

Variation and Heritability for Some Semolina Characteristics and Grain Yield and Their Relations in Durum Wheat (*Triticum Durum* Desf.)

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Abstract: The study were carried out to estimate variation, heritability and correlation for milling characteristics of 25 durum wheat (*Triticum durum* Desf.) genotypes under rainfed conditions in three environments of North West Turkey. The genetic material showed relative large variations in which all traits except test weight. The variance component due to genotype explained most of the total variation, ranging 55.94 to 0.01% of the variability associated with grain yield and each of semolina traits. Effects of G x E interaction ranged from a low of 0.09 of total variance for grain yield to a high of 65.56% of total variance for vitreous grain. Broad sense heritability estimates ranged between 89.4% (semolina colour) and 23.1% (wet gluten content) for all traits. Whereas protein content showed positive significant correlations with wet gluten content (0.708**), semolina colour (0.492**) and vitreous grain (0.445**), negative significant relationships of protein content were determined with 1000 - grain weight (-0.773**), test weight (-0.539**) and ash content (-0.446**). Vitreous grain was positively correlated with semolina colour (0.392**) and both traits showed significant correlations with wet gluten content (0.388-0.462) and protein content (0.445-0.492), respectively. The significant and negative correlation coefficient (-0.288) was observed between protein content and grain yield.

Key words: Durum wheat • Semolina value • Variance component • Heritability • Correlation

INTRODUCTION

Thrace and Marmara Region of Turkey are suitable for durum wheat production the point of view climatic conditions. Whereas the proportion of durum wheat was 60 % in 1960's, in 1980's this rate decreased to 5 % [1] and today, rate is in the level to be no. The main reasons have not improved new genotypes with high yielding and quality competing with bread wheat and wrong price policies. So, durum wheat breeding programs should be designed to ensure that any new released cultivar has acceptable agronomic characteristics to meet the demand of the farmers and the quality traits required by the processing industry.

The chief aim of a durum wheat breeding program is to obtain new cultivar with a high yielding potential and

better quality. There is genetic potential to improve the nutritional quality of durum wheat through convectional breeding techniques [2]. Durum wheat quality criteria are continually evolving in response to technological advances in durum wheat milling and secondary processing. The durum wheat physical condition is the most important factor determining wheat milling potential and end-use quality [3]. A high yield of highly refined semolina is the basic quality criteria valid today [4]. The semolina milling value can be defined as the capacity of a durum wheat to give high yields of semolina of a defined purity under industrial conditions [5] and it is of great importance regarding the physical quality character and complex because it is dependent on different factors. But, it is considered closely related with grain characteristics such as 1000 - grain weight and test

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weight characters which are the best predictors of grain size, shape and volume [6] and it is affected depending on a degree of vitreousness of endosperm [7]. Protein content, which is one of the most important traits in quality evaluation and breeding of durum wheat, are required to process semolina into a suitable final pasta product [8]. Wet gluten content of genotypes gives a good indication of protein content and because gluten strength can be judged by its extensibility and elasticity [9]. Additionally, ash content has been linked to semolina colour where it is used to assess the cleanliness of semolina [10]. Wheat quality is influenced by genotype, environment and their interaction [11-13], but relative magnitude of them on quality is unclear [14].

Quality improvement is to be possible via evaluation and selection [15], since wide variation exists in breeding material. Response to selection for quality depends on the heritability and genetic variance of quality traits and unfavourable correlated response with other important characteristics, in particular grain yield [16]. Correlations, although reliable only for the range of material tested, may point to relationships that can be utilised in making a selection programme more effective [17]. The objectives of this work were: (a) to determine the magnitude of variation present in a set of durum wheat genotypes; (b) to estimate variance components, heritability and genetic advance of the traits; (c) to investigate the relationships among the traits used with emphasis on protein content.

