

STABILITY ANALYSIS FOR GRAIN YIELD AND IT'S CONTRIBUTING TRAITS IN BREAD WHEAT (*TRITICUM AESTIVUM* L.)

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Abstract: The present investigation carried out by to assess the phenotypic stability for grain yield and its contributing traits under three varied environmental conditions using 10 diverse wheat genotypes and 45 F₁S. The mean sum of squares due to genotypes and environments were found to be significant for all the characters, indicating differential effect of environment on the genotypes. G x E interaction was found significant for all the characters except spike length and biological yield per plant. Similarly the pooled analysis of variance showed that the mean squares due to E + (G x E) interaction was partitioned into G x E (linear) and pooled deviation (nonlinear components). Mean sum of squares due to G x E (linear) component and pooled deviation (nonlinear component) was significant for all the characters. The genotypes Raj 3765, Raj 3777, Raj 4238 x WH 1021, Raj 3077 x Raj 3777 had higher yield and were suitable for variable environmental conditions. These genotypes could be utilized as a donor in routine breeding programme to improve grain yield and its contributing traits in bread wheat.

Keywords: Bread wheat, Stability, Grain yield, Genotype x environment interactions

INTRODUCTION

Bread wheat (*Triticum aestivum* L.) has covered the most cultivation area and its product is the most widely used products in the world and of primary importance for human nutrition. It is a mandatory food crop and provides as a staple food in different parts of the world. It is a major food grain in India after rice, therefore the primary objective of wheat specialists is to increase the grain yield. Wheat genotypes identification with varied adaptation to dissimilar conditions, multi-location, results in the empirical identification of superior varieties. The adaptability of a variety over diverse environments is usually tested by the degree of its interaction with different environments under which it is grown. A variety or genotype is considered to be more adaptive or stable one, if it has a high mean yield but low degree of fluctuation in yielding ability, when grown over diverse environments. According to Eberhart and Russell (1966) a stable variety should have around unit regression coefficient over environments ($b_i \approx 1$) and minimum deviation from the regression ($S^2_{di} = 0$) in addition to higher seed yield than population mean. The objectives of this study were to identify bread wheat genotypes that have both high mean yield and stable yield performance across different environments and to study the similarities, relationships and dissimilarities among yield - stability statistics carried out to quantify GE interaction effects on yield and to determine stable entries within the genotypic pool used in this study.

MATERIALS AND METHODS

Crosses among the ten genotypes were made in diallel fashion excluding reciprocals, during *Rabi* season 2016-17. A number of plants were randomly selected in a parent crossed with a number of randomly selected plants from other parent. In *Rabi* 2017-18 ten genotypes along with their 45 F₁'s progenies were evaluated in three environments created three different dates of sowing 15 Nov. (timely sown), 1 Dec. (late sown) and 15 Dec. (very late sown) with 3 replications at Research Farm, College of Agriculture, Bikaner. Row length was kept 4 meter and plant geometry (Row to row and plant to plant distance) was 22.5 cm and 10 cm, respectively. Stability parameters for grain yield and its contributing traits were worked out as per procedure suggested by Eberhart and Russell (1966).

RESULTS AND DISCUSSION

The genotype × environment (GE) interaction is a major challenge to plant breeders. Many stability parameters for genotypes grown in different environments were developed for this purpose and each has its advantages and limitations. In various methods, GE interactions are used to characterize the response of genotypes to changing environments along with mean grain yields. Accordingly, genotypes with a minimal variance for yield across environments were considered stable (Mohammadi, 2012). Improving yield and yield stability is the principal objective of wheat breeding programs (Ram *et al.*, 2007). To study of wheat genotypes with wide

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or specific adaptation to different environments, trials are grown at three different date of sowing. These have resulted to empirical identification of superior cultivars, some of which have been released in several countries (Basford *et al.*, 2004). The environments involve a wide range of photoperiods and temperatures which could cause large genotype (G) \times environment (E) interactions (GEI), specifically in the semi-arid regions. Large crossover-type GEI, especially among high yielding lines gives incorrect suggestions to farmers across the all test environments. Quantification of GEI and comprehension its physiological bases are needed to breed effectively for superior environments (Thomason and Phillips, 2006). Most yield trials are used only to determine which cultivars give the highest average grain yield and therefore valuable recommendation for planting by farmers. Understanding the structure and nature of GEI is important in plant breeding programs because a significant GEI can seriously impair efforts in selecting superior genotypes relative to new crop introductions and cultivar improvement programs (Shafii and Price, 1998). The detect of GEI in trials has led to the development of procedures that are generically called stability analyses. The available numerous stability statistics to breeders and to the production agronomist provide different strategies and approaches of dealing with GEI. Stability is an important concept for plant breeders interested in analysing GEI data (Denis *et al.*, 1996; Ayed *et al.*, 2016).

