

## Evaluation of Wheat *Triticum aestivum* L. Genotypes with High Productivity in Al-Muthanna and Thi-Qar Provinces

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### Summary

A field experiment was conducted to evaluate the performance of different genotypes of soft wheat crop at two locations, the first location in the agricultural research station and trials of the College of Agriculture / Muthanna province and the second location in Al-Garraf district / province of Thi- Qar. Ten genotypes (IR885, IR899, IR1123 , IR981, IR1131, IR969, IR1069) are used and Abu Ghraib, Furat and IBA 99 as comparison cultivars) , Three replicates of each treatment were randomized using a randomized complete block design factorial experiments. The effect of genotypes, locations and the interactions between them has been studied in some growth and yield traits and its components. IR98 genotype record higher rate of seed yield kg / donum for both Al-Muthanna and Thi- Qar locations in which reached (1847.667 and 1889.000 kg / donum), respectively while the results showed that the superiority of Furat cultivar giving the highest rate of height plant, spike length and number of spikes / m<sup>2</sup> which reached ( 105.333 cm and 17,000 cm and 805.333 spike / m<sup>2</sup> ), respectively. As well as my agriculture disagreed with each other in most of the studied traits where the second location (Thi-Qar) gave the highest averages for plant height and spike length and number of the forest and the number of spikes / m<sup>2</sup> and weight of 100 tablets and dry weight / m<sup>2</sup> and holds the grain reached (96,400 cm and 14,800 cm and 8.600 and 615.000 spike / m<sup>2</sup> and 3.725 g and 245.700 g / m<sup>2</sup> 1889.000 kg / sq.m), respectively. With regard to the interactions between the genotypes and locations, Furat cultivar had superiority in Thi- Qar location, in which gave the highest rate of plant height and spike length which reached (113.333 cm and 20.000 cm), respectively, while the cultivar surpassed EBA 99 in Thi- Qar location, giving the highest rate of Leaf area reached (76.633 cm<sup>2</sup>) while the interactions between genotype IR1069 in Thi- Qar location, the highest average of 100 seed weight reached (5.020 g).

**Keywords:** Performance evaluation , genotypes , fine wheat, agricultural locations .

### Introduction

wheat Crop *Triticum aestivum* L. Is considered the most important strategic grain crops relied on to reduce the food gap, which has become the world's problem, and the wheat are grown in most parts of the world and constitute the basic material for the manufacture of bread. Recent statistical projections indicate that global production will reach 689.8 million tons in 2015 (13), and the reason for importance of this crop is being the main food of more than 40 countries in the world (nearly 35% of the world population), Due to its high content of protein and carbohydrates (12) as it is gave more than 25% needs of calories and protein (11) in addition to contain the amounts of fat and vitamins (B1 and B2) and some mineral salts (11).

The rectification a new genotypes and election the superior of them in yield and other traits consider important to creating new varieties that can be an alternative to the local varieties that currently cultivated or in the provision of genotypes can be used to transfer genes of higher yield and resistance to diseases and high protein to local varieties cultivated through the hybridization program (5) .

The genotype Affects in the plant behavior in terms of vegetative growth, Leaf area chlorophyll content, grain yield and its components , as both (8 and 15) found that difference of wheat bread genotypes in most vegetative growth traits and grain yield and its components, due to the nature of growth, capability of branching and total dry matter production, also the good crop and soil preparation practices have a role in increasing the yield, but this increasing did not reach the optimum level. it seems that the ideal solution to this problem is to develop new varieties more suitable for the environmental conditions , so to achieve the goal of increasing a grain yield the plant breeders needs to identify the most important characteristics of growth and the components of yield associated morphologically and genetically directly or indirectly with grain yield for use it as elected evidence. so as a simple correlation measures the connectivity relationships, the coefficient of the path is the best way to identify those relationships because it determines the effects of direct and indirect of the yield traits based on genetic relationships (14) .

Wheat varieties ( Maxibak and Sabrbek ) Still prevailed cultivated in Iraq since long period and characterized in sensitivity for lodging, the incidence of certain diseases , in addition to the lack of purity because of mechanical mixing and accumulated mutations, so now it is necessary to develop new varieties replace or grown along with these varieties , and because of the lack of available studies on the breeding of wheat and the lack of sources of heterogeneity genetic that can be used for breeding purposes , So the goal of

this research is to evaluate introduced genotypes of bread wheat different in the length of the growth and maturity from the International Center for maize and Wheat Improvement in Mexico (CIMMYT) with three comparison varieties that adopted in Iraqi agriculture in terms of growth and yield traits and components in order to determine the productive of the different varieties and election of superior varieties under conditions of the southern region of Iraq.

