b_i , β_i , S^2d_i , and r^2) was performed by using simple correlation (Fisher & Yates, 1953). LSD was computed to compare the differences among means of genotypes while each regression coefficient was tested by t test using the standard error of the corresponding b value. The degree of linear relationships (r 's) was also calculated among the studied traits in this study to examine their mutual effects.

Results and Discussion

Environment-Genotype variations and Gx E interactions

Combined analysis of variance of the studied traits (Table 3) showed that all the variations in the total sum of squares were attributed to the various environmental factors (Y, D and I) and their interactions which always were statistically significant or highly significant with the exception of Y×D interaction of no. of kernels/spike, 1000-kernel weight and harvest index.

The total variations in the studied traits were mostly due to the main effect of the environmental factors (Y, D, and I) which their variations ranged from 10.39% for harvest index to 41.82% for grain weight/spike. This range among the studied wheat features descended from the highest value of the effects of water stresses (15.29 %) to planting dates (3.68%) to years (0.44%). Environmental factors interactions also contributed to a small extent to the total

variations and these contributions extended from 0.23% for harvest index to 4.14% for days to heading and most of these interaction variations were due to D×I interaction over all the studied traits with exception of 100-kernel weight which was much more sensitive to Y×I interaction (Table 3).

The previous findings reflected on the environmental variations which were highly significant and estimated by 31.63, 36.42, 27.98, 43.95, 15.28, 12.98 and 10.62 % of the total variations for days to 50% heading, no. of spikes/ plant, no. of kernels/spike, spike kernel weight, 1000 kernel weight, grain yield plot" and harvest index, respectively (Table 4). These environmental variations were a direct result of : (1) The wide variations in climatic and edaphic factors between the two winter growing seasons (Table 1 and Fig. 1), (2) The effects of optimum and late sowing dates, and (3) The influence of water stress on yield and its components (EI-Morshidy *et al.*, 1998 and 2000). So, these results emphasize that adopting the proper agricultural practices, especially sowing on the proper time with no water stress during the growing season, would visibly reduce a large amount of the environmental variations either by diminishing its main effect or by lessening its interactions or both.

The analyzed data also revealed that there were highly significant differences among genotypes for all the studied features across environments. Moreover, the contribution of the genotype variations to the total sum of squares was ranged from 6.82% (100-kernel weight) to 50.42% (no. of kernels/spike). Obviously, all degrees of G×E interactions were significant with exception of 100-kernel weight. In addition, the range of G×E contributions to the total SS were from 18.17% (no. of kernels/spike) to 88.46% (harvest index) over all the studied plant characters in used environments (Tables 3 and 4). The genetic diversity and the significant G×E interactions imply both sensitivity of genotypes and differential responses of these genotypes to various environments, suggesting the importance of stability parameters assessment of these genotypes under these conditions to identify the best stable suitable genotypes under this range of environments. Saini & Gautam (1990) stated that the range of contributions of both environmental effects and genetic differences to the total SS was from 31.0 to 72.1 % for environmental effects and from 8.3 to 34% for the genetic differences. Moreover, Nachit *et al.* (1992) showed that the mean squares of environments, genotypes and G×E interactions of the analysis of variance of wheat genotypes were highly significant and accounted for 89.3%, 0.5% and 10.2% of the treatment combinations SS, respectively. The results in this study are generally in harmony with previous studies (EI-Defrawy *et al.*, 1994; Kheiralla & Ismail, 1995; Ismail, 1995, EI-Morshidy *et al.*, 1998 and 2000; Kheiralla *et al.*, 2004 and Bose *et al.*, 2014).

TABLE 3. Mean squares of the combined analysis of variance for each of the studied traits over all wheat genotypes (22 G.) and used environments (12 E.).

Source of variation	d.f.	Mean squares						
		Days to 50% heading	Number of spikes / m ²	Number of kernels/spike	Spike kernels weight (gm)	1000-kernel weight (gm)	Grain yield m ²	Harvest index (gm)
Year (Y)	1	1089.70**	20597.88**	3846.95**	8.91**	22219.72**	51136.99**	3911.29**
Rep/Y	4	6.21	1260.50	23.45	0.029	205.12	374.162	4.04
Date (D)	1	16281.92**	396500.06**	5172.95**	4.18**	1909.66**	448113.58**	4326.85**
Y x D	1	168.21**	7622.28**	0.021 ^{NS}	0.10**	148.54*	10662.67**	0.56 ^{NS}
Error (a)	4	64.05	254.91	5.71	0.0069	82.63	57.202	1.87
Irrigation (I)	2	744.56**	131296.04**	7663.59**	23.07**	3054.91**	702178.44**	3229.30**
Y x I	2	272.67**	3725.64**	85.71**	0.16**	1052.83**	1417.51**	40.82**
D x I	2	1013.09**	11941.03**	172.83**	1.27**	283.84**	52127.87**	82.27**
Y x D x I	2	50.18 ^{NS}	678.30**	15.71*	0.03**	290.25**	2409.81**	7.32**
Error (b)	16	43.08	133.44	10.65	0.005	67.91	143.08	3.01
Genotype (G)	21	1475.60**	66153.47**	2136.55**	2.17**	715.35**	142649.44**	856.23**
Y x G	21	119.63**	1565.88**	20.65**	0.026**	265.06**	3241.46**	11.86**
D x G	21	208.59**	33309.52**	144.92**	0.244**	340.65**	21333.51**	51.68**
Y x D x G	21	94.33**	971.13**	11.81**	0.020**	249.95**	2813.33**	6.25**
I x G	42	189.23**	22492.97**	211.78**	0.352**	252.63**	40787.62**	172.14**
Y x I x G	42	76.16**	1609.69**	15.10**	0.022**	213.22**	2692.86**	9.19**
D x I x G	42	115.90**	11387.62**	60.68**	0.207**	237.07**	9655.17**	70.27**
Y x D x I x G	42	63.53**	1319.21**	8.74**	0.017**	204.08**	2322.15**	5.49*
Error (c)[Pooled Error]	504	36.71	98.91	5.24	0.0039	54.64	1277.85	2.07
Total	791							
C.V.		6.57	2.74	5.47	3.95	28.44	7.74	4.84

*P ≤ 0.05 **P ≤ 0.01 ^{NS} = not significant.