Analysis of variance for stability for grain yield and its contributing characters in bread wheat were shown in Table 1. The pooled analysis of variance across the environments revealed significant differences among genotypes and environments for grain yield, indicating differential effect of environment on the genotypes. The mean sum of squares due to G \times E interaction was found significant for all the characters except spike length and biological yield per plant. Similar findings were also reported by Sharma *et al.* (2003), Arya *et al.* (2004), Jeena *et al.* (2005), Banerjee *et al.* (2006), Singh and Chaudhary (2007), Arain *et al.* (2011), Mohammadi *et al.* (2012), Saleem *et al.* (2015) and Bavandpori *et al.* (2018). The mean sum of squares due to E + (G \times E) was also found significantly for all the characters. Similarly the pooled analysis of variance showed that the mean squares due to E + (G \times E) interaction was partitioned into G \times E (linear) and pooled deviation (nonlinear components). Mean sum of squares due to G \times E (linear) component and pooled deviation (nonlinear component) were significant for all the characters which indicated that prediction across the environments was possible for all characters. Similar results were also reported by Ashraf *et al.* (2001), Madariya *et al.* (2001), Khumkar *et al.* (2001), Singh *et al.* (2002), Sharma *et al.* (2003), Arya *et al.* (2004), Najeeb *et al.* (2004),

Jeena *et al.* (2005), Singh and Chaudhary (2007) and Patel *et al.* (2014).

Among the crosses, three best stable crosses for various environments conditions were presented in Table 2. The joint consideration of mean performance of genotypes across the environments and stability parameters revealed that out of 45 genotypes, genotypes namely Raj 1482 \times Raj 3777, Raj 4238 \times WH 1021 and Raj 3765 \times DBW 90 were suitable for harsh environment (mean < population mean, $b_i < 1$ and $S^2d_i = 0$) while the genotypes Raj 3077 \times DBW 90, Raj 1482 \times DBW 90 and Raj 3765 \times PBW 550 were most suitable for early flowering in average environment (mean < population mean, $b_i = 1$ and $S^2d_i = 0$). The present finding supported the results obtained by Badhe *et al.* (1998) and Meena *et al.* (2014). For days to maturity the cross combinations Raj 4079 \times Raj 3765 and Raj 4079 \times Raj 3777 were predictable for better environment. Among 45 genotypes, genotypes namely Raj 4079 \times Raj 1482, Raj 3077 \times Raj 3777 and Raj 4079 \times DBW 90 were suitable for poor environments (mean < population mean, $b_i < 1$ and $S^2d_i = 0$) while the crosses Raj 3777 \times HD 3086, PBW 550 \times HD 3086 and Raj 3077 \times WH 1021 were most stable and desirable for early maturity. The present finding supported the results obtained by Badhe *et al.* (1998) and Meena *et al.* (2014). Late season heat stress, often occurring during the last phases of spring wheat development, (booting, heading, anthesis and grain filling period) is considered as one of the major environmental limiting factors that drastically reduces grain number per spike, grain weight and final grain yield throughout most of the bread wheat growing areas in the region and other warm and dry regions [Modhej *et al.*, (2008) and Mohammadi *et al.* (2012)]. The crosses WH 1021 \times PBW 550, Raj 1482 \times Raj 3777 and Raj 3765 \times HD 3086 were suitable for rich environment conditions while the crosses namely Raj 1482 \times WH 1021 and Raj 4238 \times WH 1021 were suitable for varying environment conditions. Taller plants are more likely to lodge quite often. Short stature in wheat is preferred. In the present study, cross combinations Raj 4079 \times HD 3086, Raj 3777 \times WH 1021 and Raj 3765 \times DBW 90 were considered for poor environments (mean < population mean, $b_i < 1$ and $S^2d_i = 0$). While the crosses namely Raj 3765 \times PBW 550 and Raj 3777 \times PBW 550 (mean < population mean, $b_i = 1$ and $S^2d_i = 0$) were considered stable for average environments. Significant GE interaction for plant height was also reported by Sakin *et al.* (2011). Number of effective tillers per plant is important character from the point of view of straw and grain yield, respectively. In the current study, crosses Raj 3765 \times Raj 3077, Raj 4079 \times Raj 3077 and Raj 1482 \times WH 1021 were considered for poor environments (mean > population mean, $b_i < 1$ and $S^2d_i = 0$). While the crosses namely Raj 4238 \times WH 1021, DBW 90 \times PBW 550 and Raj 4079 \times Raj 3765 were best suited for average

environments. These results are also similar with the earlier findings of Jaydeep *et al.* (2006). For test weight, the crosses WH 1021 x PBW 550, DBW 90 x PBW 550 and Raj 3777 x HD 3086 were recognized for late sown while the crosses Raj 4238 x WH 1021, Raj 4079 x PBW 550 and Raj 4238 x DBW 90 were responsive for timely sown conditions. The crosses like Raj 3765 x Raj 3077, Raj 3077 x Raj 1482 and Raj 4079 x Raj 3077 were found desirable for wide range of environments.

