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CORRELATION AND PATH COEFFICIENT ANALYSES OF GRAIN YIELD AND ITS RELATED COMPONENTS FOR SOME COMMON WINTER WHEAT GENOTYPES

Gergana N. Desheva*, Manol G. Deshev

Institute of Plant Genetic Resources “Konstantin Malkov”, Sadovo, Druzhiba Str., 4122, BULGARIA

*E-mail: gergana_desheva@abv.bg

Abstract

The aim of the study was to determine the interrelationship and the direct and indirect effects of some yield components among themselves and with the grain yield in the 49 common winter wheat varieties originating from different countries. The experiment was conducted in the experimental field of IPGR-Sadovo, Bulgaria during 2017-2019 growing seasons in the randomized block design in three replications and 10 m² plot size. Data were recorded for a number of days to heading, plant height, spike length, number of spikelets per spike, number of grains per spike, grain weight per spike, thousand-kernels weight and grain yield per hectare. Grain yield per ha correlated positively and significantly with plant height, spike length, grain weight per spike and thousand grain weight. Path coefficient analysis revealed that the grain weight per spike, spike length and plant height had the highest direct positive effect on the grain yield per hectare. The results of study could be useful as selection criteria to increase the grain yield in common winter bread wheat.

Key words: correlation, path analysis, common winter wheat, yield, yield components

INTRODUCTION

One of the main goals in wheat breeding is to increase the grain yield. The wheat yield is the complex polygenic trait, depending on genetic and environmental factors and their interaction (Shara et al., 2016; Chitralkha et al., 2017). The genetic architecture of the yield can be resolved better by studying its component characters as: plant height, number of productive tillers per plant, number of spikelet per spike, number of grains per spike, grain weight per spike, thousand kernels mass and others (Khan and Dar, 2010). Initiating a breeding program based on the yield component requires knowledge of the relationship between the yield and its component characters. Therefore, the correlation is a realistic technique to develop selection criteria mostly based on phenotypic characters used for the improvement of yield through plant breeding programs (Ghallab et al., 2017; Shahidullah et al., 2017;

Noopur et al., 2019). The importance to assess the correlations present among the different traits to be improved has long been recognized by a number of works. Some of the researchers indicated the positive correlation between the grain yield and the yield components traits in wheat such as: the plant height (Kamani et al., 2017), the number of productive tillers per plant (Bhutto et al., 2016; Kamani et al., 2017; Mecha et al., 2017), the number of spikelets per spike (Mecha et al., 2017; Dagade et al., 2020), the number of spikes per plant (Fellahi et al., 2013), the number of grains per spike (Bhutto et al., 2016; Shahidullah et al., 2017; Chitralkha et al., 2017; Dagade et al., 2020), the number of spikes per m² (Sokoto et al., 2012; Renu et al., 2018), the biological yield (Kamani et al., 2017; Shahidullah et al., 2017; Verma et al., 2019) and the harvest index (Ali and Shakor, 2012; Kamani et al., 2017; Mecha et al., 2017; Verma et al., 2019).

Simple correlation analysis indicates the degree of association between the traits, but it can't provide reasons for the association. The better understanding of the association is provided by the path coefficient analysis (Shah et al., 2010; Desheva, 2016). It helps in partitioning of correlation coefficients into direct and indirect effects and in the assessment of the relative contribution of each component character to the yield (Verma et al., 2019). Noopur et al. (2019) noted that the information related to the nature and extent of association among the various yield attributes, the direct and indirect effects of each component on the yield are helpful in formulating an effective breeding strategy.

The aim of the study was to determine the interrelationship and the direct and indirect effects of some yield components among themselves and with the grain yield in the common winter wheat.

MATERIALS AND METHODS

The present study was carried out in the experimental field of the Institute of Plant Genetic Resources “Konstantin Malkov” - Sadovo during 2017-2019 growing seasons which is located at longitude 024°55'56.3''E and latitude 47°07'42.6''N with an average

altitude of 166 m above sea level in the south of Bulgaria. Forty eighth common winter wheat varieties from different countries (Bulgaria, the Czech Republic, Portugal, France, Poland, Italy, Spain, and Greece) and one accession with unknown biological status and origin were examined (Table 1). The experiment was conducted in the randomized block design with three replications and 10 m² plot size. Normal agronomic and cultural practices were applied to the experiment throughout the growing seasons. In the phase of full maturity of the plants, 20 plants were selected from each accession for biometric measurements. Data were recorded for plant height, spike length, number of spikelets per spike, number of grains per spike, grain weight per spike, thousand-kernels weight and grain yield per hectare. Number of days to heading is calculated from 1 January to the full heading.

