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STUDY OF GENETIC COMPONENTS FOR YIELD AND QUALITY TRAILS IN BREAD WHEAT (*Triticum aestivum* L.)

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ABSTRACT

Ten bread wheat parental lines and their forty five hybrids were used for studying the character association among grain yield and its related characters. The results showed that the number of grain per spike, grain weight per spike, number of productive tillers per plant and harvest index were significantly and positively associated with grains yield per plant in both parental, and F1 generation.

Keywords : Genetic components, bread wheat, Triticum aestivum

Grains yield in wheat, as in any other crop plant is a complex multicomponent character. Fonseca and Patterson (1963) emphasized the significance of component approach in formulating a successful breeding programme. The present study examnines the nature of association of association of yield with other characters by utilizing parental and their all possible F1 hybrid lines.

MATERIALS AND METHODS

Forty five hybrids were obtained by crossing ten genotypes/cultivars namely HD 2285, K 8305, UP 2121, K8565, K 8030, HUW 234, PB W 226, K 8103, HUW 300 and HP 1633 in a 10 X 10 diallel mating system excluding reciprocals. The experiment 10 parents and their 45 F1s was grown in Randomized Complete Block Design with three replication at research farm of C.S.A. University of Agriculture and Technology, Kanpur and D.G. College Kanpur as well. The experiment was conducted in two row plots. The lengths of each row 1.5 m with inter and intra row distance of 25 and 15 cm, respectively. Recommended dose of fertilizers with fine irrigation was applied to raise a good crop. Ten plants for observation were randomly taken from each parent and F1 generation Data were recorded on 13 characters namely days to reproductive phase, flag leaf area, spike area plant

height, number of productive Tillers per plant, grain weight per spike, number of grains per spike, harvest index 1000-grain weight, grain yield per plants, seed hardness, protein content and tryptophan content. Simple correlation were worked out between grain yield and other characters for each parents and hybrids as per method suggested by Al jibouri *et al* (1958). The significance of phenotypic coefficient was tested against 'r' values from 'r' table of fisher and Yates (1938) for n-2 degree of freedom where n is a number of treatment.

RESULTS AND DISCUSSION

Estimate of phenotypic and genotypic association between all possible 78 pairs involving all the 13 characters separately for parental and first filial generation were computed. The results are presented in Table 1 and 2 for these generations respectively.

In parents, the correlation coefficient were positive and significant for 15 combinations out of 78 Days to reproductive phase was positively and significantly associated with flag area, 1000-grain weight and grain yield per plant. Flag leaf area was correlated positive and highly significant with grain yield per plant. Spike area was associated positive direction with grain yield per plant and protein content. Similar was the situation between numbers of productive tillers per plant with harvest index. Grain yield was highly significant and positively associated with grain yield per spike, number of grains per spike, harvest index and 1000-grain weight. Grain weight per spike was positively and significantly associated with 1000-grain weight.

Tryptophan content was also correlated positively and significantly with seed hardness.

In F1 generation, the significant associations were recorded in 12 cases out of 78. Among these the significant and positive values were found in 10 combinations which were days to reproductive phase with flag leaf area and 1000-grain weight per spike with protein content, number of reproductive tillers per plant with harvest index and grain yield per plant, grain weight per spike with 1000-grain weight and grain yield per plant, number of grain per spike with grain yield per plant, 1000-grain weight with grain yield per plant, seed hardness with tryptophan content. Negative and Significant association were recorded for grain weight per spike with number of grains per spike; number of grains per spike with 1000-grain weight. Rest of the combinations exhibited weak association.

The extent of genotypic interrelationship were higher than phenotypic association in both the parental and F1 generations under study involving 78 characters combinations in each case. Ahmad *et al.* (1978) reported similar finding in bread wheat. Hence the significant phenotypic association between characters were primarily due to genetic causes, which might be due to pleotropic effect rather than linkage between genetic affecting direct characters. The self-pollination mechanism is a prelude the fixation of blocks of genes as Well as due to limited chance of breaking linkage compared with random mating system prevailing in cross pollinated crops. Such reports have been made by Weber and Moorthy (1952) and Johson *et al.* (1995).

Grain yield per plant possessed positive and significant association with grain weight per spike, number of productive tillers per plant, number of grains per spike and 1000-grain weight in F1 generation and similar was the situation in parental generation. Hence, the selection for these component traits; could lead to higher productivity. Positive correlation was repoted by Donald and Hamblin (1976), Ali *et al.* (1978) Mc Vetty and Evans (1980) for grain yield with same component traits, Paroda and Joshi (1974), Virk and Anand (1970), Vrik and Singh (1972) and Ahmad *et al.* (1978) reported for grain yied with 1000 grain weight and Pandey *et al.* (1983), Sharma *et al.* (1987) and Ahmad *et al.* (1990) for grain yield with harvest index.

