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# Genetic Studies in Bread Wheat Under irrigated and Drought Stress Conditions 

Ghulam Mahboob Subhani ${ }^{1}$ and Muhammad Aslam Chowdhry ${ }^{2}$<br>${ }^{1}$ Wheat Research Institute, AARI, Faisalabad, Pakistan<br>${ }^{2}$ Department of Plant Breeding and Genetics, University of Agriculture, Faisalabad, Pakistan


#### Abstract

A $6 \times 6$ diallel cross consisting of six wheat varieties/lines namely Pak 81, LU26S, Inqlab 91, Rohtas 90, 4072 and 4943 was analyzed to determine the nature of genetic mechanisms controlling some Morpho- physiological traits, like stomatal frequency, leaf venation, flag leaf area, specific flag leaf weight, days to heading, tillers per plant, plant height, spike length, grains per spike, 1000-grain weight, biomass per plant and grain yield per plant under irrigated and drought stress conditions. Adequacy test satisfied the simple additive dominance model. High estimates of narrow sense heritability were found for stomata! frequency, specific flag leaf weight, days to heading, plant height, spike length and 1000-grain weight, also observed lower average degree of dominance under both environments. In the present study additive genetic effects were found to be more important than nonadditive effects with the exception of yield per plant which appeared to be more closely linked to nonadditive genetic effects. Since the yield components like grains per spike, 1000-grain weight, tillers per plant were found to be significantly influenced by additive gene action, further progress in genetic improvement of the yielding ability of wheat varieties may be attempted through such yield-related characters, with much better prospects of fixing desirable genes in a single homozygote.


Key words: Diallel, gene action, combining ability, drought stress, morpho-physiological traits, bread wheat

## Introduction

Cereal breeders are largely concerned with gaining information regarding the genetic systems controlling morpho-physiological traits using statistical analysis techniques which enable them to test for epistasis and to obtain precise and unbiased estimates of the additive and dominance components of genetic variation. However, some earlier researchers (Lonts and Lone, 1984; Alam et al., 1991; lqbal et al., 1991; Chowdhry et al., 1992) reported the over dominance type of gene action for tillers per plant, spike length, flag leaf area, grains per spike, 1000-grain weight and grain yield per plant. While additive type of gene action along with partial dominance were observed for plant height, spike length, grains per spike, 1000 -grain weight and grain yield per plant by Alam et al. (1991), Iqbal et al. (1991) and Chowdhry et al. (1992). Whereas Prodanovic (1993) showed that tiller number, grain number per spike, 1000-grain weight and grain yield were conditioned mainly by dominance gene effects. Over dominance was found for some traits, such as grain number per spike and grain yield. High narrow sense heritability estimates were noted for spike length (94.7\%) and plant height ( $82.26 \%$ ).
General combining ability mean squares were higher than the specific combining ability mean squares for plant height, spike length, tillers per plant, grains per spike, 1000-grain weight, days to heading and grain yield per plant indicating that these characters were controlled by additive type of gene action (Khan and Bajwa, 1990; Li et al., 1991; Chaudhry et al., 1992, 1994). Whereas specific combining ability mean squares for tiller per plant and grain yield per plant were greater than those of their general combining ability mean squares suggesting that these traits were mainly controlled by non-additive type of gene action (Li et al., 1991). However, Chaudhry et al. (1994) reported that general combining ability mean squares were highly significant for all traits. Specific combining ability mean squares were non significant for days to heading, flag leaf area, tillers per plant and grain yield per plant, significant for spike length and highly
significant for plant height and 1000-grain weight.
The present study was under taken to determine the genetic systems affecting morpho-physiological traits and also to assess the relative performance of some bread wheat (Triticum aestivum L. em. Thell.) varieties for general and specific combining ability under irrigated and drought stress conditions.