TABLE 4. Summary of the joint regression analysis of variances for studied traits in this research .

S.O.V.	d.f.	Mean Squares							
		Days to 50% heading	Number of spikes/ m ²	Number of kernels/spike	Spike kernels weight (gm)	1000-kernel weight (gm)	Grain yield m ²	Harvest index (gm)	
Environment(E)	11	72.81**	338182.02**	263.24**	5.66**	3058.34**	820561.87**	1032.35**	
Rep/E	24	40.42803	341.53	11.96	0.0093	176.56	167.28	3.63	
Genotype (G)	21	475.59**	66153.48**	2136.55**	2.18**	715.35**	142649.45**	22.88*	
E x G	231	119.29**	9951.41*	69.99**	0.135**	242.70**	12573.08**	432.15**	
E+(ExG)	242	158.08**	24870.99**	169.69**	0.387**	370.68**	49299.84**	459.42**	
E (linear)	1	10700.85**	3720002.25**	24895.59**	62.29**	33641.64**	9026180.51**	11355.79**	
G x E (linear)	21	667.02**	106943.60**	665.99**	0.56**	777.93**	98600.73**	4526.70**	
Pooled deviation	220	61.58**	340.73**	9.93**	0.089**	180.58	3789.85**	21.66	
Pooled error	528	36.71	98.91	5.24	0.0039	210.64	1277.85	2.08	
Total	791								
Linear proportion of variance (%)		96.34	20.349	97.92	99.57	98.49	15.2364	83.23	

*P ≤0.05 **P≤0.01 ***P≤0.001 NS = not significant.

Joint regression analyses

Analysis of variance of the studied traits over all environments and genotypes when stability parameters are estimated for each genotype across all environments are presented in Table 4. All mean squares of E+G×E were highly significant and the contributions of their SS to the total SS over all traits ranged from 40.74% (1000-kernel weight) to 98.53% (harvest index). In fact, (E+G×E) ss for each trait is only a makeup of the two parts; Ess and G×Ess of the same trait. Ess is completely represented by E (linear) ss which its mean square was highly significant for the studied traits, emphasizing again that there were much differences among environments and their influences would remarkably reflect on the studied traits. Also, the partition of G×Ess interaction of the studied traits into its two components; *i.e.*, regression ss [G×E (linear)ss] and deviations from regression ss [pooled deviations], demonstrated that: (1) G×E (linear) ss's for five out of the six studied traits were statistically significant, implying that it could be proceeding in assessment of stability parameters using Eberhart & Russell's model (1966). (2) The contributions of G×E (linear) ss's to G×E interaction ss's over all the studied traits ranged from 29.14% (1000-kernel weight) to 96.74% (no. of spikes/plant), emphasizing the importance of the stable parameter S^2d_i as defined by the previous studies. (3) The highly significance of deviation mean squares in this research pointed out to both considerable variations among genotypes in their stabilities and also visual variability of genotypes relative ranking from one environment to another. These data are coincident with Eberhart & Russell (1966), Nachit *et al.* (1992), Kheiralla & Ismail (1995), Ismail (1995), EI-Morshidy *et al.* (1998 and 2000), Kheiralla *et al.* (2004), Mustâţea1 *et al.* (2009) and Koumber *et al.* (2011).

Estimated stability parameters

It is important to report that plant breeders in executing selection programs would prefer to select genotypes with high average performance and most stable across various environments. Our data in Table 6 suggest that it is possible to select from wheat accessions in this study using a combination of both response and stability production indices. Langer *et al.* (1979) stated the same conclusion in oat varieties. Therefore, in the present study genotype will be selected if it has higher mean performance than the grand mean, higher r^2 , low c. v. %, $b_i = 1$ and smaller S^2d_i . This is, in brief, because higher r^2 means that the linear model fits the data with the other parameters which will pronounce on well performed and the most stable suitable genotype.

Days to 50% heading

The studied genotypes appeared to have a wide range of variability in mean heading dates as shown in Table 5 and Fig. 2a. The range of heading dates among genotypes was about 23 days with an average of 91.79 days. Obviously, the C.V.'s% among genotypes was low, therefore, the stability will be determined on the basis of r^2 , b_i and S^2d_i (Table 5). Sixteen genotypes were stable due to their b_i 's and S^2d_i 's did not differ from a unit and the zero,

respectively plus showing high r^2 . Six of the 22 studied genotypes (2, 5, 6, 13, 20 and 22) are considered as ideal in stability parameters although they were slightly late in heading. This significant deviation from regression for heading date was attributed by Joppa *et al.* (1971) to specific cultivar \times location or other specific cultivar \times environment interaction. These results are generally in line with those reported by EI-Defrawy *et al.* (1994), Kheiralla & Ismail (1995), Ismail (1995) and EI-Morshidy *et al.* (1998 and 2000).

Number of spikes/plant

Out of the 22 studied genotypes, 8 (2, 6, 11, 13, 14, 15, 16 and 19) showed acceptable production statistics of both responses and stability for the number of spikes/plant (Table 5 and Fig. 2b). They demonstrated high or insignificant average comparing to the grand mean, low C.V.% values, higher r^2 , and insignificant b_i and S^2di . Similar results were reported by Salem *et al.* (1990), Ismail (1995), EI-Morshidy *et al.* (1998 and 2000) and Kheiralla *et al.* (2004).

Number of kernels/spike

The mean no. of kernels/spike ranged from 27.53 (genotype 14) to 56.57 (genotype 20) with an average of 42.10 (Table 5 and Fig. 2c). Six genotypes (3, 14, 17, 18, 21 and 22) have high average comparing to the grand mean with low C.V. values, high r^2 , and insignificant b_i and S^2di . Similar results were reported by Bansal & Sinha (1991 b). The data also revealed that genotypes with higher b_i gave higher number of kernels/spike, this is due to the positively significant association between \bar{X} and b_i ($r=0.448^*$). These findings are in agreement with those obtained by Salem *et al.* (1990), EI-Morshidy *et al.* (1998 and 2000) and Mustățea *et al.* (2009).

