

Grain size and other agronomical traits variation in a winter wheat population of doubled haploid lines

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Abstract

In wheat, the size of the grain, respectively its dimensions as well as degree of filled, are important characteristics on which depends both the weight of the grain and yield of flour, the quality of milling and baking as well as the production capacity of the respective genotype. This paper presents the results obtained by studying for three years, under field condition, 85 doubled haploid (DH) lines obtained from the F1's of 'G.603-86' (large grains genotype) × 'F.132' (normal grains genotype) crosses using biotechnological *Zea* system. The environmental conditions of the three years had an important contribution on the genotype × year interaction, which showed also a higher influence on 1000 kernel weight (TKW). The variability of plant height and ear emergence data was also affected to a similar extent by this interaction. Based on the performed results and analyses, were highlighted lines which show high and stable values of TKW (54-64 g), associated with a plants height of approximately 85-100 cm and an ear emergence from May 11 to 17, under some climatic conditions similar to the period of study. These doubled haploids lines can be considered as promising genotypes for using in wheat breeding programs in order to improve yield performances under temperate continental climate conditions.

Keywords: DH lines; flowering date; plant height; stability; TGW; wheat

Introduction

In wheat, the grain size respectively its dimensions as the degree of filling are important characteristics depending upon both the weight of the grain and the yield of the flour respectively the quality of the milling. Being also considered to be components of production with highest phenotypic stability, these attributes have been and remain constant concerns of breeding programs. Several studies argue that the progress of selection for a superior production capacity is directly related to the gradual increase of the grain size and degree of filling. Recently results based on eight years tests in different Southeast European countries including Romania with 422 prospective wheat lines and new varieties originated from the main wheat grown globally areas highlight the importance of 1000 kernel weight (TKW) and volumetric mass (VM) in achieving high production (Sharma *et al.*, 2014).

Undoubtedly, similar advances have been made in other breeding programs, given the importance of these attributes in expressing the productivity of newly created varieties. As a result, searches for identifying

new sources for TKW and VM have been recently become targets of real interest in breeding and genetic studies even at the molecular level.

It should be mentioned that genotypes with higher values for TKW and for the grain shape and size were also obtained by using mutagenic procedures with physical and chemical mutagens (Cheng *et al.*, 2015; Zhang *et al.*, 2015; Dobre and Giura, 2016). Also, in related species among which *Triticum spelta* and *Aegilops tauschii squarrosa* were identified in their hybrid progeny with common wheat forms with superior values for the grain size and grain weight (Giura, 2010; Xie *et al.*, 2015).

The first genetic analysis of grain size, their dimensions and the degree of grain filling in wheat, have stated a complex genetic determinism by genes located on the majority of chromosomes and this in relation with the other plant morphology and physiological attributes. There were also identified the chromosomes involved in controlling specific traits, the same or different ones from one genotype to another (Law, 1967; Petrovici and Worland, 1968; Halloran, 1976; Chojeki *et al.*, 1983; Snape *et al.*, 1985; Giura and Săulescu, 1996). Subsequently, through molecular analysis, molecular markers/QTL's associated with some of these attributes have also identified capitalizing that the required information can provide new chances for the creation of varieties with superior productivity (Börner *et al.*, 2002; Ramya *et al.*, 2010; Cui *et al.*, 2011; Simmonds *et al.*, 2014; Li *et al.*, 2019; Ma *et al.*, 2019).

One of the new sources of grain size is also the autumn wheat line 'G.603-86' obtained at National Agricultural Research and Development Institute (NARDI) Fundulea from the cross 'Cologna lunga' × 'F.6-75'. This line is characterized by very high values for TKW and for grain size: over 60 mg on average for several years and a grain length of 8.5-9.0 mm. Genetic analysis based on F3 disomics, cytologically extracted for each chromosome from the F2 monosomic populations ('Favorit'-monosomics × 'G.603-86') revealed a complex genetic control exerted by several chromosomes: some with positive effects, other with negative effects on weight, size and degree of grain filling (Giura and Săulescu, 1996).

However, in practical breeding this complex genetic determinism can impeded to a great extent, the transfer of the respective attributes by using classical methods without to use specific molecular markers, during selection cycles. In addition, the line 'G.603-86' of 90-93 cm height carries *RhtB1b* and *RhtD1b* loci and consequently in the crosses with genotypes carrying *RhtB1b* and/or *RhtD1b* can appear even higher forms due to allelic difference at *PpdD1* loci. The line also shows sensitivity to some foliar diseases, especially to yellow rust and brown rust as well a clear flowering and maturity delay compared to current varieties better suited to local growing conditions. In addition, in some years, the line exhibits a substantial strain elongation due probably to the presence in genome of a different other genes for photoperiod and vernalization requirements. Probably these attributes are inherited from the 'Cologna lunga' genitor originated from a local Italian old wheat population with unknown genealogy but with a presumptive ascendance in the *Triticum turgidum* sp. (Giura, 2010).

Doubled haploid (DH) production has been successfully used in breeding and genetic analysis of qualitative and quantitative traits in wheat. The main advantage of DH technology consists in the considerable reduction of time necessary to obtain homozygous lines (Ren *et al.*, 2015; Patial *et al.*, 2019), which is an important breakthrough to speed up the cultivar development programs (Dunwell, 2010). In wheat, haploids can be mainly produced by *in vitro* androgenesis and interspecific crosses with: maize, related species to maize (*Zea mays* L. spp. *Mexicana*, *Tripsacum dactyloides*); sorghum *Sorghum bicolor*), pearl millet (*Pennisetum glaucum*); Job's tears (*Coixlachryma-jobi*) (Mochida and Tsujimoto, 2001). High ploidy level and the D genome of wheat have an important role in DH production, using wide hybridization with maize (Niu *et al.*, 2014).

This paper presents and discusses the data for 2004, 2005 and 2016 regarding plant height, ear emergency and thousand kernel weights. The aim of this paper is to identify DH lines with higher TKW values indifferently of environmental conditions, with a similar vegetative period and plant height as modern Romanian cultivars.

Materials and Methods

Biological material and experimental design

For a more comprehensive analysis of the complexity of genetic control of the grain size, grain dimensions and their connection to other plant traits, we developed using the *Zea* system a mapping population of 85 DH lines. For this purpose, line 'F.603-86' was crossed with the breeding line 'F.132' with small grains but carrying the *RhtB1b* and *RhtD1a* genes for plant height control, *Lr34* and *Lr67* genes for brown rust resistance and a vegetative period corresponding to local conditions, being earlier with about 10-12 days compared to 'G.603-86' line. The F1 hybrids were then crossed by maize in greenhouse condition and haploid plants regenerated by *in vitro* culture of immature embryos. Regenerated plantlets were treated with colchicine to obtain doubled haploid (DH) lines.