Meena *et al.* (2014) and Kumar *et al.* (2014) also identified some wheat genotypes which give stable performance for 1000-grains weight across the environment. For flag leaf area, number of grains per spike and harvest index G x E interaction was significant. Similar findings were also reported by Banerjee *et al.* (2006), Singh and Chaudhary (2007), Mohammadi *et al.* (2012), Singh *et al.* (2017) and Bavandpori *et al.* (2018).

Table 1. Analysis of variance for stability for grain yield and its contributing characters in bread wheat

| Source | df | Days to Heading | Days to Maturity | Grain filling period | Plant Height | Flag leaf area (cm ²) | Number of effective tillers per plant | Spike Length (cm) | Number of grains per spike | 1000-Seed weight (g) | Biological yield per plant (g) | Harvest Index (%) | Grain yield per plant (g) |
|-------------------|------|-----------------|------------------|----------------------|--------------|-----------------------------------|---------------------------------------|-------------------|----------------------------|----------------------|--------------------------------|-------------------|---------------------------|
| Repl./Env. | 6 | 1.02 | 0.85 | 2.16 | 4.56 | 0.83 | 0.26 | 1.3 | 2.91 | 4.32 | 13.79* | 1.34 | 1.37 |
| Genotypes | 54 | 15.09** | 19.41** | 29.99** | 44.38** | 10.61** | 4.11** | 5.04** | 69.23** | 32.5** | 124.87** | 62.64** | 40.94** |
| Environment | 2 | 2243.58** | 8626.69** | 2071.61** | 3758.03** | 896.34** | 40.07** | 74.17** | 1561.85** | 1520.68** | 1814.14** | 798.39** | 727.13** |
| G x E | 108 | 3.03** | 6.91** | 9.67** | 7.54** | 4.46** | 0.17** | 0.3 | 3.28** | 3.22** | 2.06 | 7.69** | 1.7** |
| Env. + (G x E) | 110 | 43.77** | 163.64** | 47.16** | 75.73** | 20.68** | 0.9** | 1.64** | 31.62** | 30.81** | 35.01** | 22.07** | 14.89** |
| Env. (linear) | 1 | 4487.17** | 17253.39** | 4143.21** | 7516.06** | 1792.68** | 80.14** | 148.34** | 3123.7** | 3041.35** | 3628.27** | 1596.79** | 1454.25** |
| G x E (linear) | 54 | 4.49** | 10.4** | 14.72** | 12.88** | 6.41** | 0.23** | 0.27 | 4.64** | 4.23** | 2.93** | 12.34** | 2.64** |
| Pooled deviations | 55 | 1.55** | 3.36** | 4.53** | 2.17** | 2.46** | 0.11* | 0.32* | 1.89** | 2.17** | 1.16** | 2.99** | 0.76** |
| Pooled error | 324 | 0.71 | 0.76 | 1.28 | 1.32 | 1.47 | 0.11 | 0.36 | 0.95 | 2.04 | 4 | 2 | 0.47 |
| Total | 1136 | 11.54 | 39.03 | 13.95 | 21.98 | 5.84 | 0.66 | 0.93 | 14.79 | 10.54 | 21.71 | 11.91 | 7.89 |

Table 2. Best stable crosses for different environments (Ranked on the basis of means)