### Materials and methods

A field experiment was carried out during 2012-2013 in two locations. The first was agricultural research and experiments station belonging to agriculture college/Al-Muthanna university/3km north east of Samawa city center while the second location in Thi-Qar goverment Al-Gharraf district about 26 km north Nassiryah city. The experiment was laid out in a Random Complete Block Design (R.C.B.D) with three replications. The soil was prepared and the soil samples collected from (0-20) depth in order to analyzing it and limiting the physical and chemical characteristics as it showed in Table (1). A field divided into plots the area for each one was 6m<sup>2</sup> and the seeds were cultivated in lines with 6m long and the distance between them was (20 cm) in each experimental plot. The quantity of fertilizer that applied was 100 kg/ha one batch during cultivation while urea fertilizer (46%N) were applied in three batch, first batch was during cultivation, the second after tillers and the third during spikes formation stage (3). Ten Genotypes of soft wheat as in table (2) which illustrates names of Genotypes field symbols and cultivar sources. The studied traits were (plant hight (cm), leaf area (cm<sup>2</sup>), Length of spike (cm), Number of tellers /plant, Number of spikes/m<sup>2</sup>, Number of seeds/ spike, 100 seed weight (gm), Dry weight (gm/m<sup>2</sup>), Seed yield (kg/dounm).

**Table (1) physical and chemical characteristics of soil locations**

Traits Location	Texture	PH	EC des/m	Particles%			Minerals%		
				Sand	Laom	Clay	N	K	P
Al-Muthanna*	Salty clay	7.65	4.21	20	39	41	0.70	1.21	0.22
Thi-Qar**	Loauty clay	7.82	3.72	22	43	35	0.65	1.35	0.52

\*Data analyzed in Al-mothanna university/agriculture college/ soil & water Lab\*\* Data analyzed in Dhi-qar directorate / soil & water Lab

After data collection and tabulation Statistical analysis was performed according to the design above and tested using arithmetic averages less significant difference (LSD) at the average level of probability (0.05) (2).

**Table (2) Data of introduced Genotypes in study.**

Genotype	IR885	IR899	IR1123	IR981	IR1131	IR969	IR1069	Abu Ghraib	IPA99	FURAT
Code	W1	W2	W3	W4	W5	W6	W7	W8	W9	W10
Source	International Corn & Wheat improvement Centre in Mexico (CIMMYT)/ General body of researches							Certified cultivars from Agriculture Ministry		

### Results and discussion

#### The effect of genotypes, locations and interaction in the growth traits

##### Plant height (cm)

The data presented in table (3) indicated that genotypes significantly affected the plant height . Maximum plant height (105.333) cm was observed for W10 genotype, whereas minimum value (78.833) cm was observed for W7 genotype. The reason for that may be due to the genetic nature of genotype , The result is in agreement with(1,16). also the location of study were differed significantly between them in this trait in which Thi- Qar location, gave the highest rate was ( 96.400cm), while Muthanna location gave the lowest rate for this trait was ( 86.300cm) and the reason for this is the nature of the soil Table (1) and environmental conditions . As regards to the effect of interaction W10 genotype gave highest rate for this trait was (113.333) cm in Thi- Qar location .

##### Leaf area (cm 2)

The results of table (3) showed that the significant effect of genotypes on leaf area where W9 genotype given highest rate of leaf area ( 64.617 cm 2) while W6 genotype given lowest rate for this trait ( 39.917 cm 2) and the reason for this is the difference of genotypes among them in leaf area in addition to the viability among genotypes, this result is in agreement with the findings of the ( 7), which showed that in order to produce high yield of crop it should produce sufficient leaf area index to intercept the largest amount of light to produce the largest amount of dry matter , also the interaction among W9 genotype in Thi- Qar location recorded the highest

rate for this trait ( 76.633 cm<sup>2</sup>) and the reason for this is due to the interaction effect of environmental and genetic factors .

### Length of spike

The results of table (3) showed that the significant effect of genotypes on length of spike where W10 genotype given highest rate of tall spike ( 17.000 cm) while W5 genotype given lowest rate for this trait ( 12.167 cm),Where in spite of the variation in the rates of this trait among genotypes, this cannot indicate that the increase in the length of the spike is significant to increase the number of grains / spike and this result is in agreement with the findings of (6). Also Thi- Qar location was superior, giving higher rate for this trait was (14.800 cm), Regarding the interaction of Thi- Qar location and the W10 genotype has given highest rate for this trait was (20.000 cm).

### Number of tellers

The results of table (3) showed that the significant effect of genotypes on number of tellers where W1 genotype given highest rate of tall spike ( 11.333 teller) while W7 genotype given lowest rate for this trait ( 4.667 teller) and the reason for that is the formation of tellers (branches) depend on cultivar, applied fertilizers, cultivation date and environmental conditions, this result is in agreement with the findings of (4, 19). Also Thi-Qar location was superior to giving highest rate of this trait (8.600 teller) and the reason for that is the nature of the soil and its texture and minerals content.