Statistical analyses were performed using the statistical program SPSS 19.0. Phenotypic correlations among the various characters were calculated by using phenotypic variances and covariance (Lidansky, 1988). The path coefficient analysis was performed to assess the direct and indirect effects of the measured traits on the grain yield according to Dewey & Lu (1959).

Table 1. List of accessions included to the study

№	Accession number	Plant species	Subspecies	Name of cultivar	Origin
1	A1BM0256	<i>Triticum aestivum</i> L.	var. <i>erythrospermum</i> (Koern.) Mansf.	Enola	BGR
2	B2000213	<i>Triticum aestivum</i> L.	var. <i>lutescens</i> (Alef.) Mansf.	Ines	CZE
3	B2000217	<i>Triticum aestivum</i> L.	var. <i>lutescens</i> (Alef.) Mansf.	Bohemia	CZE
4	B2000218	<i>Triticum aestivum</i> L.	var. <i>lutescens</i> (Alef.) Mansf.	Baletka	CZE
5	B2000226	<i>Triticum aestivum</i> L.	var. <i>lutescens</i> (Alef.) Mansf.	Secese	CZE
6	B2000227	<i>Triticum aestivum</i> L.	var. <i>lutescens</i> (Alef.) Mansf.	Seladon	CZE
7	B2000228	<i>Triticum aestivum</i> L.	var. <i>lutescens</i> (Alef.) Mansf.	Silueta	CZE
8	B2000229	<i>Triticum aestivum</i> L.	var. <i>erythrospermum</i> (Koern.) Mansf.	Coa	PRT
9	B2000230	<i>Triticum aestivum</i> L.	var. <i>erythrospermum</i> (Koern.) Mansf.	Jordao	PRT
10	B2000234	<i>Triticum aestivum</i> L.	var. <i>lutescens</i> (Alef.) Mansf.	Andalou	FRA
11	B2000236	<i>Triticum aestivum</i> L.	var. <i>lutescens</i> (Alef.) Mansf.	Muza	POL

12	B2000239	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Kobra Plus	POL
13	B2000241	<i>Triticum aestivum</i> L.	var. erythrosperrum (Koern.) Mansf.	Alcione	ITA
14	B2000242	<i>Triticum aestivum</i> L.	var. erythrosperrum (Koern.) Mansf.	Bilancia	ITA
15	B2000243	<i>Triticum aestivum</i> L.	var. erythrosperrum (Koern.) Mansf.	Carisma	ITA
16	B2000244	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Delfino (P 204 A)	ITA
17	B2000245	<i>Triticum aestivum</i> L.	var. erythrosperrum (Koern.) Mansf.	Esperia	ITA
18	B2000246	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Gemini	ITA
19	B2000247	<i>Triticum aestivum</i> L.	var.milturum (Alef.) Mansf.	Primoasi	ITA
20	B2000248	<i>Triticum aestivum</i> L.	var. erythrosperrum (Koern.) Mansf.	Accor	FRA
21	B2000182	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Navarro 150	ESP
22	B2000189	<i>Triticum aestivum</i> L.	var.ferrugineum (Alef.) Mansf.	Albimonte	ESP
23	B2000209	<i>Triticum aestivum</i> L.	var. erythrosperrum (Koern.) Mansf.	Amarok	FRA
24	B9BM0070	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Geya-1-2010	BGR
25	B2BM0106	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Geya-1	BGR
26	B2000249	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Allister	FRA
27	B2000252	<i>Triticum aestivum</i> L.	var. erythrosperrum (Koern.) Mansf.	Autan	FRA
28	B2000253	<i>Triticum aestivum</i> L.	var. erythrosperrum (Koern.) Mansf.	Colfiorito	ITA
29	B2000254	<i>Triticum aestivum</i> L.	var. erythrosperrum (Koern.) Mansf.	Collerosso	ITA
30	B2000255	<i>Triticum aestivum</i> L.	var. erythrosperrum (Koern.) Mansf.	Vittorio	ITA
31	B2000257	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Bardotka	CZE
32	B2000258	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Bazilika	CZE
33	B2000260	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Nikol	CZE
34	B2000261	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Bodycek	CZE
35	B2000262	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Elly	CZE
36	B2000263	<i>Triticum aestivum</i> L.	var. erythrosperrum (Koern.) Mansf.	Jindra	CZE
37	B2000265	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Matylda	CZE
38	B2000266	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Canadair	FRA
39	B2000268	<i>Triticum aestivum</i> L.	var.ferrugineum (Alef.) Mansf.	Oropos	GRC
40	B2000269	<i>Triticum aestivum</i> L.	var.aureum (Link) Mansf.	Nestor	GRC
41	B2000275	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Altigo	FRA
42	B2000276	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Aldric	FRA
43	B2000274	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Acienda	FRA
44	B2000169	<i>Triticum aestivum</i> L.	var. erythrosperrum (Koern.) Mansf.	Saturnus	AUT
45	B2000171	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Mahissa-1	ESP
46	B2000176	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Dimas	ESP
47	B2000177	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Estrella	ESP
48	B3000080	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.		unknown
49	A1BM0322	<i>Triticum aestivum</i> L.	var.lutescens (Alef.) Mansf.	Sadovo 772	BGR