The resulted 85 DH lines and parental forms were then studied at NARDI Fundulea in a field trial in 2004, 2005 and 2016 years respectively, using a randomized block design with three replications. The average monthly temperatures and rainfall in Fundulea, during the period of study are presented in Figure 1. The genotypes were sown in pair rows of 1 m long, spaced apart at 25 cm between rows and 50 cm between pairs. Plant height was measured from the soil level to the top of the spike, without awns, using ten plants /replicate for each genotype. Ear emergence date was recorded when 50% of the spikes from each plot had visible awns, and counted as number of days after/before 1 May. At maturity, 30 spikes (10 spikes/plot) were randomly selected and harvested for each DH line and parental form. The TKW was estimated by weighing the quantity of the seeds per each spike and divided by seed number.

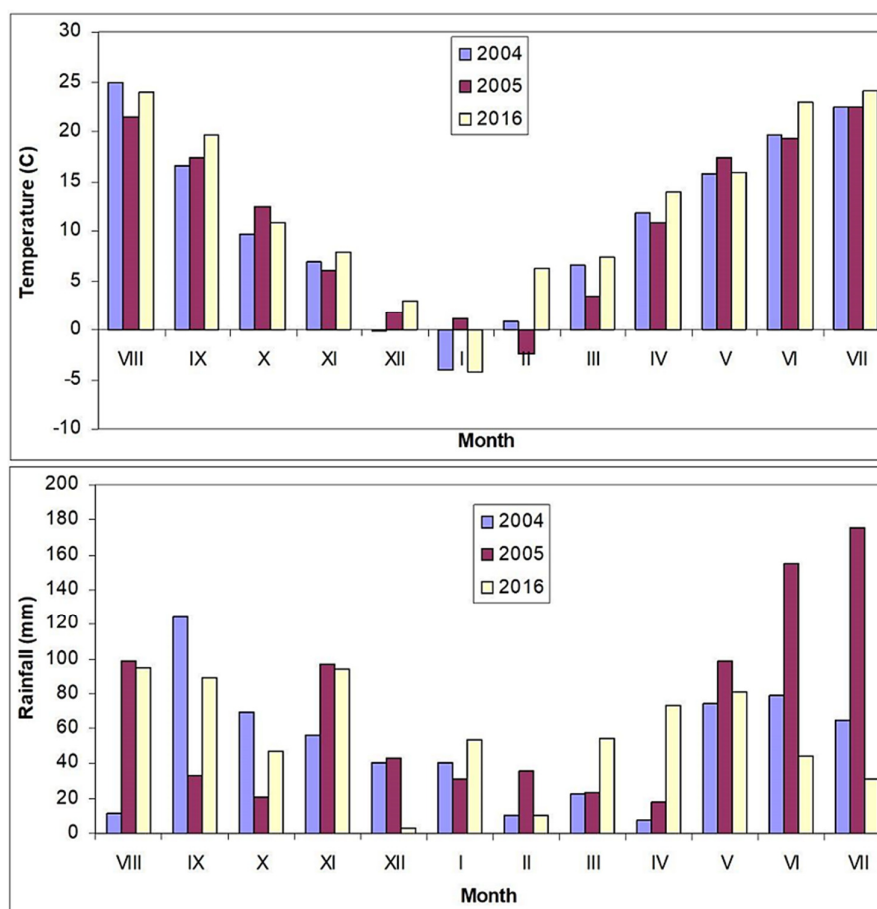


Figure 1. Average monthly temperatures and rainfall in Fundulea during 2004-2005 and 2016

Statistical analysis

The data collected for plant height, ear emergence date and TKW were statistically processed by combined ANOVA and AMMI analysis using MATMODEL Version 3. The means for each trait were compared using Multiple Range Test (Ciulca, 2006). The significance of differences between means was presented based on letters, being considered as significant the differences between means marked with different letters (a, b, c - for genotype \times years comparisons; A, B, C - for years comparisons).

The AMMI stability value (ASV) was calculated as previously described by Purchase *et al.* (2000). It represents the distance to the origin of each line in a two-dimensional space based on interaction principal component axis 1 (IPCA1) and interaction principal component axis 2 (IPCA2) coordinates, considering that lower ASV indicate higher stability.

Results

The combined analysis of the variance based on the AMMI 2 model for wheat genotypes over three years (Table 1) indicates that both genotype and climatic conditions, respectively their interaction had significant effects on TKW. The genotype showed the highest influence (64.24%) on the variability of this trait, followed by genotype \times year interaction (28.46%), amid lower influence (7.3%) of climatic conditions during the study period. The high contribution of the genotype to the variation of the TKW indicates the existence of major differences between wheat genotypes. This model based on the first two principal components fully expresses the effect of genotype \times year interaction on this trait. As such, it's relevant to assess the stability of TKW for wheat genotypes based on the first two principal components.

Table 1. Combined analysis of variance according to the AMMI 2 model for TKW of wheat genotypes

Source of variation	SS	DF	MS	F	SS % ¹
Total	52418	782			
Genotype	30822	86	358.40	42.13**	64.24
Year	3503	2	1751.44	205.87**	7.30
Genotype \times Year	13651	172	79.37	9.33**	28.46(100)
IPCA 1	8634	87	99.24	11.84**	63.24
IPCA 2	5018	85	59.03	7.04**	36.76
IPCA residuals	0	0			
Error	4441	522	8.51		

1- % of model sum of squares for genotype, year, and genotype \times year; ** significant at $P \leq 0.01$.

Based on the data from Table 2 it is noted that generally the climatic conditions of 2005 have favoured the achievement of significantly higher values of this trait, while in 2016 the values of TKW were lower than other years. Under the conditions from 2004 TKW recorded values between 25.6 g at 'DH 6-27' and 66.4 g at 'DH 5-10', amid a relatively symmetrical distribution of genotypes: 9.20% with values over 60 g; 37.93% with values of 50-60 g; 40.23% with values of 40-50 g; 12.64% with values below 40 g. This year the lines 'DH 5-10', 'DH 6-7' and 'DH 5-11' were highlighted, which achieved a TKW of over 65 g.