MATERIALS AND METHODS

Plant Material and Experimental Procedure: The genetic material of durum wheat (25 genotypes), which was comprised of 13 advanced lines, 8 mutant lines and 4 commercial varieties (Checks) were evaluated in the study.

Cite Characteristics and Agronomic Details: The study was carried out at Edirne (26°35' E, 41°38' N and elev. 32 m), Tekirdağ (27°34' E, 40°59' N and elev. 10 m) and Lüleburgaz (27°16' E, 41°22' N and elev. 41 m) locations of the Thrace Region during 2001-2002 growing year.

The chosen locations also differed, for instance, in the height above sea level, chemical composition of the soil and climate conditions. The height above sea level of Tekirdağ is H = 10 m which is a low-lying area, while Lüleburgaz's height above sea level is H = 41 m. The other location's altitude (Edirne) is H = 32 m. Soils characteristics of experiments were determined in soil analyses. Soils of the experiments were loamy clay, fine loamy and silty clay textured, neutral pH and low

salt concentrations in Tekirdağ, Lüleburgaz and Edirne, respectively. Contents of phosphorus, potassium, calcium, magnesium, iron, copper, manganese and zinc were higher than the required. Organic matter also was quite higher than 1%.

The Thrace Region is a peninsula and the central parts of Thrace Region is under effect of continental climate, along with Mediterranean climate and Black Sea climate, with lower temperatures during winters and hot, dry summers. The total cumulative precipitation recorded during growth season (November-July) for Tekirdağ, Lüleburgaz and Edirne were 513.4 mm, 486.2 mm, 380.4 mm in experiment year [18].

The plots were fertilized with 36 kg N (20-20-0) ha⁻¹ and 36 kg P₂O₅ ha⁻¹ at sowing, 87 kg N (Urea) ha⁻¹ at tillering and 41 kg N (NH₄NO₃) ha⁻¹ at pre-heading. Standard cultural practices were followed for raising the crop.

Methods: The experiments were designed in a random completed block design with three replications. Each experimental unit consisted of 6 rows, 5 m long and with 20-cm spaces between two rows. Standard cultural practices were followed for raising the crop.

The traits studied were protein content (PC), 1000 grain weight (TGW), test weight (TW), percentage of vitreous grain (VG), wet gluten content (WGC), ash content (AC), semolina colour b (SC) and grain yield (GY). Quality tests were performed on the harvested grains of each genotype for each location and traits were calculated at 14% moisture level and 100% purity. Protein content was determined as process by ICC Standard Method 105 [19]. Test weight was weighted using the Approved AACC Method 55-10 [20] using portable hectolitre test weight kit; results were reported in kilograms/hectolitre. Thousand grain weight was calculated from the weight (g) of 4 lots of 100 seeds, multiplying by 10. The percentage of vitreous grain was determined using the ICC Standard Method 129 [21]. A scalpel was used for cutting grains transversely. Vitreous grains (those completely free of starchy or speckled appearance) were separated from a 20 g clean grain sample and weighed. Vitreousness was calculated visually as a percentage of vitreous grains (w/w) in the sample. Ash content was fixed using the ICC Standard Method 104/1 [22]. The ICC Standard Method 106/2 [23] was used for wet gluten content and semolina colour b values of the genotypes were obtained using the ICC Standard Method 152 [24]. Grain yields obtained from each plot were adjusted to ton per hectare for biometrical evaluation.

Statistical Analysis: Estimates of broad sense heritability (h_{BS}^2) for different traits were computed using the variance components method based on the combined analyses over the three test locations.