## Materials and Methods

Study was conducted at University of Agriculture, Faisalabad, during the year 1994-96. Six varieties/lines of bread wheat viz., Pak 81, LU26S, Inglab 91, Rohtas 90, 4072 and 4943 were crossed in a diallel fashion. The thirty $F_{1}$ 's including reciprocals and their parents were space planted in randomized complete block design with three replications. A single row of 3.75 meter served as an experimental plot. Two seeds per hill were sown with the help of a dibble and later thinned to one seedling per site with a distance of 15 centimeters within rows and 30 centimeters between rows.
For two sets of experiments, one under regular irrigation and the other under non-irrigation (drought stress), the fields were irrigated for seed bed preparation. After planting of experimental population, four canal irrigations were applied to normal experiment during the active growing period. Whereas the other experiment entirely depends on natural precipitation and no surface irrigation was applied to drought experiment for maintaining moisture stress conditions. Normal agronomic practices like fertilizer application and weed control were applied to both experiments.
Measurements were made on only competitive plants under both environments for morpho-physiological traits like stomatal frequency, leaf venation, flag leaf area ( $\mathrm{cm}^{2}$ ), specific flag leaf weight $\left(\mathrm{mg} / \mathrm{cm}^{2}\right)$, days to heading, tillers per plant, plant height (cm), spike length (cm), grains per spike, 1000-grain weight ( g ), biomass per plant ( g ) and grain yield per plant (g). The data were subjected to analysis of variance for all the characters for individual

## Subhani and Chowdhry: Diallel, gene action, combining ability, drought stress

environments (irrigated and drought stress conditions) according to the method of Steel and Torrie (1980). The diallel cross method developed by Hayman (1954) and applied by Mather and Jinks (1982) was used for genetic analysis of the data. Further analysis for combining ability effects was performed by using Griffing (1956) Method I, Model II.

## Results and Discussion

Gene action: Highly significant mean squares among parents and crosses were obtained for all traits under irrigated as well as drought stress conditions. This confirmed the presence of genetic variability in the material grown under both environments and suggested that detailed analysis of gene action was warranted. The analysis of variance for the diallel cross (Table 1) shows that the mean squares of component, a, were highly significant for all the traits except biomass per plant and grain yield per plant under irrigated conditions. But under drought stress conditions mean squares of component, a, were significant for all traits except tillers per plant. This indicates that there was a wide variation caused by the additive genetic effects in all these traits under both environments. The significant mean squares of item, $b_{1}$ indicated the directional dominance for the expression of stomatal frequency, flag leaf area, spike length and 1000-grain weight under irrigated conditions whereas for tillers per plant, plant height, spike length, grains per spike and 1000-grain weight under drought stress conditions. The mean squares due to item, $b_{2}$, significant for flag leaf area, days to heading, plant height, grains per spike and 1000-grain weight indicating asymmetrical gene distribution among the parents under irrigated conditions. Similarly, asymmetrical gene distribution was observed for days to heading, biomass per plant and grain yield per plant under drought stress conditions, The significant mean squares due to item, $b_{3}$, for days to heading, tillers per plant, grains per spike, 1000-grain weight, biomass per plant and grain yield per plant suggested the importance of specific gene effects controlling these traits under both conditions. Different levels of dominance were observed for most of the traits under drought stress conditions.

Tests of assumption for the additive dominance model: The regression analysis for stomatal frequency, leaf venation, flag leaf area, specific flag leaf weight, days to heading, plant height, spike length, and 1000-grain weight under irrigated conditions and for leaf venation, flag leaf area, days to heading, plant height, spike length, grains per spike, 1000-grain weight and biomass per plant under drought stress conditions confirmed the validity of the additive dominance model.
The regression and array variance analysis depicted the partial adequacy of the model for tiller per plant, grains per spike, biomass per plant and grain yield per plant under drought stress conditions and tillers per plant under irrigated conditions. Partial failure of the assumptions indicated a more complex genetic system than that described by theoretical model (Hayman, 1954). However, Hayman (1954) stated that it is possible to make estimates of the population parameters and genetic components of such traits. Therefore, Wilson et al. (1978) computed the genetic components for partially adequate traits. Still it must be realized that such estimates were less
reliable than they would have been had all the assumptions been satisfied.