Given the conditions of 2005 the wheat genotypes showed smaller amplitude of this trait, with the limits from 34.5 g at 'DH 6-27' to 68 g at 'DH 6-7', associated with the following distribution: 13.79% of the lines with TKW over 60 g; 33.33% with values of 50-60 g; 50.57% with values of 40-50 g; 2.30% with TKW below 40 g. The amplitude of TKW in 2016 was close to that recorded in 2005, amid lower values ranging from 27.7 g to 62.6 g 'DH 6-12', and 'G 603/86'. Thus, the distribution of genotypes showed a clear left asymmetry: only 2.30% of genotypes with TKW over 60 g; 28.74% with values of 50-60 g; 43.68% with values of 40-50 g; 25.29% with values below 40 g. Next to line 'G 603/86', only line 'DH 6-31' achieved an TKW over 60 g. Regarding the annual values of TKW (Table 1), it is found that about 20% of the lines did

not show significant variations during the study, while for 5.75% of the lines, the TKW differ significantly from one year to another.

Table 2. Values of TKW for wheat genotypes in 2004, 2005 and 2016

No.	Genotype	2004	2005	2016	Mean	ASV	No.	Genotype	2004	2005	2016	Mean	ASV
P1	G 603/86	60.5a	64.0a	62.6a	62.4	0.404	45	DH 6-20	46.4b	66.2a	43.3b	51.9	1.113
P2	F 132	40.9a	41.2a	33.1b	38.4	0.301	46	DH 6-21	55.2ab	57.8a	51.4b	54.8	0.081
3	DH 5-1	44.2a	48.5a	47.2a	46.6	0.457	47	DH 6-22	45.4a	46.7a	36.4b	42.8	0.437
4	DH 5-2	46.2a	43.6a	43.9a	44.6	0.333	48	DH 6-23	64.8a	60.7a	52.3b	59.3	0.578
5	DH 5-3	44.3a	40.7a	39.7a	41.5	0.265	49	DH 6-24	36.9b	41.4a	37.8b	38.7	0.292
6	DH 5-4	43.0b	48.7a	50.5a	47.4	0.782	50	DH 6-25	52.7a	49.0a	51.4a	51.0	0.459
7	DH 5-5	47.0a	41.9b	48.1a	45.7	0.717	51	DH 6-26	49.6a	44.6ab	39.1b	44.4	0.437
8	DH 5-6	50.7a	49.3ab	44.7b	48.2	0.153	52	DH 6-27	25.6b	34.5a	30.3a	30.1	0.535
9	DH 5-7	47.3a	46.7a	48.7a	47.5	0.510	53	DH 6-28	52.4a	50.4a	39.7b	47.5	0.632
10	DH 5-8	39.0c	61.5a	49.6b	50.0	1.070	54	DH 6-29	54.7b	61.5a	56.3b	57.5	0.344
11	DH 5-9	48.7a	47.9a	34.6b	43.7	0.788	55	DH 6-30	56.6a	52.7a	35.8b	48.4	1.231
12	DH 5-10	66.4a	55.5b	50.2c	57.4	0.800	56	DH 6-31	54.5b	48.0c	60.3a	54.2	1.154
13	DH 5-11	65.6a	60.6b	59.0b	61.7	0.301	57	DH 6-32	61.1a	60.4a	44.2b	55.2	1.014
14	DH 5-12	35.1b	52.9a	48.6a	45.5	1.101	58	DH 6-33	47.5a	55.1a	40.3b	47.6	0.613
15	DH 5-13	50.9a	50.2a	40.2b	47.1	0.511	59	DH 6-34	45.7b	44.1b	51.5a	47.1	0.910
16	DH 5-15	41.7a	44.7a	42.4a	42.9	0.307	60	DH 6-35	51.7a	47.1a	48.3a	49.0	0.382
17	DH 5-17	45.8a	47.2a	44.3a	45.8	0.176	61	DH 6-36	45.4a	46.7a	47.6a	46.5	0.493
18	DH 5-18	53.1a	46.6b	47.8b	49.1	0.413	62	DH 6-37	52.1b	60.7a	54.1b	55.6	0.393
19	DH 5-19	48.6a	51.7a	41.2b	47.1	0.388	63	DH 6-38	55.2a	53.7ab	49.1b	52.6	0.161
20	DH 5-20	56.4a	44.9b	58.5a	53.2	1.161	64	DH 6-39	52.4a	53.1a	45.4b	50.3	0.254
21	DH 5-21	61.9a	66.2a	47.1b	58.4	1.044	65	DH 6-40	46.7b	49.8ab	52.5a	49.7	0.720
22	DH 5-22	37.8c	57.8a	50.2b	48.6	1.068	66	DH 6-41	46.9b	47.1b	54.4a	49.4	0.968
23	DH 5-23	55.9a	56.6a	48.3b	53.6	0.303	67	DH 6-42	57.2a	55.2a	50.7b	54.4	0.186
24	DH 5-25	49.0b	56.9a	38.9b	48.3	0.846	68	DH 6-43	48.9b	54.4a	52.1ab	51.8	0.446
25	DH 5-26	38.3a	40.2a	32.8b	37.1	0.176	69	DH 6-44	57.4a	49.5b	51.8b	52.9	0.486
26	DH 5-27	50.8a	51.5a	39.4b	47.2	0.615	70	DH 6-45	56.6a	50.6b	48.5b	51.8	0.359
27	DH 6-1	58.1a	57.2a	59.5a	58.3	0.518	71	DH 6-46	41.9b	49.3a	52.4a	47.8	0.974
28	DH 6-3	52.8a	52.3a	55.5a	53.5	0.609	72	DH 6-47	57.2a	54.6a	42.6b	51.5	0.772
29	DH 6-4	49.7a	48.8ab	45.0b	47.8	0.107	73	DH 6-48	52.2a	47.6b	48.8b	49.5	0.380
30	DH 6-5	46.7a	44.2a	42.4a	44.4	0.210	74	DH 6-49	48.7b	53.9a	40.3c	47.6	0.572
31	DH 6-6	62.3a	63.5a	57.3b	61.0	0.103	75	DH 6-50	50.0a	50.9a	41.4b	47.4	0.386
32	DH 6-7	66.3a	68.0a	59.3b	64.5	0.290	76	DH 6-51	51.7a	48.3a	47.1a	49.0	0.254
33	DH 6-8	46.2a	44.8a	38.3b	43.1	0.269	77	DH 6-52	52.5a	53.6a	45.3b	50.5	0.286
34	DH 6-9	30.8b	42.9a	32.8b	35.5	0.513	78	DH 6-53	53.5b	62.8a	40.4c	52.2	1.152
35	DH 6-10	45.9a	44.4a	30.9b	40.4	0.832	79	DH 6-55	39.0b	41.1b	46.3a	42.1	0.881
36	DH 6-11	47.0b	53.9a	33.6c	44.8	1.055	80	DH 6-56	43.7b	45.2b	50.6a	46.5	0.864
37	DH 6-12	38.5a	39.8a	27.7b	35.3	0.583	81	DH 6-59	55.8a	57.4a	56.3a	56.5	0.335
38	DH 6-13	45.9a	45.0a	42.9a	44.6	0.182	82	DH 6-60	52.8a	54.1a	53.9a	53.6	0.397
39	DH 6-14	48.5a	46.2ab	43.1b	45.9	0.174	83	DH 6-61	53.1a	53.6a	42.5b	49.7	0.542
40	DH 6-15	41.5b	44.7ab	47.6a	44.6	0.740	84	DH 6-64	58.1a	59.4a	36.3b	51.3	1.497
41	DH 6-16	54.4a	47.4b	48.6b	50.1	0.424	85	DH 6-65	44.6a	44.3a	37.2b	42.0	0.258
42	DH 6-17	30.6b	46.6a	44.0a	40.4	1.087	86	DH 6-66	43.1a	41.5a	33.7b	39.4	0.381
43	DH 6-18	52.3a	51.7a	38.6b	47.5	0.752	87	DH 6-67	49.1b	48.0b	56.3a	51.1	1.003
44	DH 6-19	39.7b	44.9a	38.0b	40.8	0.182	Mean		49.3B	50.7A	45.7C	48.6	