Spike kernels weight (gm)

Average spike kernels weight ranged from 1.19 (genotype 12) to 2.08 gm (genotype 14) with an average of 1.59 gm (Table 5 and Fig. 2d). Using the parameters b_i , S^2di , C.V.% and r^2 as selection criteria to the stability in this trait associated with high mean. Six stable genotypes (2, 6, 8, 14, 18 and 22) were selected when compared with the average over all genotypes. According to Eberhart & Russell (1966), these genotypes may be considered superior. Again, the relationship between \bar{X} and b_i was highly positively significant ($r=0.523^{**}$) (Table 6). Similar results were reported by EI-Morshidy *et al.* (1998 and 2000).

1000-kernels weight (gm)

The studied accessions differed in their averages of 1000-kernels weight which ranged from 29.83 (genotype 3) to 45.62 gm (genotype 13) with an average of 37.76 gm (Table 5, and Fig. 3e). Eighteen genotypes could be defined as the most stable suitable genotypes according to selection criteria. These genotypes were characterized by having low C.V. %, high r^2 , insignificant b_i and S^2di . Additionally, eight genotypes (2, 8, 12, 11, 13, 14, 16 and 22) were the most desired genotypes for 1000-kernels weight and showed high mean

performance when compared with grand mean beside their stability. Noticeably, the relationship between \bar{X} and b_i , for this trait (Table 6) was positively significant (0.446*), indicating that the well performed genotypes (with higher b_i) across varying environments would produce higher 1000-kernels weight. Similar results were obtained by Salem *et al.* (1990), Ismail (1995), EI-Morshidy *et al.* (1998 and 2000) and Mustățea1 *et al.* (2009).

Grain yield/m² (gm)

The studied genotypes appeared to have a wide range of variability in average grain yield as shown in Table 5 and Fig. 3f. Mean grain yield ranged from 362.31 gm/m² (genotype 12) to 579.78 gm/m² (genotype 2) with an average of 462.05 gm/m². Concerning the estimated stability parameters (C.V. %, r^2 , b_i and S^2d_i) for this trait, most of the C.V.'s % for the studied genotypes were close to the acceptable upper limit in the agriculture research (<25%), this was due to the sensitivity of yield to different environments as well it is actually a net product of the physiological processes within a plant. Coefficient of determinations was also so high and ranged from 0.80 to 0.97 over all genotypes. In a simultaneous consideration to the stability parameters b_i and S^2d_i , out of the 22 genotypes 15 were stable over all the studied environments; *i.e.* their b_i and S^2d_i were insignificant. More than half of these stable genotypes (7) showed high yield; *i.e.* above the grand mean. According to ascending orders of yields to these genotypes, the stable genotypes were 8 (523.56 gm), 10 (494.69 gm), 11 (525.11 gm), 13 (521.06 gm), 14 (502.39 gm), 18 (477.97 gm) and 22 (567.75 gm), (Table 5). It was clear to notice that genotypes no. 8, 10, 12, 13, 15, 18 and 20 were stable and exhibited low average response to different environments ($b_i < 1.0$), they considered relatively better in stressed environments. The genotypes no. 1, 3, 11, 17, 19 and 21 performed consistently better in favorable environments ($b_i > 1$). The most desired and stable genotypes can be considered when their regression coefficient equal one ($b_i = 1$) with lower values of S^2d_i (Eberhart & Russell, 1966), accordingly in this study both genotypes no. 14 and 22 were considered as desired and stable for grain yield when compared with grand mean. The large variation in mean grain yield, C.V. %, b_i and S^2d_i indicated different responses of genotypes to environmental changes (Akçura *et al.*, 2005). Our results are in line with those obtained by Bansal & Sinha (1991b), Abd EI-Ghani *et al.* (1994), Kheiralla & Ismail (1995), Ismail (1995), EI- Morshidy *et al.* (1998 and 2000), Mustățea1 *et al.* (2009), Anwar *et al.* (2011) and Koumber *et al.* (2011).

Harvest index

Data in Table 5 and Fig. 3g indicated that the mean of harvest index ranged from 23.13 (genotype 4) to 39.47 gm (genotype 22) with an average of 29.78 gm. The results showed that sixteen genotypes were matched with selection criteria to be defined as the most stable suitable genotypes. These genotypes showed low C.V. %, high r^2 , and insignificant b_i and S^2d_i . The most desired genotypes for harvest index were 3, 8, 10, 14, 16, 18 and 22 due to their high mean performance when compared with grand mean and their stability.

TABLE 5. Means and estimated stability parameters in this study of the studied traits of each genotype (G) of wheat over all the used environments (E) .