Genetic components: fkifi estimated components of genetic variance, $D, H_{1}, H_{2}, h^{2}, F$ and some parameters derived from these estimates are presented in Table 2. The narrow sense heritability for stomatal frequency, specific flag leaf weight, days to heading, plant height, spike length, and 1000-grain weight were higher than that of leaf venation, flag leaf area, tillers per plant, grains per spike, biomass per plant and grain yield per plant. These results were confirmed by the findings of Prodanovic (1993) who reported the high narrow sense heritability for plant height and spike length. These traits with higher narrow sense heritability also showed lower average degree of dominance ( $\left.\left(\mathrm{H}_{1} / \mathrm{D}\right) 1 / 2\right)$, ranging from 0.15 to 0.84 than the other traits. Under drought stress conditions, narrow sense heritability for flag leaf area, specific flag leaf weight, days to heading, plant height, spike length and 1000-grain weight were higher than other traits. The traits with higher narrow sense heritability also showed lower average degree of dominance ranging from 0.24 to 0.35 .
The $D$ and $H_{1}$ components were significant in irrigated conditions revealing that both additive and non-additive type of gene action were involved in the inheritance of leaf venation, flag leaf area, days to heading, spike length and grains per spike. The significant estimates of $D$ indicated that there were additive genetic differences among the parents for stomatal frequency, plant height. The $\mathrm{H}_{1}$ was positive and significant for tillers per plant, biomass per plant and grain yield per plant under irrigated conditions revealed that only non-additive type of gene effects were involved for the expression of these traits. Positive and significant values of $F$ indicated that dominant genes (alleles) were important for the expression of stomata! frequency, flag leaf area, days to heading, spike length, grains per spike and 1000-grain weight. The significant value of $E$ showed important share of environmental effects in the expression of all traits except tillers per plant and 1000-grain weight under irrigated conditions. The significant value of $h^{2}$ revealed a substantial contribution of dominant genes in controlling stomatal frequency, flag leaf area, days to heading, spike length, grains per spike and 1000-grain weight due probably to the loci marked by heterogeneity.
The significant and positive values of $D$ revealed that additive type of gene effects were important for leaf venation, flag, leaf area, specific flag leaf weight, days to heading and plant height whereas significant $H_{1}$ revealed the non-additive effects for tillers per plant under drought stress conditions (Table 2). The significant values of $D$ and $H_{1}$ exhibited that both additive and non-additive effects were involved for the inheritance of grains per spike, biomass per plant and grain yield per plant. The negative and significant values of $F$ showed the presence of recessive alleles for stomatal frequency, flag leaf area and days to heading. The significant values of $h^{2}$ revealed a substantial contribution of dominant genes in controlling flag leaf area, days to heading, tillers per plant, plant height, spike length, grains per spike and 1000 grain weight. Environment played an important role for the expression of all traits under drought stress conditions (Table 2). High narrow sense heritability estimates were

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observed for specific flag leaf weight, days to heading, plant height, spike length and 1000-grain weight under both environments. These traits also showed the partial dominance type of gene action under both environments (lqbal et al., 1991; Chowdhry et al., 1992; Khan et al., 1992). Flag leaf area, tillers per plant, grains per spike, biomass per plant and grain yield per plant were controlled by over dominance. Similar finding were reported by Lonts and Lone (1984), Alam et al. (1991), lqbal et al. (1991) and Prodanovic (1993) who also reported the over dominance for these traits.