Genotypes LSD5% =2.7; Years LSD5% =0.5; Genotype × Year LSD5% =4.7

Depending on the values of the ASV parameter (Table 2), it is noted that the highest stability of the TKW was presented by the lines: 'DH 6-21', 'DH 6-6', 'DH 6-4', 'DH 5-6', 'DH 6-38', 'DH 6-14', 'DH 5-26', 'DH 5-17', 'DH 5-26', 'DH 6-13', 'DH 6-19'. The high stability of lines 'DH 6-6', 'DH 6-21' and 'DH 6-38' was also associated with high levels of TKW, over 52 g.

Considering that the IPCA1 axis expresses approximately 63.24% of the genotype \times year interaction from Figure 2 it is observed that the highest values of TKW were registered under the conditions from 2004-2005, while in 2016 the values were considerably lower. Also, according to the coordinates of each year it turns out that the conditions from 2016 showed a higher contribution to the genotype \times year interaction, compared to those from 2004-2005.

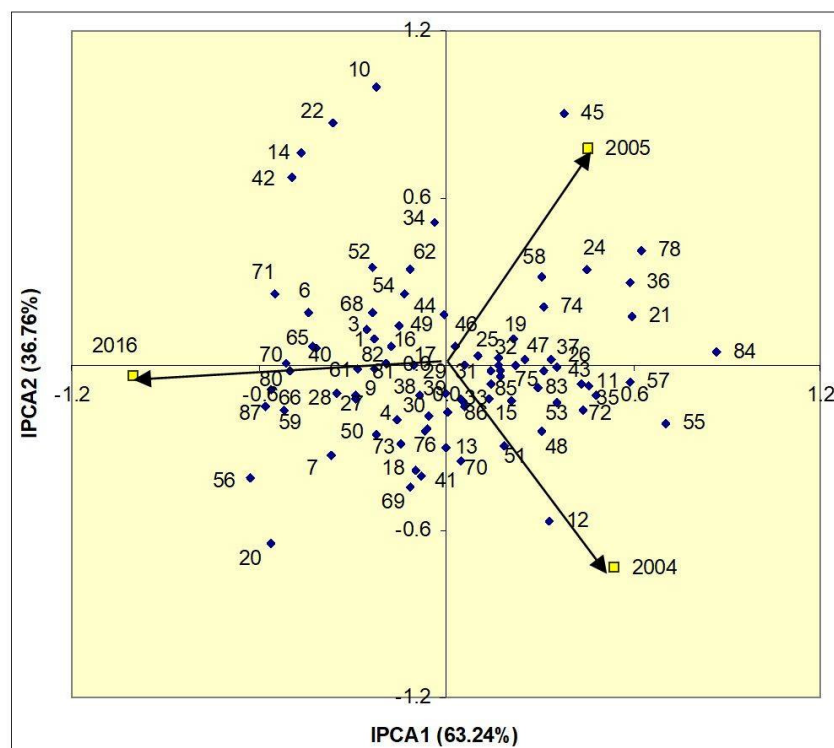


Figure 2. Biplot of interaction for principal component axis (IPCA1 and IPCA2) for TKW of wheat genotypes during 2004-2005 and 2016

The genotypes close to the vector of a given year indicate a strong association with this one. Thus, the line 'DH 6-20' showed a specific adaptation to the conditions of 2005 when it registered a TKW significantly higher with 20-23 g compared to other years. Also, the line 'DH 5-10' showed a strong specific adaptation to the conditions of 2004, achieving a significantly higher TKW by 11-16 g compared to the values of 2005 and 2016.

In case of lines: 'DH 6-21', 'DH 6-6', 'DH 6-4', 'DH 5-6', 'DH 6-38', the close position to the origin indicates high stability of TKW. Considering the distance from the origin, from Figure 2 it is observed that the lines 'DH 6-64', 'DH 6-30' and 'DH 6-53' shows a low stability associated with high values of TKW under the conditions from 2004-2005 and significantly lower values in 2016. For lines 'DH 6-31' and 'DH 5-20', the high instability is associated with a significantly lower TKW in 2005 compared to the values of 2004 and 2016. The lines 'DH 5-8', 'DH 5-12', 'DH 5.22', 'DH 6-17', presented a high interaction with the climatic conditions during the study, recording different values of this trait from year to year, higher in 2005 and lower in 2004.

According to the analysis of the variance based on the first two components of the interaction (Table 3), it is found that all three main sources of variation had a significant influence on plant height of wheat genotypes during the three years. For this trait, the contributions of the three sources of variation are more balanced, so that the variability of plant height was influenced to a high extent by the genotype (49.88%), while the genotype \times year interaction had a lower influence (16.03%). Given that the first two main components fully express the effect of genotype \times year interaction, it turns out that the AMMI2 model is suitable for the assessment of plant height at this set of genotypes.