RESULTS AND DISCUSSION

Phenotypic correlation

The phenotypic correlation reveals the degree of association between different characters and thus aids in the selection to improve the yield and the yield attributing characters simultaneously (Ghallab et al., 2017; Verma et al., 2019). The positive correlation between desirable traits is supposed to be favourable and help to breeder in selection whereas, the negative correlations hinder the recovery of the combinations in both characters (Chitralkha et al., 2017). The phenotypic correlation coefficients of the grain yield and its related components in the investigated 49 common winter wheat genotypes are presented in Table 2. The grain yield had a positive and significant correlation with the plant height ($r=0.1821^*$), spike length ($r=0.301^{**}$), grain weight per spike ($r=0.204^*$) and thousand-kernels weight ($r=0.180^*$). The selection for these characters could be directly followed by an immediate yield improvement of the wheat crop (Nukasani et al., 2013). Generally, in those characters in which the grain yield showed a positive and significant correlation, there were component interactions in which a gene conditioning an increase in one character will also influence another character provided the other conditions are kept constant (Mecha et al., 2017). Sokoto et al. (2012), Nasri et al. (2014) and Ojha et al. (2018) also confirm that the plant height, spike length and 1000-grains weight have an importance in yield determination. Mecha et al. (2017), noted that the grain yield had a positive correlation with the spike length and the thousand-kernels weight, while according to Shara et al. (2016), the 1000 kernel weight was in a negative relationship with the grain yield. The positive correlation of the plant height with the grain yield was recorded by Nasri et al. (2014), while the negative by Joshi et al. (2008) and Chitralkha et al. (2017). Tsenov et al. (2009), Rachovska and Uhr (2010), Chamurliyski et al. (2011) and Nukasani

et al. (2013) confirmed the positive relationship between the yield and the grain weight per spike, while Mohsin et al. (2009) between the yield and the spike length. Al-Najjar and Al-Zubaidy (2020) observed a significant and positive correlation between the yield, the plant height and the thousand-kernels weight. Chamurliyski et al. (2014) found a positive relationship between the grain yield, the grain weight per spike and the thousand-kernels weight.

The plant height correlated positively and significantly with the number of days to heading ($r=0.152^*$), which was also observed by Yadav et al. (2019) (Table 2). The spike length was in a positive relationship with the number of days to heading ($r=0.141^*$) and the plant height ($r=0.221^{**}$). The last correlation was in agreement with our previous study (Desheva, 2016) and the reports of Sokoto et al. (2012), Ghallab et al. (2017), Chitralkha et al. (2017), Stojsin et al. (2018), Yadav et al. (2019) and Kumari et al. (2020).

The positive correlations appeared between the number of spikelets per spike, the number of days to heading and the spike length, respectively ($r=0.147^*$ and $r=0.233^{**}$) (Table 2). The correlation between the number of spikelets per spike and the spike length was in line with the findings made by Nikolova-Andreeva (2011), Bhutto et al. (2016), Mecha et al. (2017), Stojsin et al. (2018) and Dimitrov et al. (2019).