No. of Genotype	Days to 50% heading						Number of spikes/ m ²					
	\bar{X}_1	C.V. %	b \pm S.E.	β_1	S ² d _i	r ² %	\bar{X}_2	C.V. %	b \pm S.E.	β_1	S ² d _i	r ² %
1	100.2	1.28	1.14 \pm 0.16	0.14	0.59	0.79	333.33	6.19	1.41 \pm 0.10	0.4*1	6.23	0.60
2	87.58	1.06	0.99 \pm 0.16	-0.01	0.57	0.84	424.03	5.10	0.70 \pm 0.07	-0.30	2.58	0.86
3	93.97	1.12	0.49 \pm 0.18	-0.51*	0.67	0.56	325.17	4.44	1.27 \pm 0.24	0.27	31.75	0.92
4	96.28	2.82	0.62 \pm 0.41	-0.38	3.70	0.80	323.97	4.32	1.53 \pm 0.17	0.53*	16.91	0.49
5	76.94	0.98	1.14 \pm 0.18	0.14	0.71	0.81	363.19	9.39	2.60 \pm 0.24	1.60*	31.49	0.82
6	91.28	1.63	1.02 \pm 0.26	0.02	1.49	0.85	264.14	7.48	1.33 \pm 0.33	0.33	63.07	0.90
7	92.97	1.86	0.04 \pm 0.29	-0.96*	1.82	0.53	373.86	9.09	0.16 \pm 0.05	-0.84*	1.65	0.96
8	95.36	3.01	0.79 \pm 0.48	-0.21	5.03	0.90	359.50	8.26	1.37 \pm 0.16	0.37*	14.92	0.54
9	94.19	1.65	0.54 \pm 0.26	-0.46	1.47	0.74	386.83	4.50	1.43 \pm 0.32	0.43*	59.40	0.65
10	93.50	2.38	0.57 \pm 0.33	-0.43	2.34	0.75	417.72	5.12	0.49 \pm 0.08	-0.51*	3.90	0.79
11	82.28	1.76	0.48 \pm 0.27	-0.52*	1.64	0.56	382.67	2.36	1.11 \pm 0.23	0.11	29.65	0.92
12	95.17	2.00	1.20 \pm 0.32	0.20	2.20	0.92	328.50	7.42	1.45 \pm 0.23	0.45*	29.87	0.65
13	89.67	1.25	0.95 \pm 0.19	-0.05	0.77	0.93	411.22	5.21	1.09 \pm 0.14	0.09	11.38	0.74
14	95.53	2.75	1.53 \pm 0.44	0.53*	4.21	0.53	419.14	5.29	0.72 \pm 0.13	-0.28	9.38	0.87
15	94.78	2.11	1.37 \pm 0.33	0.37	2.45	0.71	395.28	4.31	0.81 \pm 0.20	-0.19	22.82	0.80
16	88.11	3.54	1.83 \pm 0.52	0.83*	5.93	0.50	429.42	9.16	0.71 \pm 0.11	-0.29	7.00	0.85
17	92.17	1.04	0.76 \pm 0.16	-0.24	0.56	0.88	324.75	6.27	0.28 \pm 0.15	-0.72*	12.42	0.70
18	90.39	1.69	2.01 \pm 0.25	1.01*	1.42	0.61	330.14	4.21	0.59 \pm 0.11	-0.49*	7.69	0.55
19	95.22	2.87	1.45 \pm 0.46	0.45	4.56	0.90	365.94	2.26	0.83 \pm 0.16	-0.17	14.30	0.92
20	87.94	3.03	1.06 \pm 0.44	0.06	4.33	0.70	350.47	7.13	0.52 \pm 0.07	-0.48*	3.27	0.80
21	97.72	2.25	1.15 \pm 0.37	0.15	2.94	0.85	359.42	1.35	1.25 \pm 0.21	0.25	24.04	0.93
22	88.11	1.97	0.86 \pm 0.29	-0.14	1.84	0.84	314.67	6.14	0.36 \pm 0.07	-0.64*	2.80	0.89
Grand mean	91.79						362.88					
LSD _{0.05}	2.79						4.59					

TABLE 5 .Cont .

No. of Genotype	Number of kernels/spike			Spike kernels weight (gm)			r ²					
	\bar{X}_3	C.V. %	b \pm S.E.	β_i	S ² d _i	r ² %		\bar{X}_4	C.V. %	b \pm S.E.	β_i	S ² d _i
1	42.37	1.89	0.60 \pm 0.12	-0.40*	0.26	0.84	1.32	6.19	0.48 \pm 0.11	-0.52*	0.001	0.85
2	34.66	1.77	0.99 \pm 0.16	-0.01	0.47	0.92	1.74	4.31	0.76 \pm 0.10	-0.24	0.004	0.83
3	49.12	3.03	0.85 \pm 0.25	-0.15	1.16	0.85	1.60	10.36	0.44 \pm 0.22	-0.56*	0.057**	0.75
4	53.56	1.25	0.55 \pm 0.07	-0.45*	0.10	0.81	1.40	15.59	0.23 \pm 0.29	-0.77*	0.084**	0.74
5	39.17	3.89	1.64 \pm 0.25	0.64*	1.22	0.84	1.60	16.83	0.90 \pm 0.36	-0.10	0.121**	0.91
6	47.45	2.93	0.64 \pm 0.23	-0.36*	1.02	0.95	1.68	8.93	0.95 \pm 0.20	-0.05	0.004	0.92
7	39.78	1.92	0.75 \pm 0.13	-0.25	0.30	0.92	1.55	5.81	0.78 \pm 0.12	-0.22	0.002	0.85
8	35.87	5.25	0.94 \pm 0.31	-0.06	1.86	0.87	1.66	5.96	0.87 \pm 0.13	-0.13	0.002	0.90
9	43.22	3.50	1.70 \pm 0.25	0.70*	1.20	0.75	1.65	6.55	1.45 \pm 0.14	0.45*	0.024**	0.96
10	30.56	1.97	1.06 \pm 0.14	0.06	0.36	0.83	1.60	6.61	1.11 \pm 0.14	0.11	0.022**	0.76
11	32.62	1.55	1.00 \pm 0.08	0.00	0.13	0.91	1.48	10.34	0.89 \pm 0.20	-0.11	0.004	0.83
12	53.87	5.08	0.38 \pm 0.23	-0.62*	1.03	0.75	1.19	8.70	0.39 \pm 0.14	-0.61*	0.027**	0.66
13	35.42	3.17	1.00 \pm 0.24	0.00	1.10	0.78	2.06	7.46	1.73 \pm 0.21	0.73*	0.049**	0.79
14	46.87	5.51	0.96 \pm 0.50	-0.04	4.63	0.65	2.08	12.26	1.30 \pm 0.34	0.30	0.006	0.70
15	27.53	2.47	1.43 \pm 0.19	0.43*	0.70	0.94	1.83	5.82	1.49 \pm 0.14	0.49*	0.002	0.82
16	38.55	4.01	1.63 \pm 0.38	0.63*	2.69	0.68	1.80	5.33	1.36 \pm 0.13	0.36*	0.028**	0.94
17	41.69	3.51	1.28 \pm 0.24	0.28	1.12	0.93	1.38	7.61	1.32 \pm 0.14	0.32*	0.027**	0.85
18	47.23	2.41	1.16 \pm 0.19	0.16	0.68	0.72	1.67	9.03	1.24 \pm 0.20	0.24	0.004	0.94
19	42.05	3.55	0.80 \pm 0.18	-0.20	0.62	0.86	1.23	10.49	1.12 \pm 0.17	0.12	0.033**	0.76
20	56.57	1.96	0.68 \pm 0.13	-0.32*	0.30	0.69	1.34	3.64	0.95 \pm 0.07	-0.05	0.001	0.93
21	45.84	2.68	0.93 \pm 0.16	-0.07	0.47	0.85	1.28	3.87	1.25 \pm 0.07	0.25	0.001	0.89
22	42.12	2.08	1.03 \pm 0.15	0.03	0.41	0.97	1.81	8.37	0.98 \pm 0.20	-0.02	0.004	0.86
Grand mean	42.10						1.59					
LSD _{0.05}	1.06						0.03					