Table 3. Combined analysis of variance according to the AMMI 2 model for plant height

Source of variation	SS	DF	MS	F	SS % ¹
Total	200941	782			
Genotype	98870	86	1150	6466.58**	49.88
Year	67560	2	33780	220.08**	34.08
Genotype \times Year	31783	172	185	35.37**	16.03 (100)
IPCA 1	25347	87	291	55.92**	79.75
IPCA 2	6435	85	76	14.53**	20.25
IPCA residuals	0	0			
Error	2727	522	5		

1- % of model sum of squares for genotype, year and genotype \times year; ** significant at $P \leq 0.01$.

The climatic conditions during the study showed a strong influence on plant height, causing significant variations from year to year (Table 4). Thus, the conditions from 2016 were significantly more favourable for increasing the plants height compared to those of 2004-2005. The amplitude of the plant height for wheat lines in 2004 was 53.5 cm, ranging from 45.5 cm in 'DH 6-27' to 99 cm in 'DH 6-23'. Thus, the distribution of genotypes showed a low asymmetry with the following structure: only 16.09% of the lines with a plant height below 60 cm; 14.94% with values of 60-70 cm; 32.18% with values of 70-80 cm; 27.59% with values of 80-90 cm; 9.2% of the lines with a plant height over 95 cm.

Under the conditions from 2005 the plants height recorded values between 100 cm in 'DH 6-23' and 46.3 cm in 'DH 6-27', on the background of a left asymmetry of the genotypes distribution: 12.64% of the genotypes with a height of plants over 90 cm; 31.03% with values of 80-90 cm; 27.59% with values of 70-80 cm; 14.94% with values 60-70 cm; 13.79% with a height below 60 cm. Also, and this year lines 'DH 6-23' and 'DH 6-29' showed the highest plant (99-100 cm).

Given the conditions of 2016, there was a considerable increase of plants height for all genotypes. The amplitude of this trait (68 cm) was higher than the previous years, ranging from 59.5 cm in 'DH 6-12' to 127.5 cm in 'DH 5-10' and 'DH 5-20'. The distribution of genotypes in this year showed a left asymmetry, with the following structure: 13.79% of genotypes with a plant height over 110 cm; 22.99% with values of 100-110 cm; 26.44% with values of 90-100 cm; 20.69% with a values of 80-90 cm; 10.34% with values of 70-80 cm; only five genotypes have had a height below 70 cm. Higher values of plant height of over 120 cm were recorded this year by the lines: 'DH 5-10', 'DH 5-20', 'DH 6-29', 'DH 6-21', 'DH 5-11', 'DH 6-23'.

Considering the annual averages of plants height, it is observed that lines 'F 132', 'DH 6-33' and 'DH 6-60' did not registered significant variations during the study. Most lines (78.16%) have better used the conditions from 2016, showing significantly higher plant compared to 2004-2005. About 15% of the lines have a very different reaction to the conditions during the study, recording significant differences of plant height from year to year.

According to ASV parameter (Table 4), it is noted that the highest stability was recorded by the lines: 'DH 6-24', 'G 603/86', 'DH 6-31', 'DH 6-39', 'DH 6-42', 'DH 6-23', 'DH 5-3', 'DH 6-28'. The good stability of these lines was associated with a high size of plants at 'DH 6-23' (106.5 cm), 'G 603/86' (98.17 cm) and 'DH 6-42' (90.22 cm), and with lower values at 'DH 6-28' (65.50 cm) and 'DH 5-3' (72.67 cm), respectively.

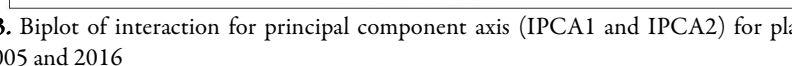
Table 4. Values of plant height for wheat genotypes in 2004, 2005 and 2016