Initial the data were subjected to randomised complete block design analyses of variance. Appropriate transformations (logarithmic, square root) were performed when necessary. Entry means were used in the combined analyses across environments. Variance components were estimated according to Snedecor and Cochran [25] as follows:

$$\begin{aligned}V_g &= (MS_g - MS_{gl})/rl, \\V_{gl} &= (MS_{gl} - MS_e)/r, \\V_e &= MS_e \text{ and} \\V_{ph} &= V_g + V_{gl} + r + V_e / rl\end{aligned}$$

Where V_g , V_{gl} , V_e and V_{ph} are the variances due to genotypes (lines), genotype x location (G x L) interaction, experimental error and phenotypes, respectively. MS_g , MS_{gl} and MS_e are the mean squares of genotypes (i.e. lines), genotype x location (G x L) interaction and pooled error and l denotes the number of environments (i.e. locations) and r the number of replications.

$$\text{Broad sense heritability } (h_{BS}^2) = [V_g / V_{ph}] \times 100$$

Genetic advance (GA), expressed as a percentage of the mean was calculated as:

$$GA = k \times (\text{Phenotypic variance})^{0.5} \times h_{BS}^2 (100/\bar{x}) [26],$$

where 'k' was the selection intensity at 5 % level (value = 2.06)

Simple correlation coefficients were determined among traits studied.

RESULTS AND DISCUSSION

Variations in Traits: The results of combined variance analysis made on data obtained from 3 locations are given in Table 1. Highly significant differences were detected among environments and genotypes for each of investigated traits. In general G x E interactions were significant, but relatively small than genotype and environment. While it was assumed that if the mean square for the genotype-environment interaction is solely determined by random reasons, CV is lower or equal to 5%, whilst the presence of this interaction increases values of the coefficient [27]. In addition,

Dotlacil *et al.* [28] considered a minimum 10% CV is a sign of wide diversity in wheat. Thus, the genetic material used are amendable to selection for improving grain yield and semolina characters for relative large variations in which all traits except test weight (Table 1).

There was a wide range between the minimum and maximum values of most traits, but test weight had low ranges (14% of the means) (Table 2). In contrast, wet gluten content showed the highest range of variation (112% of the means). The remaining traits exhibited wide variation (>55%). Wide phenotypic variations have been reported by Lerner *et al.* [29] and Paul *et al.* [30].

Component of variance for each trait expressed as percentage illustrates the relative contribution of each source to total variance in Table 2. The variance component due to genotype explained most of the total variation, ranging 55.94 to 0.01% of the variability associated with grain yield and each of semolina characters. Genotypic variance is a parameter which represents the magnitude of heritable effects. High degree of it indicates that selection can be successfully applied in this population [31]. The availability of significant genetic variability concerning SC, TGW, VG, AC and PC in population points out that selection might be conducted taking into consideration these traits in direction of increases or reductions.

Effects of G x E interaction ranged from a low of 0.09 of total variance for GY to a high of 65.56% of total variance for VG. Relatively large G x E interaction for the traits except SC and GY indicates that selection should be carried out in range of environments and it is compulsory to breed different genotypes for every specific environment [32]. Taking into account these results, it is recommended to develop of new genotype/cultivar for different environments regarding important semolina traits and grain yield which is considered as a major aim in durum wheat breeding studies. The main effects of genotype environment and their interaction, were statistically significant source of variation for all traits. Of greater interest was the relative magnitude of variation attributable to these effects, which were compared using variance component ratios [14] (Table 2). The σ_g^2/σ_e^2 ratio differed among the traits. A ratio > 1.0 indicates greater influence and stability of genetic factors relative to the variability associated with the interaction G x E. The ratios for all traits ranged from 10.0 (SC) to 0.41 WGC). The highest value calculated signifies a larger influence on variability by the genotype than the G x E interaction for SC. The ratios for WGC (0.41), PC (0.50) and AC (0.77) were < 1.0, indicating the important

Table 1: Mean square value of grain yield and semolina traits of 25 durum wheat genotypes over three environments