Combining ability analysis: The general combining ability (gee) mean squares were significant ( $p<0.05$ ) for all traits except flag leaf area, grains per spike, biomass per plant and grain yield per plant (Table 3). Whereas specific combining ability (sca) mean squares was significant for all traits except stomata! frequency and specific flag leaf weight. The mean squares due to general combining ability were higher than the mean squares for specific combining ability for all traits, indicating the prevalence of additive gene action for these characters with the exception of grain yield per plant under irrigated conditions where non-additive effects appeared more important. Similar results were also reported by Khan and Bajwa (1990), Li et al. (1991) and Chaudhry et al. (1992, 1994). General combining ability variance was higher than specific combining ability variance for stomatal frequency, specific flag leaf weight, days to heading, plant height, spike length, and grains per spike under irrigated conditions (Table 3). The separation of general combining ability effects by the standard error ( g , ) are given in Table 4, Inglab 91 had consistently the highest general combining ability effects for specific flag leaf weight, spike length, biomass per plant and grain yield per plant. Variety LU26S had also highest values for 1000-grain weight and specific flag leaf weight. Whereas Rohtas 90 showed maximum general combining ability for tillers per plant and grains per spike under irrigated conditions (Table 4). These varieties were good general combiner for those traits which show maximum general combining ability effects. Promising parents and sca effects of the crosses for grain yield and other morpho-physiological traits under irrigated conditions are presented in Table 4. Crosses LU26S $\times$ Roh 90 and Pak81 $\times$ Inq; 91 had high and significant sca effects for grain yield per plant, 1000-grain weight, spike length, tillers per plant and days to heading under irrigated conditions. Although the cross $4072 \times 4943$ with highest sca effects for grain yield per plant and high grain yield per plant also had high sca effects for flag leaf area. Most of the crosses with high sca had at least one high sca parent. However, some of the crosses with high sca had one or both parent with average gca (Singh and Chatrath, 1997).
Significant general combining ability mean squares (Table 3) were obtained for all characters revealed a greater involvement of additive effects in their inheritance except in the case of tillers per plant, grains per spike that produced insignificant results. Khan and Bajwa (1990), Li et al. (1991) and Chaudhry et al. $(1992,1994)$ reported similar results, as were found in the present study under drought stress conditions.
General combining ability variance was higher than specific combining ability variance for all the traits except tillers per plant, grains per spike, biomass per plant and grain yield
per plant (Table 3), indicating the additive gene action for these traits under drought stress conditions. All varieties/genotypes had high mean values for these traits, indicating that selection among progeny should produce good responses. In case of drought stress conditions variety Pak 81 was good general combiner only for plant height (Table 3). Whereas variety LU26S was good general combiner for 1000-grain weight and grain yield per plant. Variety Inqlab 91 was good general combiner for spike length, grains per spike and biomass per plant. Variety Rohtas 90 was only good combiner for tillers per plant. Genotype 4072 was good general combiner for flag leaf area, specific flag leaf weight and days to heading. In contrast to irrigated conditions genotype 4943 was good general combiner for stomatal frequency under drought stress conditions. Crosses Pak $81 \times$ LU26S, Inq $91 \times$ Roh 90 and $4072 \times 4943$ had high sca effects for grain yield per plant, biomass per plant, grains per spike, spike length and tillers per plant under drought stress conditions (Table 4). Most of the crosses with high sca effects for grain yield and other morpho-physiological traits had at least one high gca parent. However, some of the crosses with high sca had one or both parent with average gca (Singh and Chatrath, 1997).
It is obvious from the present results that the cross Pak. $81 \times$ LU26S showed specific combination for leave venation, spike per length, grains per spike, biomass per plant and grain yield per plant under drought stress conditions. Whereas under irrigated conditions Pak. $81 \times$ Inqlab 91, LU265 $\times$ Rohtas 90 and LU26S $\times 4072$ were specific crosses for most of the traits. The cross Inglab $91 \times 4072$ had maximum specific combining ability effects for 1000 -grain weight under both environments.
The best performing cross may be produced by crossing the two parents showing the highest general combining ability. For example under irrigated conditions, LU26S and Inglab 91 had the lowest general combining ability estimates for days to heading and plant height and highest estimates for spike length, and grain yield. Their cross is expected to produce a population with less days to heading, dwarf plants and with improved grain yieid. Populations that reveal high general combining ability estimates are good candidates to be used as parents in a population improvement programme (Baker, 1978). Under drought stress conditions Pak. $81 \times$ Inqlab 91 cross is expected to produced a population with improved biomass and grain yield.
General combining ability was found to be more important than specific combining ability for most of the traits except grain yield per plant under both environments. The presence of predominantly large amount of non-additive gene action for grain yield would necessitate the maintenance of heterozygosity in the population. Such genetic variability is non-fixable, The superior parents with high combining abilities have been used extensively in a breeding programme for widening the genetic base of the breeding material.
In conclusion, it may be stated that the diallel cross evaluated by different methods including the analysis of variance and genetic components analysis has yielded identical or comparable information on genetic structure of the characters studied. Adequacy test satisfied the additive dominance model. Genetic components were also computed for those traits who were partially fulfilled the assumptions

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for the validity of the additive dominance model. Invariably, in the present study additive genetic effects were found to be more important than nonadditive effects with the exception of grain yield per plant which appeared to be more closely linked to nonadditive genetic effects. Since the yield components like grains per spike, 1000-grain weight, tillers per plant were found to be significantly influenced by additive gene action, further progress in genetic improvement of. the yielding ability of wheat varieties may be attempted through such yield-related characters, with much better prospects of fixing desirable genes in a single homozygote.

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