No.	Genotype	2004	2005	2016	Mean	ASV	No.	Genotype	2004	2005	2016	Mean	ASV
P1	G 603/86	92.5b	90.5b	111.5a	98.2	0.098	45	DH 6-20	90.0b	73.5c	104.0a	89.2	0.578
P2	F 132	75.0a	76.3a	77.0a	76.1	1.776	46	DH 6-21	87.5b	88.0b	125.5a	100.3	1.749
3	DH 5-1	73.5b	75.0b	92.0a	80.2	0.187	47	DH 6-22	63.5b	64.7b	76.0a	68.1	0.752
4	DH 5-2	53.0b	52.0b	67.5a	57.5	0.459	48	DH 6-23	99.0b	100.0b	120.5a	106.5	0.128
5	DH 5-3	68.0b	64.0c	86.0a	72.7	0.156	49	DH 6-24	61.0b	63.0b	82.5a	68.8	0.084
6	DH 5-4	88.5b	90.5b	104.0a	94.3	0.502	50	DH 6-25	85.0b	86.3b	110.0a	93.8	0.451
7	DH 5-5	54.0b	55.3b	76.5a	61.9	0.209	51	DH 6-26	79.5b	80.0b	92.5a	84.0	0.672
8	DH 5-6	80.0b	78.5b	100.5a	86.3	0.169	52	DH 6-27	45.5b	46.3b	60.0a	50.6	0.542
9	DH 5-7	71.0b	74.0b	98.5a	81.2	0.616	53	DH 6-28	58.5b	60.5b	77.5a	65.5	0.164
10	DH 5-8	79.5b	80.5b	107.0a	89.0	0.709	54	DH 6-29	96.5b	99.0b	126.5a	107.3	0.880
11	DH 5-9	80.5b	84.5a	84.0ab	83.0	1.760	55	DH 6-30	83.5b	85.5ab	87.5a	85.5	1.614
12	DH 5-10	91.0b	92.0b	127.5a	103.5	1.580	56	DH 6-31	70.0b	71.3b	91.5a	77.6	0.112
13	DH 5-11	77.5c	87.5b	124.0a	96.3	2.137	57	DH 6-32	79.5b	81.5b	109.0a	90.0	0.855
14	DH 5-12	56.5b	57.3b	72.5a	62.1	0.397	58	DH 6-33	82.5a	84.0a	77.5b	81.3	2.462
15	DH 5-13	79.0b	80.3b	95.0a	84.8	0.421	59	DH 6-34	62.5b	64.0b	96.0a	74.2	1.266
16	DH 5-15	67.5c	74.0b	92.5a	78.0	0.261	60	DH 6-35	57.5b	59.0b	92.5a	69.7	1.411
17	DH 5-17	74.0b	72.0b	98.5a	81.5	0.568	61	DH 6-36	72.5c	79.0b	106.5a	86.0	1.088
18	DH 5-18	74.0c	87.5a	80.0b	80.5	2.002	62	DH 6-37	87.5b	88.7b	97.5a	91.2	0.994
19	DH 5-19	61.5b	74.0a	77.0a	70.8	1.057	63	DH 6-38	69.5b	70.7b	101.0a	80.4	1.088
20	DH 5-20	72.5b	61.5c	127.5a	87.2	3.958	64	DH 6-39	76.0b	77.3b	97.5a	83.6	0.112
21	DH 5-21	88.0b	73.5c	104.5a	88.7	0.603	65	DH 6-40	81.5b	83.0b	100.0a	88.2	0.187
22	DH 5-22	57.0c	88.0b	106.0a	83.7	1.632	66	DH 6-41	67.5b	68.7b	103.0a	79.7	1.475
23	DH 5-23	81.0a	56.5b	84.5a	74.0	0.867	67	DH 6-42	83.5b	84.7b	102.5a	90.2	0.122
24	DH 5-25	70.0c	81.0b	87.5a	79.5	0.793	68	DH 6-43	71.0b	72.0b	95.0a	79.3	0.370
25	DH 5-26	62.5b	64.0b	86.0a	70.8	0.298	69	DH 6-44	72.5b	74.0b	96.5a	81.0	0.346
26	DH 5-27	79.5b	80.7b	102.0a	87.4	0.216	70	DH 6-45	80.5b	82.5b	93.5a	85.5	0.743
27	DH 6-1	78.0b	79.3b	101.0a	86.2	0.257	71	DH 6-46	53.0b	54.0b	100.5a	69.2	2.645
28	DH 6-3	85.5b	86.3b	110.0a	93.9	0.426	72	DH 6-47	89.5a	91.0a	89.0a	89.8	2.026
29	DH 6-4	73.5b	75.0b	91.0a	79.8	0.284	73	DH 6-48	73.0b	74.0b	103.5a	83.5	0.999
30	DH 6-5	54.5b	56.5b	79.5a	63.5	0.420	74	DH 6-49	89.5b	90.7b	94.5a	91.6	1.478
31	DH 6-6	90.5b	92.0b	115.5a	99.3	0.443	75	DH 6-50	67.0b	69.5b	91.5a	76.0	0.349
32	DH 6-7	89.5b	90.3b	113.0a	97.6	0.329	76	DH 6-51	55.5b	57.3b	101.5a	71.4	2.460
33	DH 6-8	78.0b	80.0b	96.5a	84.8	0.212	77	DH 6-52	78.5b	80.3b	112.5a	90.4	1.298
34	DH 6-9	70.5b	74.0b	80.0a	74.8	1.155	78	DH 6-53	83.0a	84.7a	68.5b	78.7	3.390
35	DH 6-10	73.0b	76.5b	88.5a	79.3	0.577	79	DH 6-55	58.5b	61.0b	88.0a	69.2	0.832
36	DH 6-11	82.0b	82.0b	105.0a	89.7	0.322	80	DH 6-56	58.5b	59.7b	89.5a	69.2	1.039
37	DH 6-12	75.0a	50.0c	59.5b	61.5	2.353	81	DH 6-59	80.0b	82.3b	91.0a	84.4	0.953
38	DH 6-13	60.0b	61.0b	88.0a	69.7	0.757	82	DH 6-60	92.7a	91.8a	90.0a	91.5	2.126
39	DH 6-14	61.5b	62.7b	76.5a	66.9	0.510	83	DH 6-61	92.0b	93.0b	105.5a	96.8	0.647
40	DH 6-15	71.5b	74.0b	80.0a	75.2	1.203	84	DH 6-64	76.5b	77.7b	88.0a	80.7	0.849
41	DH 6-16	83.5b	72.0c	99.5a	85.0	0.423	85	DH 6-65	80.0b	81.3b	89.0a	83.4	1.099
42	DH 6-17	64.0c	87.5b	96.5a	82.7	0.682	86	DH 6-66	59.0b	57.5b	87.5a	68.0	0.928
43	DH 6-18	80.0b	74.0c	99.0a	84.3	0.305	87	DH 6-67	85.0b	86.7b	102.0a	91.2	0.340
44	DH 6-19	56.0c	61.5b	66.5a	61.3	1.159	Mean		74.3C	75.5B	94.6A		

Genotypes LSD5% =2.1; Years LSD5% =0.4; Genotype × Year LSD5% =3.7

The length and position of the vectors for lines 'DH 5-22' and 'DH 6-17' indicate a high instability of plant height, associated with significant variations (over 20 cm) from year to year, with the highest values in 2016 and the lowest in 2004. The line 'DH 6-53' showed a high stability of plant height under the conditions from 2004-2005 and a significantly lower value in 2016. A special reaction is also observed in the case of the line 'DH 5-23' which registered a very small variation of plants height in 2004 and 2016, associated with significantly higher values compared to 2005.

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The combined analysis of the variance based on the AMMI 2 model for wheat genotypes over three years (Table 5) indicates that both genotype and climatic conditions, respectively their interaction had significant effects on ear emergence. The climatic conditions showed the highest influence (62.22%) on the variability of this trait, while the genotype (19.94%) and the genotype \times year interaction (17.84%) had similar influences. This model based on the first two principal components fully expresses the effect of genotype \times year interaction on the emergence precocity, being appropriate for evaluating the stability of this trait in wheat lines.

Table 5. Combined analysis of variance according to the AMMI 2 model for ear emergence date

Source of variation	SS	DF	MS	F	SS % ¹
Total	30211	782			
Genotypes	5980	86	69.53	163.86**	19.94
Years	18658	2	9329.00	21985**	62.22
Genotypes x Years	5351	172	31.11	73.32**	17.84 (100)
IPCA 1	3812	87	43.82	104.87**	71.23
IPCA 2	1540	85	18.11	43.35**	28.77
IPCA residuals	0	0			
Error	221	522	0.42		

¹% of model sum of squares for genotype, year and genotype \times year; ** significant at $P \leq 0.01$.