The number of grains per spike correlated positively and significantly with the spike length ($r=0.293^{**}$) and the number of spikelets per spike ($r=0.291^{**}$) (Table 2). Our findings were in conformity with Sokoto et al. (2012) and Desheva et al. (2016). Meena et al. (2014), Dutamo et al. (2015), Ojha (2018) and Dimitrov et al. (2019) also noted a positive correlation between the number of grains per spike and the number of spikelets per spike. Stoyanov (2013), Shrief et al. (2019), Haydar et al. (2020) and Bonchev (2020) observed a positive linear relationship of the number of

grains per spike with the spike length.

The grain weight per spike correlated positively with the spike length ($r=0.432^{**}$), the number of spiklets per spike ($r= 0.306^{**}$) and the number of grains per spike ($r=0.717^{**}$) (Table 2). The results were coincident with those reported by Stojsin et al. (2018). Kamani et al. (2017) and Dimitrov et al. (2019) also found a positive relationship between the grain weight per spike and the number of grains per spike. Meena et al. (2014) in contrast to our study found a negative relationship between the grain weight per spike and the number of spiklets per spike and the number of grains per spike.

The thousand-kernels weight showed a positive correlation with the plant height ($r=0.218^{**}$), the spike length ($r=0.218^{**}$) and the grain weight per spike ($r=0.471^{**}$), while it was negative with the number of grains per

spike ($r=-0.250^{**}$) (Table 2). The negative relationship found in this study was in agreement with previous reports of Fellahi et al. (2013), Shara et al. (2016), Mecha et al. (2017), Bonchev (2020) and in contraversaly with Kamani et al. (2017). A positive correlation between the thousand-kernels weight and the grain weight per spike was recorded by Nikolova-Andreeva (2011), Nikolova (2012), Stoyanov (2013), Bhutto et al. (2016) and Stoyanov and Dunchev (2020), while Yadav et al. (2019) observed a negative correlation of the thousand-kernels weight with the spike length. Dragov and Dechev (2016) found a positive relationship between the thousand-kernels weight and the plant height in durum wheat.

Shrief et al. (2019) confirmed a positive relationship between the thousand-kernels weight and the spike length.

Table 2. Phenotypic correlation coefficients of grain yield and its related components in common winter wheat

Variable	NDH	PH	SL	NSS	NGS	GWS	TKW	GY
NDH	1							
PH	0.152*	1						
SL	0.141*	0.221**	1					
NSS	0.147*	0.086	0.233**	1				
NGS	-0.023	-0.134	0.293**	0.291**	1			
GWS	0.056	0.046	0.432**	0.306**	0.717**	1		
TKW	0.117	0.218**	0.218**	0.019	-0.250**	0.471**	1	
GY	0.086	0.182*	0.301**	-0.028	0.067	0.204**	0.180*	1

*. Correlation is significant at the 0.05 level (2-tailed), **. Correlation is significant at the 0.01 level (2-tailed).

NDH- number of days to heading, PH- plant height, SL- spike length, NSS-number of spiklets per spike, NGS-number of grains per spike, GWS- grain weight per spike, GY- grain yield per hectare, TKW-thousand-kernels weight

Path coefficient analysis

The path coefficients studied the causes and effects and also studied the relationships between the variables which are differentiated the associations into direct and indirect effects through other dependent variables (Kumari et al., 2020). Generally, the characters that exerted a positive direct effect and a positive and

significant correlation coefficient with the grain yield were known to affect the grain yield in the favourable direction and need much attention during the process of selection cases (Ojha et al., 2018). The direct and indirect effects of the investigated 7 characters (independent variables) on the grain yield per hectare (dependent variable) are presented in Table 3.