**Table (3) Effect of Genotypes, locations and interaction on Growth traits for Al-Muthanna & Thi-Qar Locations.**

Trait	Cultivar	Location		Mean Cultivar	
		Al-Mothanna	Dhi-qar		
Plant height (cm)	W1	93.333	96.000	94.667	
	W2	88.333	85.000	86.667	
	W3	84.000	97.667	90.833	
	W4	88.667	92.333	90.500	
	W5	82.333	97.333	89.833	
	W6	82.333	100.667	91.500	
	W7	72.333	85.333	78.833	
	W8	94.000	111.667	102.833	
	W9	80.333	84.667	82.500	
	W10	97.333	113.333	105.333	
	Mean locations	86.300	96.400		
	L.S.D (0.05)	Cultivars = 2.74		Locations =1.22	Interactions =3.87
	Leaf Area (cm <sup>2</sup> )	W1	51.267	49.667	50.467
W2		48.900	45.133	47.017	
W3		65.233	51.600	58.417	
W4		53.833	54.167	54.000	
W5		52.313	51.933	52.123	
W6		39.000	40.833	39.917	
W7		47.167	48.133	47.650	
W8		53.200	53.200	53.200	
W9		52.600	76.633	64.617	
W10		42.867	57.000	49.933	
Mean locations		50.638	52.830		
L.S.D (0.05)		Cultivars =6.97		Locations N,S	Interaction =9.86
Spike Tall (cm)		W1	13.667	13.667	13.667
	W2	11.667	14.333	13.000	
	W3	16.333	15.000	15.667	
	W4	10.667	14.000	12.333	
	W5	11.000	13.333	12.167	
	W6	11.667	13.333	12.500	
	W7	12.000	15.000	13.500	
	W8	12.667	15.000	13.833	
	W9	12.333	14.333	13.333	
	W10	14.000	20.000	17.000	
	Mean locations	12.600	14.800		
	L.S.D (0.05)	Cultivars =1.01		Locations =0.45	Interactions =1.43
	Tellers No.	W1	10.000	12.667	11.333
W2		5.333	8.000	6.667	
W3		7.333	7.333	7.333	
W4		8.667	11.667	10.167	
W5		5.000	7.000	6.000	
W6		7.000	9.000	8.000	
W7		4.000	5.333	4.667	
W8		4.333	6.667	5.500	
W9		7.333	10.000	8.667	
W10		5.000	8.333	6.667	
Mean locations		6.400	8.600		

L.S.D (0.05)	Cultivars =1.27	Locations =0.57	Interactions =N.S.
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### The effect of genotypes and locations in the yield traits

#### Number of spikes/ m<sup>2</sup>

The results of table (4) showed that the significant effect of genotypes on number of spikes/m<sup>2</sup> where W10 genotype given maximum number of spikes (805.333 spike/m<sup>2</sup>) while W7 genotype given minimum number for this trait ( 447.167 spike/m<sup>2</sup>) this result is in agreement with the findings of (1, 10) also Thi- Qar location was superior in this trait giving maximum number reach (615.000 spike/m<sup>2</sup>).

#### Number of seeds/ spike

The results of table (4) showed that the significant effect of Number of seeds/ spike where W7 genotype given highest rate of seed (97.000 seed/spike) while W2 genotype given lowest rate for this trait ( 37.000 seed/spike) this result is in agreement with the findings of (1, 12).

#### 100 seed weight (gm)

The results of table (4) showed that the significant effect of 100 seed weight (gm) where W7 genotype given highest rate (4.918 gm) while W1 genotype given lowest rate for this trait ( 2.445 gm) may be due to the genetic nature of cultivar in which cultivars differ in Germplasm, this result is in agreement with the findings of (4, 1) also Thi- Qar location was superior in this trait giving highest rate reach (3.725 gm), Regarding the interaction between genotypes and locations W7 genotype gave highest rate for this trait reach (5.020 gm).

#### Dry weight (gm/m<sup>2</sup>)

The results of table (4) showed that the significant effect of Dry weight (gm) where W2 genotype given highest weight (335.833 gm) while W3 genotype given lowest weight for this trait ( 47.500 gm) also Thi- Qar location was superior to giving highest weight reached (245.700 gm).

#### Seed yield (kg/dounnm)

The results of table (4) showed that the significant effect of Seed yield (kg/dounnm) where W4 genotype given highest yield (1868.333 kg/dounnm) while W8 genotype given lowest yield for this trait reached (844.323 kg/dounnm) may be due to the genetic nature of cultivar in which cultivars differ in Germplasm in seed yield and this result in agreement with (1, 4, 18) also the results showed the significant effect of for locations in this traits in which Thi- Qar location gave highest yield (1244.159 kg/dounnm).