According to the data from Table 6 it is observed that generally the climatic conditions from 2016 have caused a significant precocity of ear emergence, while in 2005 there was a significant delay. Under the conditions from 2004, the ear emergence date registered amplitude of about 14 days. In this sense the earliest genotypes were: 'F 132', 'DH 5-21', 'DH 6-31', which emerged around May 11, while the line 'DH 6-45' which emerged around May 25 was the latest. The precocity of ear emergence in this year showed a symmetrical distribution of lines: 3.45% of the lines emerged in May 10 to 11; 41.38% emerged in May 11 to 16; 45.98% emerged in May 16 to 21; 9.20% of the lines emerged after May 21.

Under the conditions from 2005 there was a delay of ear emergence, amid amplitude of 10 days between the earliest lines ('F 132' and 'DH 5-1') which emerged on May 12 and the lines 'DH 5-4', 'DH 5-8', 'DH 5-10', 'DH 5-21', 'DH 6-20' which emerged on May 22, respectively. Thus, 29.89% of the lines emerged until May 16, while 59.77% emerged in May 16 to 21, and 10.34% emerged after May 21.

Amid a precocity compared to previous years, the ear emergence amplitude in 2016 was higher (17 days), ranging between April 28 for 'DH 6-33' and 'DH 6-46' lines, and May 15 for 'DH 5-4', 'DH 5-8' and 'DH 5-22' lines. Thus, 13.79% of the lines emerged before May 1; 3.45 % in May 1; 37.93 % in May 2 to 5; 16.09 % in May 6 to 11; 28.74% after May 11.

Compared to the parental forms, during the study 62.5% of the 'DH 5' lines presented a later ear emergence than the parents mean, and four lines were later than both parents, without being registered earlier lines than 'F.132'. In the case of the 'DH 6' series, 49.18% of the lines showed a later ear emergence than the parents mean, in this respect the lines 'DH 6-3' and 'DH 6-20' were later than paternal parent 'G 603/86', while only the line 'DH 6-31' was earlier than maternal parent 'F.132'.

Regarding the annual values of this trait, it is found that about 11.5% of the lines did not show significant variations of ear emergence in 2004-2005, while about 90% of the lines have shown significant differences of this trait from one year to another.

According to values of ASV parameter (Table 6) it is noted that the highest stability of ear emergence was presented by the lines: 'DH 6-8', 'DH 6-67', 'DH 6-30', 'DH 6-19', 'DH 5-13', 'DH 6-28', 'DH 6-4', 'DH 6-13'. The high stability of the lines: 'DH 6-30', 'DH 6-13', 'DH 6-28', 'DH 6-67', 'DH 6-19', has been associated with a good precocity, an ear emergence around May 9-10, respectively.

Table 6. Ear emergence date for wheat genotypes in 2004, 2005 and 2016

No.	Genotype	2004*	2005*	2016*	Mean*	ASV	No.	Genotype	2004*	2005*	2016*	Mean*	ASV
P1	G 603/86	16.7b	20.3a	12.0c	16.3	1.007	45	DH 6-20	20.3a	21.0a	12.3b	17.9	0.431
P2	F 132	9.7b	11.0a	2.0c	7.6	0.481	46	DH 6-21	18.7a	11.3c	12.7b	14.2	1.529
3	DH 5-1	17.3a	11.0b	0.7c	9.7	1.255	47	DH 6-22	12.3b	17.7a	9.3c	13.1	1.181
4	DH 5-2	20.7a	16.0b	6.0c	14.2	0.814	48	DH 6-23	11.7b	19.0a	11.0b	13.9	1.717
5	DH 5-3	17.3a	15.3b	3.7c	12.1	0.725	49	DH 6-24	17.3a	12.0b	-2.0c	9.1	1.727
6	DH 5-4	18.7b	21.0a	14.0c	17.9	1.058	50	DH 6-25	21.3a	17.7b	7.0c	15.3	0.807
7	DH 5-5	11.3b	14.7a	3.7c	9.9	0.364	51	DH 6-26	15.3b	19.7a	12.3c	15.8	1.207
8	DH 5-6	18.7a	12.0b	-1.7c	9.7	1.805	52	DH 6-27	13.0b	15.7a	2.7c	10.4	0.321
9	DH 5-7	16.3a	16.0a	2.7b	11.7	0.775	53	DH 6-28	10.5b	13.0a	1.5c	8.3	0.210
10	DH 5-8	17.7b	21.0a	14.0c	17.6	1.223	54	DH 6-29	20.0a	19.3a	9.3b	16.2	0.284
11	DH 5-9	15.3b	19.7a	11.3c	15.4	1.106	55	DH 6-30	11.3a	11.7a	1.3b	8.1	0.080
12	DH 5-10	16.3b	21.0a	12.7c	16.7	1.094	56	DH 6-31	9.7b	13.7a	-2.0c	7.1	0.621
13	DH 5-11	15.7b	20.3a	11.7c	15.9	1.084	57	DH 6-32	17.0a	13.0b	9.7c	13.2	0.908
14	DH 5-12	16.3b	20.7a	12.0c	16.3	1.007	58	DH 6-33	16.3b	18.7a	-3.0c	10.7	2.079
15	DH 5-13	13.7b	16.0a	4.0c	11.2	0.186	59	DH 6-34	13.0a	11.7b	1.0c	8.6	0.449
16	DH 5-15	12.7b	17.3a	3.3c	11.1	0.390	60	DH 6-35	14.7a	15.7a	2.7b	11.0	0.478
17	DH 5-17	14.3b	18.0a	3.7c	12.0	0.480	61	DH 6-36	19.3a	15.3b	4.3c	13.0	0.985
18	DH 5-18	10.7b	14.3a	-0.3c	8.2	0.592	62	DH 6-37	20.0a	16.3b	-2.0c	11.4	2.360
19	DH 5-19	13.7a	11.3b	0.0c	8.3	0.629	63	DH 6-38	17.3a	12.3b	-1.7c	9.3	1.738
20	DH 5-20	19.0a	14.0b	12.3c	15.1	1.225	64	DH 6-39	14.3b	15.7a	2.3c	10.8	0.440
21	DH 5-21	9.7c	21.0a	13.7b	14.8	2.519	65	DH 6-40	13.7b	16.0a	0.0c	9.9	0.867
22	DH 5-22	15.3a	11.3c	14.0b	13.6	2.127	66	DH 6-41	11.3b	15.3a	-2.3c	8.1	1.098
23	DH 5-23	13.7b	17.0a	5.3c	12.0	0.378	67	DH 6-42	16.0a	15.7a	0.7b	10.8	1.167
24	DH 5-25	12.3b	15.0a	4.0c	10.4	0.275	68	DH 6-43	14.3b	18.3a	1.0c	11.2	0.999
25	DH 5-26	16.3b	18.7a	8.7c	14.6	0.463	69	DH 6-44	17.3a	12.0b	1.0c	10.1	1.163
26	DH 5-27	14.7b	17.3a	7.3c	13.1	0.549	70	DH 6-45	23.3a	14.0b	1.3c	12.9	2.150
27	DH 6-1	15.3b	16.7a	7.7c	13.2	0.379	71	DH 6-46	11.7b	14.7a	-2.7c	7.9	1.011
28	DH 6-3	21.0a	19.3b	11.0c	17.1	0.269	72	DH 6-47	18.3a	15.7b	2.3c	12.1	1.095
29	DH 6-4	13.7b	16.3a	3.7c	11.2	0.218	73	DH 6-48	15.0b	16.3a	12.0c	14.4	1.375
30	DH 6-5	14.7b	16.0a	3.3c	11.3	0.274	74	DH 6-49	20.3a	18.7b	-1.3c	12.6	2.366
31	DH 6-6	15.7b	17.7a	10.3c	14.6	0.955	75	DH 6-50	18.7a	14.7b	-1.7c	10.6	1.868
32	DH 6-7	14.3b	17.3a	9.7c	13.8	0.936	76	DH 6-51	12.3b	14.3a	1.7c	9.4	0.321
33	DH 6-8	15.7a	16.0a	5.7b	12.4	0.020	77	DH 6-52	11.3b	15.0a	11.0b	12.4	1.865
34	DH 6-9	15.3b	18.3a	11.0c	14.9	1.038	78	DH 6-53	13.0b	19.7a	4.0b	12.2	0.586
35	DH 6-10	16.0a	14.3b	3.0c	11.1	0.541	79	DH 6-55	11.3b	16.0a	-2.0c	8.4	1.046
36	DH 6-11	17.7b	19.7a	12.3c	16.6	0.977	80	DH 6-56	14.7b	16.0a	0.0c	10.2	0.998
37	DH 6-12	16.7b	18.0a	9.7c	14.8	0.585	81	DH 6-59	12.3b	15.0a	2.0c	9.8	0.286
38	DH 6-13	11.3b	12.7a	0.7c	8.2	0.232	82	DH 6-60	13.3b	15.7a	3.0c	10.7	0.230
39	DH 6-14	14.7b	16.0a	8.7c	13.1	0.793	83	DH 6-61	16.7b	20.0a	11.7c	16.1	0.921
40	DH 6-15	18.3a	17.3a	7.7b	14.4	0.232	84	DH 6-64	17.3a	16.0b	4.0c	12.4	0.638
41	DH 6-16	17.3a	18.0a	10.3b	15.2	0.641	85	DH 6-65	16.0b	19.3a	13.0c	16.1	1.265
42	DH 6-17	17.7a	15.0b	13.0c	15.2	1.389	86	DH 6-66	14.3b	16.0a	3.3c	11.2	0.375
43	DH 6-18	18.3a	11.7b	12.3b	14.1	1.529	87	DH 6-67	12.7a	13.3a	2.3b	9.4	0.039
44	DH 6-19	12.7b	14.3a	2.7c	9.9	0.158		Mean	15.4B	16.1A	5.5C		