Traits	Source of variation					
	Environment	Genotype	G x E	Error	CV%	S.E.(±)
TGW (g)	2886.14**	100.39**	28.61**	2.69	13.90	0.407
TW (kg*hl ⁻¹)	136.52**	12.25**	4.22**	1.06	2.59	0.137
VG (%)	4126.72**	701.83**	198.37**	21.80	13.15	0.727
AC (%)	1.71**	0.13**	0.04**	0.004	10.46	0.013
SC	73.85**	40.09**	1.37**	0.07	13.48	0.154
WGC (%)	3034.15**	129.93**	61.87**	4.99	26.23	0.531
PC (%)	552.91**	10.42**	3.61**	0.06	18.37	0.164
GY (t*ha ⁻¹)	46.75**	2.93**	0.52**	0.24	16.33	0.062

TGW: Thousand grain weight, TW: Test weight, VG: Vitreous grain percent, AC: Ash content, SC: Semolina colour (b), WGC: Wet gluten content, PC: Protein content and GY: Grain yield

Table 2: Range, means, estimates of components of variance, broad-sense heritability (h_{BS}^2) and genetic advance for the characters of 25 durum wheat genotypes

	Range			Estimates of components of variance*					
	Min-Max	% of mean	Mean	σ_{ph}^2	σ_g^2	σ_{gl}^2	$\sigma_g^2 / \sigma_{gl}^2$	h_{BS}^2 (%)	GA as % of mean
TGW	33.5-62.0	65	43.9	19.30	7.98	8.65	0.92	41.3	8.52
TW	73.0-84.0	14	79.1	3.04	0.89	1.03	0.86	29.3	1.33
VG	42.0-98.0	68	82.9	143.19	55.94	65.56	0.85	39.1	11.63
AC	1.25-2.24	55	1.81	0.028	0.010	0.013	0.77	35.7	6.63
SC	12.1-23.6	67	17.1	4.81	4.30	0.43	10.0	89.4	23.63
WGC	14.0-48.0	112	30.4	32.78	7.56	18.33	0.41	23.1	8.98
PC	8.3-17.5	69	13.4	1.84	0.59	1.18	0.50	32.1	6.64
GY	3.11-7.50	77	5.7	0.61	0.27	0.09	3.00	44.3	12.46

* σ_{ph}^2 , σ_g^2 and σ_{gl}^2 are phenotypic, genotypic and genotype x environment interaction, respectively. $\sigma_g^2 / \sigma_{gl}^2$ is ratio of variances estimated for cultivar main effect and interaction.

influence of G x E interaction on these traits. The genotype and G x E interaction effects were almost similar for TGW, TW and VG due to their ratios ≈ 1.0 . Grain semolina value is a complex character that depends on numbers of traits and the individual contribution of each trait varies depending on the specific reaction to environmental conditions.

Heritability and Genetic Advance: The processing quality of durum wheat and our understanding of the factors that determine quality, has improved recently. This was accomplished despite limited knowledge of the mode of inheritance and heritability of the traits [2]. Heritability is often used by plant breeders and geneticists as a measure of precision of a trial or a series of trials for standardization of their selection units. Its main use is for computing the response to selection [33]. The magnitude of heritability was affected by the type of genetic material and yield level of the environment [34].

It has emphasized that heritability alone is not enough to make sufficient improvement through selection unless accompanied by substantial amount of genetic advance [35]. Broad sense heritability estimated on the basis of genotypic and phenotypic variances ranged between 89.4 and 23.1% for all traits. WGC had the lowest heritability value (23.1%) followed by TW (29.3%). The heritability estimates were small for most of the characters due to larger phenotypic variances, indicating the growing areas effect. Only heritability of SC was greater than that of GY. In the present study SC had the highest (89.4%) broad sense heritability. The heritabilities for TGW, VG, AC and PC were 41.3%, 39.1%, 35.7% and 32.1%, respectively. The heritability estimate was 44.3% for GY (Table 2). The highest heritability (89.4%) coupled with high genetic advance (23.6%) of SC suggested that selection could be practised for this trait. Low heritability and low genetic advance were observed for WGC and TW indicating low transfer of these traits in the subsequent generations.