The grain weight per spike had the greatest positive effect (0.525) on the grain yield per hectare, followed by the spike length (0.242) and the plant height (0.111). Therefore a slight increase in one of these traits may directly contribute to the grain yield per ha. These traits are very important components of the grain yield and should be given a high weightage in any selection process aimed at improving grain yield in common winter wheat. Nukasani et al. (2013), Meena et al. (2014) also exhibited a maximum positive direct effect of the grain weight per spike on the yield. Mohsin et al. (2009) also indicated that the grain yield had a positive direct effect with the spike length and the number of grains per spike. Al-Maliky (2018) found the highest direct effect of the plant height on the grain yield variation. Shara et al. (2016) in contraversion to our study found a negative direct effect of the grain weight per spike on the grain yield, while Ojha et al. (2018) noted a negative direct effect of the plant height. The deviations of our results from those of the above mentioned authors may be due to the different set of samples included in the

experiments, on the one hand, and on the other, due to the fact that the yield as a polygenic nature is strongly influenced by fluctuations in the environment.

The highest negative direct effect was observed for the number of grains per spike (-0.383). The direct effects of the thousand-kernels weight and the number of spikelets per spike on grain yield per ha were also negative, respectively (-0.243 and -0.145) (Table 3). Mecha et al. (2017) in contrast to our study pointed out that the thousand grains weight and the number of grains per spike showed a positive direct effect.

The highest positive total indirect effect showed the number of grains per spike (0.450), followed by the thousand-kernels weight (0.423) and they were controlled via the grain weight per spike (0.376 and 0.247). Only the grain weight per spike had a negative total indirect effect (-0.321) via the number of grains per spike, the thousand-kernels weight, and the spike length, respectively (-0.275, -0.114 and 0.104) (Table 3).

Table 3. Direct (Bold and Underline) and indirect effect of the 7 characters (independent variables) on the grain yield per hectare (dependent variable) in 49 common winter wheat genotypes. The last column shows the phenotypic correlations (*r*) of the independent variables with the grain yield per hectare

Variable	NDH	PH	SL	NSS	NGS	GWS	TKW	Total indirect effect	r
NDH	<u>0.047</u>	0.017	0.034	-0.021	0.009	0.029	-0.028	0.039	0.086
PH	0.007	<u>0.111</u>	0.053	-0.012	0.051	0.024	-0.053	0.071	0.182
SL	0.007	0.025	<u>0.242</u>	-0.034	-0.112	0.227	-0.053	0.059	0.301
NSS	0.007	0.010	0.056	<u>-0.145</u>	-0.111	0.161	-0.005	0.117	-0.028
NGS	-0.001	-0.015	0.071	-0.042	<u>-0.383</u>	0.376	0.061	0.450	0.067
GWS	0.003	0.005	0.104	-0.044	-0.275	<u>0.525</u>	-0.114	-0.321	0.204
TKW	0.005	0.024	0.053	-0.003	0.096	0.247	<u>-0.243</u>	0.423	0.180

NDH- Number of days to heading, PH- Plant height, SL- Spike length, NSS-Number of spikelets per spike, NGS-Number of grains per spike, GWS- Grain weight per spike, TKW- thousand-kernels weight

The phenotypic correlation analysis indicated that the thousand-kernels weight and the number of grains per spike had positive

influences on the grain yield per ha (Table 2), but the path coefficient analysis suggested that they had a direct negative influence on the grain

yield per ha (Table 3). The results showed that the direct negative effect of the thousand-kernels weight and the number of grains per spike on the grain yield per ha were masked from the positive indirect effect of these characters through the grain weight per spike (Desheva, 2016). Therefore, their positive effect on improving the yield was indirect through the increment of grain weight per spike.

Generally, the results of the correlation and path analysis obtained plant height, spike length and grain weight per spike, showed a significant positive correlation with the maximum positive direct effect on the grain yield. Therefore, a direct selection on the bases of these traits could be effective for the grain yield improvement of bread wheat for the studied location.

CONCLUSION

The results obtained from 49 bread wheat genotypes showed that the grain yield per hectare was significantly and positively correlated at a phenotypic level with the plant height, the spike length, the grain weight per spike and the thousand-kernels weight. The path coefficient analysis indicated that the grain weight per spike had the greatest positive effect on the grain yield per hectare, followed by the spike length and the plant height. The number of grains per spike, the thousand-kernels weight and the number of spikelets per spike had a direct negative effect on the yield grain per ha. Therefore, the characters - grain weight per spike, spike length and plant height are the most important characters affecting the grain yield variation and consequently may be considered as effective criteria to increase grain yield in common winter wheat.

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