TABLE 5 .Cont .

No. of Genotype	1000-kernel weight (gm)					Grain yield/ m ²						
	\bar{X}_5	C.V. %	bi±S.E.	β_1	S ² d _i	r ² %	\bar{X}_6	C.V. %	bi±S.E.	β_1	S ² d _i	r ² %
1	34.14	2.28	0.86±0.13	-0.14	0.86	0.84	372.92	24.25	1.27±0.19	0.27	69.51	0.87
2	41.08	1.93	0.77±0.11	-0.23	0.60	0.79	579.78	5.17	0.65±0.05	-0.35*	18.61	0.83
3	29.83	3.32	0.77±0.17	-0.23	1.39	0.92	434.81	10.76	1.04±0.80	0.04	28.96	0.85
4	35.53	2.80	0.68±0.17	-0.32*	1.40	0.88	367.67	17.95	0.55±0.11	-0.45*	40.66	0.91
5	34.53	2.90	0.93±0.17	-0.07	1.43	0.92	466.69	22.00	2.18±0.21	1.18*	77.69	0.86
6	37.81	9.65	0.22±0.14	-0.78*	14.9	0.75	453.83	8.46	0.26±0.06	-0.74*	23.68	0.93
7	33.58	1.52	0.91±0.10	-0.09	0.55	0.70	438.28	9.04	0.44±0.06	-0.56*	24.44	0.88
8	40.22	3.53	0.95±0.21	-0.05	2.28	0.91	523.56	20.39	0.96±0.24	-0.04	88.34	0.90
9	36.53	3.68	1.40±0.22	0.40*	2.55	0.84	396.17	18.17	1.44±0.12	0.44*	44.34	0.88
10	44.95	9.85	1.81±0.34	0.81*	11.1	0.73	494.69	9.34	0.75±0.08	-0.25	28.37	0.81
11	40.45	2.60	0.93±0.18	-0.07	1.56	0.95	525.11	17.48	1.18±0.15	0.18	56.41	0.92
12	32.21	3.73	1.22±0.27	0.22	3.84	0.96	362.31	21.80	0.79±0.19	-0.21	70.94	0.93
13	45.62	3.20	1.29±0.24	0.29	3.02	0.82	521.06	23.38	0.87±0.20	-0.13	75.20	0.94
14	42.51	1.70	1.09±0.11	0.09	0.58	0.87	502.39	9.55	1.05±0.08	0.05	29.56	0.82
15	36.04	1.85	1.03±0.12	0.03	0.79	0.90	444.67	14.98	0.90±0.11	-0.10	40.91	0.80
16	44.06	3.11	1.14±0.17	0.14	1.43	0.97	528.00	12.84	1.46±0.11	0.46*	41.96	0.95
17	34.86	1.60	0.95±0.09	-0.05	0.44	0.90	459.61	11.88	1.05±0.09	0.05	33.70	0.83
18	35.08	2.67	0.92±0.16	-0.08	1.24	0.82	477.97	15.06	0.95±0.12	-0.05	44.34	0.97
19	35.93	3.91	1.12±0.23	0.12	2.79	0.75	425.53	14.24	1.17±0.10	0.17	37.25	0.93
20	35.22	2.39	0.77±0.14	-0.23	0.99	0.91	422.78	8.37	0.75±0.06	-0.25	21.80	0.84
21	36.02	3.08	1.08±0.19	0.08	1.75	0.79	399.44	15.92	1.21±0.11	0.21	39.06	0.89
22	44.45	2.36	1.16±0.18	0.16	1.57	0.78	567.75	7.40	1.04±0.07	0.04	26.03	0.94
Grand mean	37.76						462.05					
LSD _{0.05}	3.41						16.51					

TABLE 5. Cont .

No. of Genotype	Harvest index (gm)					S ² di	r ² %
	\bar{X}_7	C.V.%	b±S.E.	β_i	S ² di		
1	26.99	11.56	1.21±0.52	0.20	6.02	0.92	
2	34.61	5.55	1.68±0.32	0.68*	2.34	0.91	
3	31.83	4.52	0.88±0.24	-0.12	1.26	0.85	
4	23.13	5.71	0.49±0.22	-0.51*	1.08	0.79	
5	25.61	6.79	1.19±0.29	0.19	1.96	0.90	
6	23.61	3.81	0.60±0.15	-0.40*	0.49	0.75	
7	32.51	3.69	0.48±0.20	-0.52*	0.86	0.98	
8	33.26	2.16	0.73±0.12	-0.27	0.33	0.93	
9	24.43	7.86	0.69±0.32	-0.31	2.28	0.90	
10	30.34	4.94	1.00±0.25	0.00	1.39	0.84	
11	26.04	5.30	0.88±0.23	-0.22	1.18	0.83	
12	26.33	2.96	1.25±0.13	0.25	0.37	0.87	
13	27.94	3.44	0.76±0.16	-0.24	0.57	0.78	
14	37.15	5.98	1.31±0.37	0.31	3.08	0.90	
15	30.17	4.77	0.51±0.24	-0.49*	1.28	0.93	
16	38.88	4.94	1.42±0.32	0.42	2.34	0.70	
17	30.32	6.53	1.45±0.33	0.45*	2.48	0.76	
18	32.98	4.37	1.30±0.24	0.30	1.32	0.84	
19	29.32	4.30	1.18±0.21	0.18	0.96	0.77	
20	24.52	5.14	0.64±0.21	-0.36	0.98	0.93	
21	25.64	3.28	0.80±0.14	-0.20	0.47	0.96	
22	39.47	3.50	1.24±0.23	0.24	1.23	0.90	
Grand mean	29.78						
LSD _{0.05}	0.66						