Positive and significant correlation between tryptophan content and seed hardness was observed and protein content was found positively associated with tryptophan content. The similar finding was reported by Gill and Brar (1973), Pandey (1980) and Kerkhi (1983).

A positive association between protein content and seed hardness provided a check as the seed hardness could be taken as indicator of protein level in grain. Therefore, a large number of germplasm could be screened for protein the help of seed hardness which is quickly *measurable* in the Labortory and field conditions.

The positive and signifivant association between protein content and tryptophan content in both the parental and F1 generations was accordance with Srivastava *et al.* (1971) who reported the increase of concentration with the increase in protein content in percent which might be fertilizers to experimental crops.

On generalization of the result, it may be concluded that maximum emphasis during selection should be given for grain weight per spike, number of grain per spike, number of prodctive tillers per plant and harvest index for the improvement of grain yield in bread wheat. Table1: Phenotypic (upper diagonal) and genotypic(lower diagonal) among 13 attributes in parental generation of a 10-parent-diallel cross in bread wheat

Days to reproductive phase • 0.74** 0.42 0.13 Flag leaf area 0.78 gp' 0.38 0.2 Flag leaf area 0.78 gp' 0.38 0.2 spike area 0.52 0.42 Gp' 0.15 Plant height 0.21 0.27 0.18 gp' No of productive 0.27 0.23 0.38 0.3 Itlers/plant 0.27 0.23 0.38 0.3 No of productive 0.15 0.23 0.3 0.3 Spike 0.15 0.23 0.3 0.3 No of grains per 0.15 0.22 0.14 0.3 Spike 0.01 0.22 0.14 0.3 Ionograin weight 0.72 0.32 0.07 0.1 Ionograin weight 0.12 0.12 0.12 0.1 Spike 0.72 0.12 0.12 0.07 0.1 Spike 0.12 0.12 0.12 <		tillers/plant per spike	grains per spike	Index	weight	yieldper plant	hardness	content	content
0.78 gp' 0.38 0.52 0.42 6p' 0.51 0.27 0.18 0.21 0.27 0.18 0.27 0.23 0.18 0.15 0.23 0.38 0.15 0.23 0.38 0.15 0.23 0.38 0.15 0.23 0.38 0.15 0.23 0.38 0.15 0.23 0.38 0.15 0.23 0.38 0.15 0.23 0.38 0.15 0.14 0.25 0.01 0.22 0.14 0.12 0.32 0.07 1 0.12 0.07 1 0.12 0.08 0.89 0.14 0.14	42 0.12	0.23 0.12	-0.01	-0.04	0.68*	0.84**	-0.10	0.23	0.24
0.52 0.42 Gp' 0.21 0.27 0.18 0.27 0.28 0.38 0.27 0.23 0.38 0.15 0.23 0.38 0.15 0.23 0.38 0.15 0.23 0.38 0.15 0.23 0.38 0.15 0.23 0.38 0.15 0.23 0.38 0.15 0.14 0.25 0.01 0.22 0.14 0.07 0.32 0.07 1 0.12 0.07 0.12 0.12 0.07 1 0.12 0.14 0.12 0.08 0.89	38 0.21	0.18 -0.08	0.17	0.24	0.01	0.82**	0.31	0.21	0.21
0.21 0.27 0.18 0.27 0.23 0.18 0.15 0.23 0.38 0.15 -0.74 0.25 0.01 0.22 0.14 0.01 0.22 0.14 0.01 0.22 0.14 0.01 0.22 0.14 1 0.22 0.14 0.01 0.22 0.14 0.01 0.22 0.14 0.12 0.32 -0.07 1 0.12 0.12 -0.07 1 0.12 0.13 -0.17 1 0.12 0.08 0.89	p' 0.15	0.32 0.21	0.12	-0.02	-0.01	0.84**	-0.17	0.65*	0.20
0.27 0.23 0.38 0.15 -0.74 0.25 0.15 -0.74 0.25 0.01 0.22 0.14 -0.02 0.32 -0.07 -0.02 0.32 -0.07 -0.12 0.12 -0.07 nt 0.12 0.08 0.89 0.87 0.89 -0.14	18 gp'	0.31 0.17	0.31	0.21	-0.16	-0.08	-0.31	-0.27	0.17
0.15 -0.74 0.25 0.01 0.22 0.14 0.01 0.22 0.14 -0.02 0.32 -0.07 .0.12 0.12 -0.07 .0.12 0.12 -0.07 .0.12 0.12 -0.07 .0.12 0.12 -0.07 .012 0.12 -0.07 .012 0.12 -0.07 .012 0.12 -0.07 .012 0.12 -0.07 .012 0.12 -0.07 .012 0.12 -0.07 .012 0.08 0.89	38 0.34	gp' 0.37	-0.16	0.72**	0.21	0.69*	0.17	0.31	0.21
0.01 0.22 0.14 -0.02 0.32 -0.07 -0.12 0.12 -0.07 ant 0.12 0.08 0.89 0.87 0.38 0.14	25 0.20	0.39 gp'	0.72**	0.31	0.87**	0.74**	0.32	0.12	0.18
-0.02 0.32 -0.07 0.72 0.12 -0.07 -0.07 0.12 0.18 0.38 -0.14 0.87 0.89 -0.14 -0.14	14 0.33	-0.14 0.74	,db	0.12	0.21	0.82**	0.07	0,02	0.05
0.72 0.12 -0.07 0.12 0.08 0.89 0.89 0.87 0.89 -0.14 -0.14	07 0.24	0.82 0.35	0.14	gpʻ	-0.62	0.72**	0.35	0.27	0.38
0.12 0.08 0.89 0.87 0.89 -0.14	07 -0.13	0.26 0.91	0.24	0.17	,db	0.71**	0.08	0.32	-0.21
0.87 0.89 -0.14	-0.06	0.78 0.78	0.85	0.80	0.75	,db	0.17	0.21	0.15
	14 -0.21	0.21 0.34	0.15	0.40	0.20	0.20	ġb	0.06	0.68*
Protein content 0.40 0.24 0.69 -0.2	59 -0.23	0.34 0.14	0.04	0.32	-0.21	0.27	0.10	gp'	0.39
Trytophan content 0.32 0.25 0.24 0.2	24 0.24	0.24 0.20	0.09	0.43	-0.12	0.18	0.74	0.28	gp