| Characters | Better environment | Poor environment | Average environment |
|------------------------------------|---|---|--|
| Days to heading | 1. Raj 4079 x Raj 3765 2. Raj 4079 x Raj 3777 | 1. Raj 1482 x Raj 3777 2. Raj 4238 x WH 1021 3. Raj 3765 x DBW 90 | 1. Raj 3077 x DBW 90 2. Raj 1482 x DBW 90 3. Raj 3765 x PBW 550 |
| Days to maturity | 1. Raj 3777 x HD 3086 2. PBW 550 x HD 3086 3. Raj 3077 x WH 1021 | 1. Raj 4079 x Raj 1482 2. Raj 3077 x Raj 3777 3. Raj 4079 x DBW 90 | 1. DBW 90 x PBW 550 2. Raj 3077 x Raj 1482 3. Raj 3765 x Raj 3077 |
| Grain filling period | 1. WH 1021 x PBW 550 2. Raj 1482 x Raj 3777 3. Raj 3765 x HD 3086 | 1. Raj 4079 x Raj 3765 2. Raj 3765 x DBW 90 3. Raj 3765 x PBW 550 | 1. Raj 1482 x WH 1021 2. Raj 4238 x WH 1021 |
| Plant height | 1. Raj 3077 x WH 1021 2. Raj 1482 x WH 1021 3. PBW 550 x HD 3086 | 1. Raj 4079 x HD 3086 2. Raj 3777 x WH 1021 3. Raj 3765 x DBW 90 | 1. Raj 3765 x PBW 550 2. Raj 3777 x PBW 550 3. |
| Flag leaf area | 1. Raj 4079 x WH 1021 2. Raj 3765 x Raj 3777 3. DBW 90 x HD 3086 | 1. Raj 4238 x DBW 90 2. Raj 3777 x PBW 550 3. Raj 3765 x PBW 550 | - |
| No. Of effective tillers per plant | 1. WH 1021 x PBW 550 2. Raj 3765 x HD 3086 3. Raj 3777 x HD 3086 | 1. Raj 3765 x Raj 3077 2. Raj 4079 x Raj 3077 3. Raj 1482 x WH 1021 | 1. Raj 4238 x WH 1021 2. DBW 90 x PBW 550 3. Raj 4079 x Raj 3765 |
| No. of grains per spike | 1. WH 1021 x PBW 550 2. Raj 3765 x HD 3086 3. Raj 3777 x HD 3086 | 1. Raj 3765 x Raj 3077 2. DBW 90 x PBW 550 3. Raj 3077 x Raj 3777 | 1. Raj 4238 x WH 1021 2. Raj 3077 x Raj 1482 3. WH 1021 x DBW 90 |
| 1000-seed weight | 1. Raj 4238 x WH 1021 2. Raj 4079 x PBW 550 3. Raj 4238 x DBW 90 | 1. WH 1021 x PBW 550 2. DBW 90 x PBW 550 3. Raj 3777 x HD 3086 | 1. Raj 3765 x Raj 3077 2. Raj 3077 x Raj 1482 3. Raj 4079 x Raj 3077 |
| Harvest index | 1. DBW 90 x PBW 550 2. Raj 3777 x HD 3086 3. Raj 1482 x PBW 550 | 1. WH 1021 x PBW 550 2. Raj 3765 x Raj 3077 3. Raj 3765 x DBW 90 | 1. Raj 3077 x Raj 3777 |
| Grain yield per plant | 1. Raj 3765 x Raj 3077 2. WH 1021 x PBW 550 3. Raj 3765 x HD 3086 | 1. Raj 3765 x DBW 90 2. Raj 4238 x Raj 4079 3. Raj 4079 x PBW 550 | 1. Raj 4238 x WH 1021 2. Raj 3077 x Raj 3777 |

Better environment (Below average stability): mean > μ , bi > 1 and S²di = 0

Poor environment (Above average stability): mean > μ , bi < 1 and S²di = 0

Average environment (Average stability): mean > μ , bi = 1 and S²di = 0

Grain yield is a complex character and the analysis of individual yield component can lead to simplification

in explaining the stability for grain yield. The joint consideration of mean performance of genotypes

across the environments and stability parameters revealed that out of 45 genotypes, genotypes namely Raj 3765 x DBW 90, Raj 4238 x Raj 4079 and Raj 4079 x PBW 550 were suitable for harsh environment (mean > population mean, $b_i < 1$ and $S^2d_i = 0$) while the genotypes Raj 4238 x WH 1021 and Raj 3077 x Raj 3777 was most suitable for high grain yield in all types of environment (mean > population mean, $b_i = 1$ and $S^2d_i = 0$). Similar results for harsh environments were reported by Hassaan (2003), Rane *et al.* (2007), Bhoite *et al.* (2011), Arain *et al.* (2011) and Maeng (2019). For varying environments conditions similar findings were obtained by Ranjana and Kumar (2013), Patel *et al.* (2014), Singh *et al.* (2017) and Bavandpori *et al.* (2018). The cross combinations Raj 3765 x Raj 3077, WH 1021 x PBW 550 and Raj 3765 x HD 3086 were favourable for better environment. The present results were also confirmed by Mondal and Khajuria (2002), Hassaan (2003), Arya *et al.* (2004), Mishra (2007) and Bhoite *et al.* (2011).

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