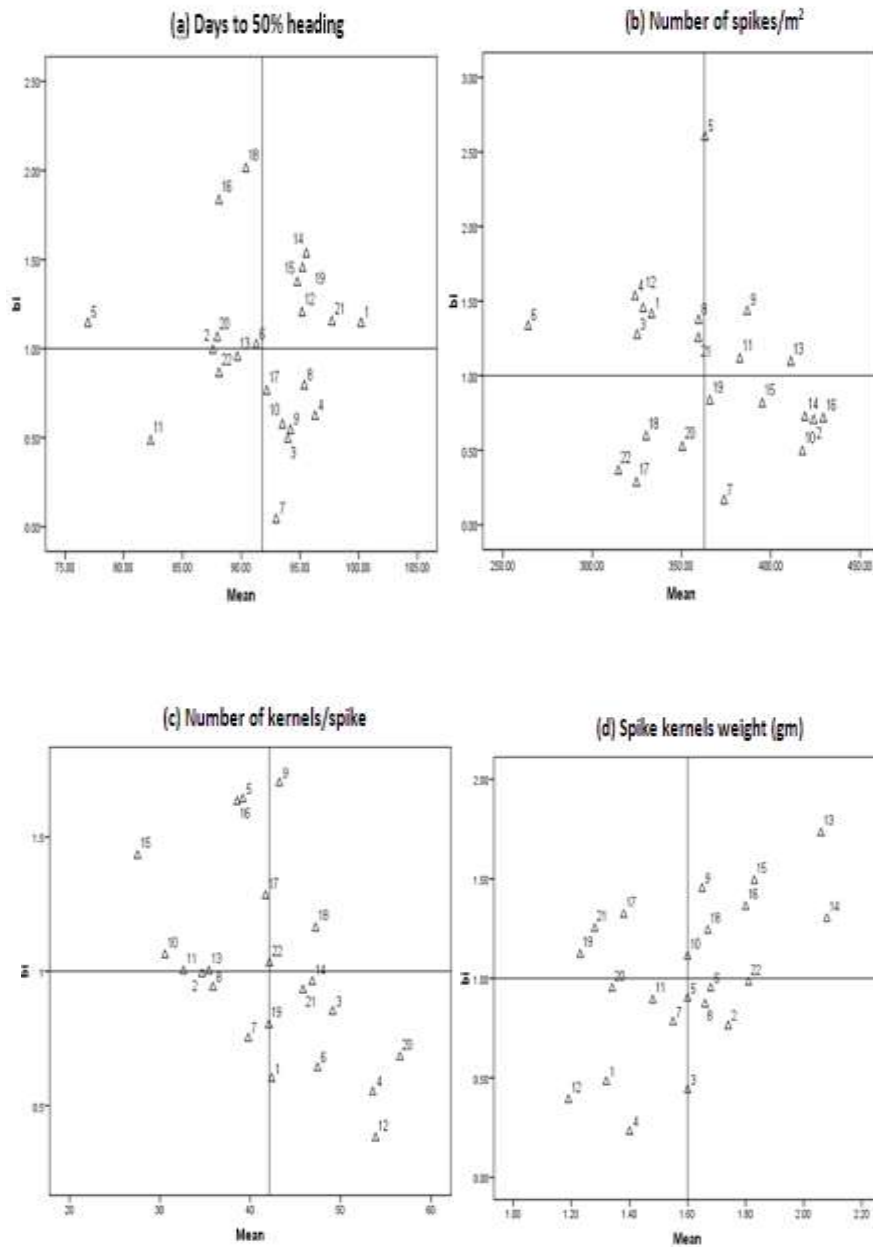


Fig. 2. Present graphically the relationships between the stability parameters (b_i) and its mean performance of each genotype of the 22th genotypes for (a) Days to 50% heading, (b) Number of spikes/plant, (c) Number of kernels/spike and (d) Spike kernels .