**Table (4) Effect of genotypes, locations and interaction on Yield and its compenents traits for Al-Mothanna & Thi-Qar Locations**

Traits	Cultivars	Locations		Mean Cultivars
		Al-Mothanna	Dhi-Qar	
Spikes No./m <sup>2</sup>	W1	497.000	514.667	505.833
	W2	662.667	685.333	674.000
	W3	606.667	635.333	621.000
	W4	641.667	646.000	643.833
	W5	513.333	544.667	529.000
	W6	661.667	685.667	673.667
	W7	433.333	461.000	447.167
	W8	531.000	581.667	556.333
	W9	558.333	567.333	562.833
	W10	782.333	828.333	805.333
	Mean Locations	588.800	615.000	
L.S.D (0.05)		Cultivars =21.03	Locations =9.40	Interactions = N.S.
Seed No./spike	W1	73.000	76.333	74.667
	W2	35.000	39.000	37.000
	W3	49.333	53.667	51.500
	W4	66.333	68.333	67.333
	W5	61.000	58.000	59.500
	W6	46.333	44.333	45.333
	W7	82.000	76.000	79.000
	W8	48.667	44.667	46.667
	W9	56.000	54.667	55.333
	W10	48.000	45.000	46.500
	Mean Locations	56.567	56.000	

<b>100 seed weight (gm)</b>	<b>L.S.D (0.05)</b>	<b>Cultivars = 3.98</b>	<b>Locations = N.S.</b>	<b>Interactions = N.S.</b>
	W1	2.410	2.480	2.445
	W2	4.433	4.457	4.445
	W3	4.030	4.073	4.052
	W4	4.343	4.283	4.313
	W5	4.417	4.083	4.250
	W6	3.200	3.273	3.237
	W7	4.817	5.020	4.918
	W8	3.183	3.343	3.263
	W9	2.767	2.827	2.797
	W10	2.950	3.413	3.182
	<b>Mean Locations</b>	<b>3.655</b>	<b>3.725</b>	
<b>Dry weight/m<sup>2</sup> (gm)</b>	<b>L.S.D (0.05)</b>	<b>Cultivars = 2.66</b>	<b>Locations = 1.19</b>	<b>Interactions = 3.76</b>
	W1	208.333	243.333	225.833
	W2	316.667	355.000	335.833
	W3	133.333	161.667	147.500
	W4	203.333	213.333	208.333
	W5	238.333	250.000	244.167
	W6	290.000	306.667	298.333
	W7	206.667	216.667	211.667
	W8	193.333	208.667	201.000
	W9	245.000	286.667	265.833
	W10	200.000	215.000	207.500
	<b>Mean Locations</b>	<b>223.500</b>	<b>245.700</b>	
<b>Seed yield (kg/donum)</b>	<b>L.S.D (0.05)</b>	<b>Cultivars =16.27</b>	<b>Locations = 7.28</b>	<b>Interactions = N.S.</b>
	W1	873.857	969.323	921.590
	W2	1027.150	1190.000	1108.575
	W3	1203.333	1387.667	1295.500
	W4	1847.667	1889.000	1868.333
	W5	1383.667	1288.333	1336.000
	W6	981.823	994.550	988.187
	W7	1711.667	1758.000	1734.833
	W8	820.320	868.327	844.323
	W9	863.110	876.057	869.583
	W10	990.067	1220.333	1105.200
	<b>Mean Locations</b>	<b>1170.266</b>	<b>1244.159</b>	
	<b>L.S.D (0.05)</b>	<b>Cultivars = 98.55</b>	<b>Locations = 44.07</b>	<b>Interactions = N.S.</b>

### Conclusions and Recommendations:

In conclusion W4 Genotype has given the highest rate of seed yield reached ( 1868.333 ) kg / donum , which shows the appropriate environmental conditions with the nature of genotype as a result led to a superiority in the quantity of yield and also Thi- Qar location was superior to giving highest rate of most of the studied traits ( plant height, length of Spike, number of tellers, number of spikes , 100 seed weight, dry weight, and seed yield ), while the interaction between the location of Thi- Qar and genotype W7 gave highest rate of 100 seed weight reached ( 5.020 g ), while giving genotype W9 highest rate of leaf area was ( 76.633 cm<sup>2</sup>) and W10 genotype gave highest rate of plant height and spike length was ( 113.333 cm and 20.000 cm ), respectively. So we recommend cultivation W4 genetic due to the superiority of this genotype by giving the highest rate seed yield and the possibility of distributing the cultivation of these genotypes at the Thi- Qar location, because of the superiority of this location most of the studied traits and replant these genotypes at other locations within the study area to reach the stability of genetic and submitted for approval and registration.

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