Table 3: Coefficients of correlation among traits studied of durum wheat grown in 3 sites of Northwest Region of Turkey during 2001-2002 growing season (n = 25)

	TW	VG	AC	SC	WGC	PC	GY
TGW	0.545**	-0.437**	0.463**	-0.409**	-0.680**	-0.773**	0.339**
TW		-0.336**	0.282**	-0.391**	-0.431**	-0.539**	0.381**
VG			-0.334**	0.392**	0.388**	0.445**	-0.377**
AC				0.033	-0.343**	-0.446**	0.391**
SC					0.462**	0.492**	-0.203**
WGC						0.708**	-0.324**
PC							-0.288**
GY							

Moderate heritability and high genetic advance estimates for TGW and VG reflect the possibility of effective selection for these traits. Moreover moderate heritability and genetic advance for AC and PC recommended that selection should be delayed to more advanced generations for those traits. But, selection for PC in wheat is complicated due to the negative relationship with GY and the influence of environmental conditions on PC. Our findings are in agreement with Clarke *et al.* [36] who stated that heritability for PC ranged from 29% to 53%. Bilgin *et al.* [37] determined the heritability and genetic advance of 71.7-32.5%, 49.3-11.7% and 62.4-10.1% for TGW, TW and VG, respectively. Johnston *et al.* [38] notified broad sense heritability of 31 to 69% for semolina colour and also they stated that it was a highly heritable trait and improvement of colour should be possible through early generation selection. Budak [39] determined the heritability of 29% for TW. Akçura [40] found heritability values of 35.4%, 39.9%, 32.6% and 50.0% for TGW, PC, AC and SC, respectively. The heritability estimates in present study were much lower than these ones exhibited differences from these findings. But, Yağdı and Sözen [41] estimated lower heritability values for TGW, GC, PC and TW than that of ours.

Correlations among Traits: Nearly all correlations among traits except correlation between AC and SC were significant (Table 3). Whereas PC showed positive significant correlations with WGC (0.708), SC (0.492) and VG (0.445), negative significant relations of PC were determined with TGW (-0.773**), TW (-0.539**) and AC (-0.446**). Rharrabti *et al.* [42], Eslami *et al.* [43] and El-Khayat *et al.* [44] reported that positive correlations among PC, WGC, SC and VG. The degree of vitreousness of durum wheat grains are related to protein composition of the grains [45], this observation was to be expected. Grain physical quality of durum wheat, which depends on TGW and TW influence the semolina production [6]. Oak *et al.* [46] reported similar relations for PC, TGW and TW.

VG was positively correlated with SC (0.392**) and both traits showed significant correlations with WGC and PC. Thus, it appears that increasing for VG and SC could lead to increases in PC under the conditions of our study.

The significant and negative correlation coefficient (-0.288) was observed between PC and GY. This relationship had been encountered in barley [47], in spring wheat [48], in durum wheat [49] and in triticale [50]. This negative association between PC and GY mainly caused from dilution of protein by non-nitrogen compounds in the grain [51].

CONCLUSIONS

The availability of significant genetic variability concerning SC, TGW, VG, AC and PC in population points out that selection might be conducted taking into consideration these traits in direction of increases or reductions. Genotypic main effects were found to have a significant on semolina traits, but contributed a smaller proportion of variability as compared with either environment or genotype x environment interaction effects. The high variance ratio from 1.0, heritability (89.4%) coupled with high genetic advance (23.6%) of SC suggested that selection could be successfully practised for that trait. The results of our correlation analyses indicate that VG should be considered as a selection criterion for future durum wheat quality breeding programmes.

ACKNOWLEDGMENT

We acknowledge financial support from Agriculture, Forestry and Food Technologies Research Grant Committee of The Scientific and Technical Research Council of Turkey. Thrace Agricultural Research Institute and Barilla G.E.R. Fratelli-Società per Azioni Company deserve gratitude for helping in the carrying the experiments and for their contribution to the quality analyses.

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