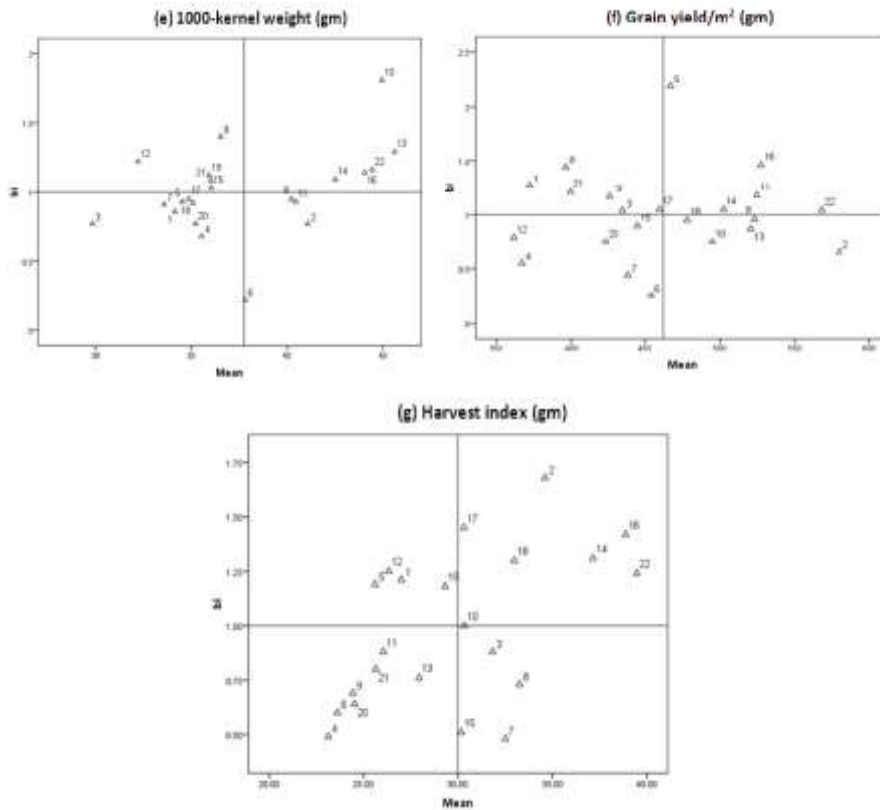


Fig. 3. Present graphically the relationships between the stability parameters (b_i) and its mean performance of each genotype of the 22th genotypes for (e) 1000-kernels weight, (f) Grain yield/m² and (g) Harvest index.

The data also indicated that genotypes with high b_i , gave higher harvest index (Table 6), indicated by highly positive significant association between \bar{X} and b_i ($r = 0.556^{**}$). These results are in agreement with those obtained by Salem *et al.* (1990).

Further investigation to the data was also done using all possible combinations of pairwise simple correlations (r) among the studied traits on the basis of the average of each genotype over all environments (Table 7). Understanding these correlations among traits is of paramount importance to increase the yield. Wherever these correlations serve as a guide for incorporating the economic characters and often a reduction in one yield component may be compensated by an increase in another (Bansal & Sinha, 1991 b and EI-Defrawy *et al.*, 1994).

TABLE 6. Simple correlations among all combinations of each trait' s.

Index 1	Index 2	Plant characters							
		Days to 50% heading	Number of spikes / m ²	Number of kernels/spike	Spike kernels weight (gm)	1000-kernel weight (gm)	Grain yield m ²	Harvest index (gm)	
χ	c.v.%	-0.168 ^{NS}	-0.101 ^{NS}	0.063 ^{NS}	0.026 ^{NS}	0.427*	-0.391 ^{NS}	-0.216 ^{NS}	
	bi or β_i	0.054 ^{NS}	0.033 ^{NS}	0.448*	0.523**	0.446*	0.026 ^{NS}	0.556**	
	S ² di	-0.039 ^{NS}	0.189 ^{NS}	0.511**	0.358 ^{NS}	0.455*	-0.101 ^{NS}	0.145 ^{NS}	
	r ²	-0.032 ^{NS}	-0.193 ^{NS}	0.180 ^{NS}	-0.152 ^{NS}	-0.289 ^{NS}	-0.072 ^{NS}	-0.054 ^{NS}	
c.v.%	bi or β_i	0.288 ^{NS}	0.664**	-0.318 ^{NS}	-0.305 ^{NS}	0.173 ^{NS}	0.407 ^{NS}	0.236 ^{NS}	
	S ² di	0.979**	0.917**	0.789**	0.0906**	0.981**	0.942**	0.909**	
	r ²	-0.169 ^{NS}	0.647**	-0.193 ^{NS}	-0.318 ^{NS}	-0.181 ^{NS}	-0.304 ^{NS}	-0.411 ^{NS}	
bi or β_i	S ² di	0.340 ^{NS}	0.579**	0.285 ^{NS}	-0.089 ^{NS}	0.277 ^{NS}	-0.013 ^{NS}	0.479*	
	r ²	-0.099 ^{NS}	-0.305 ^{NS}	-0.132 ^{NS}	-0.039 ^{NS}	-0.332 ^{NS}	0.164 ^{NS}	-0.341 ^{NS}	
S ² di	r ²	-0.197 ^{NS}	-0.581**	0.247 ^{NS}	-0.362 ^{NS}	-0.185 ^{NS}	-0.353 ^{NS}	-0.379 ^{NS}	

*P ≤ 0.05 **P ≤ 0.01 NS = not significant.

TABLE 7. All possible relations between the averages of the studied traits on the basis of genotype mean over all the used environments in this study.

The average	\bar{X}_1	\bar{X}_2	\bar{X}_3	\bar{X}_4	\bar{X}_5	\bar{X}_6	\bar{X}_7
\bar{X}_1	-	-0.140 ^{NS}	0.228 ^{NS}	-0.229 ^{NS}	-0.226 ^{NS}	-0.529 ^{**}	-0.043 ^{NS}
\bar{X}_2	-	-	-0.611 ^{**}	0.443 [*]	0.539 ^{**}	0.419 [*]	0.355 ^{NS}
\bar{X}_3	-	-	-	-0.566 ^{**}	-0.720 ^{**}	-0.719 ^{**}	-0.525 ^{**}
\bar{X}_4	-	-	-	-	0.637 ^{**}	0.669 ^{**}	0.538 ^{**}
\bar{X}_5	-	-	-	-	-	0.757 ^{**}	0.456 ^{**}
\bar{X}_6	-	-	-	-	-	-	0.681 ^{**}
\bar{X}_7	-	-	-	-	-	-	-

*P ≤ 0.05 **P ≤ 0.01 NS = not significant.

For instance, the relationship between yield and each of its other six studied components [days to heading (\bar{x}_1), No of spikes plant" (\bar{x}_2), No. of kernels/spike (\bar{x}_3), spike kernels weight (\bar{x}_4), 1000 kernels weight (\bar{x}_5) and harvest index (\bar{x}_7)] in this work were -0.53**, 0.41*, -0.72**, 0.67**, 0.76** and 0.68**, respectively (Table 7).