*Significant at 5% level;**Significant at 1% level

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Table2: Phenotypic (upper diagonal) and genotypic(lower diagonal) among 13 attributes in parental generation of a 10-parent-diallel cross in bread wheat

Character	Days to Reproduce tive phase	flag laef area	spike area	Plant height	No of productive tillers/plant	Grain weight per spike	No of grains per spike	Harvest index	1000grain weight	Grain yieldper plant	Seed hardness	Protein content	Trytophan content
Days to reproductive phase	,db	0.27*	0.12	-0.14	90.0	-0.13	-0.02	0.07	0.28*	60.0-	0.13	0.09	0.07
Flag leaf area	0.25	gp,	0.12	0.06	0.06	-0.11	0.06	0.14	-0.17	-0.04	0.05	0.14	-0.11
spike area	0.14	0.16	ġb	-0.05	0.10	0.10	-0.08	-0.08	-0.05	0.15	0.19	0.25	0.02
Plant height	-0.14	0.07	-0.06	db'	0.05	-0.01	0.10	0.08	-0.15	-0.09	0.01	-0.21	0.06
No of productive tillers/plant	60.0	0.07	0.17	0.06	,dɓ	0.13	-0.06	0.34**	60.0	0.39**	-0.11	0.05	0.07
Grain weight per spike	-0.15	-0.12	0.12	0.01	0.15	ġb	-0.59	0.17	0.45**	0.62**	0.10	0.02	0.13
No of grains per spike	-0.04	0.08	-0.10	0.12	-0.04	-0.68	ġp,	-0.04	-0.42**	0.26**	-0.29	-0.04	-0.01
Harvest index	-0.08	0.15	-0.09	0.08	0.00	0.17	-0.04	gp'	-0.11	0.16	0.22	0.15	0.20
1000grain weight	0.33	-0.18	-0.06	-0.15	0.12	0.52	-0.46	-0.10	gp'	0.29**	-0.06	-0.04	60.0-
Grain yieldper plant	-0.11	-0.66	0.17	-0.10	0.40	0.67	0.28	0.14	0.34	gp'	0.06	0.14	0.05
Seed hardness	0.10	0.08	-0.21	0.01	-0.18	0.13	-0.13	0.29	-0.07	0.08	gpʻ	-0.01	0.26*
Protein content	0.23	0.25	0.35	-0.28	0.02	0.04	-0.03	0.18	-0.02	0.29	-0.08	gp'	0.21
Trytophan content	0.08	-0.14	0.02	0.07	0.06	0.16	-0.01	0.23	-0.11	0.06	0.31	0.25	gp'

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*Significant at 5% level;**Significant at 1% level

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