These results indicate that wheat breeders should select the earlier plants which have improved number of spikes, spike kernels weight and, in particular, heavier 1000 kernels weight to improve grain yield using both production index (\bar{x}) and the estimated stability parameters in this work. Where, Bansal & Sinha (1991b) stated that the stability in grain yield of *T. aestivum* (Landraces and improved wheat cultivars) under stress conditions was strongly depended on the stability in spikes either per unit area or per plant. In addition, our results were in agreement with those obtained by EI-Morshidy *et al.* (1998 and 2000) and Yan & Hunt (2001).

Moreover, our data emphasize that both mean performance of a genotype and its stability parameters should be taken together into consideration to recommend such new genotype to be used in varying environments. Whereas, previous studies illustrated that the promising genotypes were Ahgaf, Giza 163 and Giza 160, these genotypes which all showed to have at most higher average of performance than the grand mean and also acceptable stability parameters to its studied traits (EI-Morshidy *et al.*, 1998 and 2000). Although selection based upon yield *per se* should be the most efficient method for increasing the mean yield of a population (Wells & Kofoid, 1986). Parveen *et al.* (2010) noticed some cultivars as stable on the basis of overall mean yields and stability parameters viz., regression coefficients and minimum deviations from regression.

Thus, according to Eberhart & Russell (1966), the genotypes no. 13, 14 and 22 may be considered superior under abiotic stresses (heat and drought) because they showed high mean performance when compared with grand mean beside acceptable stability parameters to the studied traits under these conditions as follows:

No. of genotype	Item	Characters						
		Days to heading	No. of spikes/ m ²	No. of kernels/ spike	Spike kernels weight	1000 kernel weight	Grain yield/ m ²	Harvest index
13 (NGB10893)	Mean Stability	Equal to stable	Higher stable	Medium stable	Higher unstable	Higher stable	Higher stable	Medium stable
14 (Sids 1)	Mean Stability	Late unstable	Higher stable	Higher stable	Higher stable	Higher stable	Higher stable	Higher stable
22 (Giza 168)	Mean Stability	Early stable	Lower unstable	Higher stable	Higher stable	Higher stable	Higher stable	Higher stable

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In reality, the genotypes no. 13 (NGB10893), 14 (Sids 1) and 22 (Giza 168) are recommended and adapted to use in abiotic stresses (heat and drought) environments. The breeder should compromise the relationship between an average of performance of a genotype and its stability parameters.. Thus, the breeders are often requested to recommend the highest yielding genotypes irrespective of whether a genotype is stable over all traits or no.

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مقاييس الثبات لمقارنة بعض التراكيب الوراثية لمقمح الخبز تحت ظروف الجفاف والحرارة

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اجري هذا البحث لمقارنة ٢٢ تركيب وراثي محلي ومستورد لتحمل الحرارة والجفاف وذلك باستخدام تحليل التباين والثبات وكذلك معاملي الاختلاف والتقدير. تم ذلك باجراء تجربتين حقليتين بالمزرعة البحثية لكلية الزراعة جامعة سوهاج خلال موسمي ٢٠١٢/٢٠١١ و ٢٠١٣/٢٠١٢ حيث نمت النباتات في تصميم القطاعات المنشقة في ثلاث مكررات في كل تجربة في كل موسم وزرعت التراكيب الوراثية في القطع المنشقة في كل موسم في ميعادين مختلفين للزراعة (مبكر ومتأخر). وطبق في كل تجربة ثلاث معاملات ري (اجهاد مائي) في القطع الكاملة حيث كانت المعاملة الاولى (I₁) كنترول (للمقارنة) لا يوجد اجهاد مائي (١٠ ريات) طول موسم النمو خلال جميع مراحل نمو النبات، المعاملة الثانية (I₂) ٧ ريات حيث خفضت ٣ ريات (اجهاد مائي) عن الكنترول وذلك من بداية مرحلة التزهير حتى الحصاد ، المعاملة الثالثة (I₃) ٥ ريات فقط حيث خفضت ٥ ريات عن الكنترول وذلك من مرحلة الامتلاء حتي مرحلة النضج اللبني. ولذا شكلت ١٢ بيئة نتيجة استخدام ٢ سنة x ٢ ميعاد زراعة x ٣ معاملات ري لمقارنة تلك التراكيب الوراثية وكانت النتائج كالتالي :

معظم الاختلافات الموجودة في الاختلافات الكلية للصفات المدروسة كانت ترجع إلى تأثير البيئة والتي كانت ترجع إلى تأثير العوامل البيئية (السنوات ومواعيد الزراعة ومعاملات الري) وكذلك التفاعل البيئي (السنوات x معاملات الري). لذا كانت الاختلافات البيئية معنوية وامتدادت من ١٠,٦٢% (معامل الحصاد) إلى ٤٣,٩٥% (وزن حبوب السنبله) .

كانت هناك فروق احصائية بين التراكيب الوراثية في جميع الصفات المدروسة وكان مدى تأثير هذه الفروق في الاختلافات الكلية بين ٦,٨٢% (وزن الالف حبة) إلى ٥٠,٤٢% (عدد الحبوب / سنبله) .

كانت التفاعلات الوراثية x البيئية للصفات المدروسة معنوية جدا والتي قدرت مساهمتها في الاختلافات الكلية ب ٤٠,١٧% ، ٣٠,٧٨% ، ١٨,١٧% ، ٢٢,٠٢% ، ٢٥,٤٦% ، ١٨,٦١% و ٨٨,٤٦% وذلك للتزهير ، عدد السنابل / نبات ، عدد الحبوب / سنبله ، وزن حبوب السنبله ، وزن الالف حبة ، محصول الحبوب و معامل الحصاد على التوالي .

أكدت النتائج ضرورة استخدام كل من متوسط أداء التراكيب الوراثية ومقاييس الثبات الخاصة به معا للتوصية باستخدام اي تركيب وراثي في بيئات مختلفة. حيث امكن تحديد ٣ تراكيب وراثية عالية المحصول وأكثر ثباتا تحت ظروف الجفاف والحرارة وهذه التراكيب هي جيزة ١٦٨ ، سدس ١ ، NGB10893 والتي يمكن استخدامها في برامج التربية على نطاق واسع تحت هذه الظروف .