* Days after 1 May Genotypes LSD5% = 0.6; Years LSD5% = 0.1; Genotype × Year LSD5% = 1.1

Based on the biplot from Figure 4 and considering that the IPCA1 axis expresses about 71.23% of the genotype \times year interaction, it can be seen that the lines showed a considerable delay of the ear emergence in 2004-2005 compared with 2016. Also, according to the coordinates of each year it turns out that the conditions from 2016 showed a higher contribution to the genotype \times year interaction, while the conditions from 2005 had a smaller effect on the respective interaction.

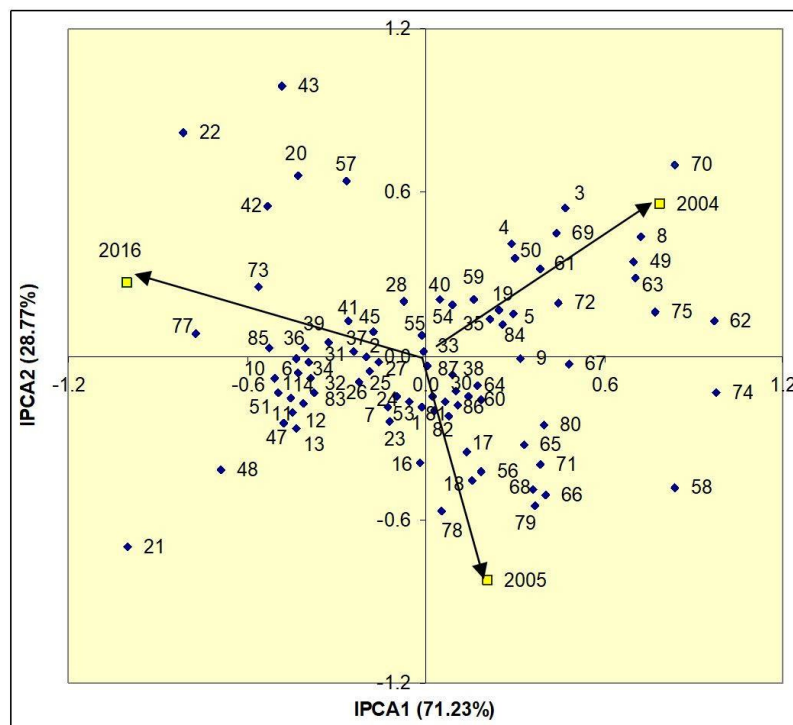


Figure 4. Biplot of interaction for principal component axis (IPCA1 and IPCA2) for ear emergence of wheat genotypes during 2004-2005 and 2016

The close position to the origin in case of lines: 'DH 6-21', 'DH 6-6', 'DH 6-4', 'DH 5-6', 'DH 6-38', indicates high stability of ear emergence (Figure 4). Considering the distance from the origin, it is observed that the lines: 'DH 6-18', 'DH 5-22', 'DH 5-20', 'DH 6-17', 'DH 6-32', shows a low stability associated with a late ear emergence. For lines 'DH 6-37', 'DH 6-49', 'DH 6-33', 'DH 6-50' and 'DH 6-45', the high instability is associated with an earlier ear emergence in 2016 compared to 2004-2005. The line 'DH 5-21', presented a high interaction with the climatic conditions during the study expressing a particular reaction, namely a considerably earlier ear emergence in 2004 compared to 2005 and 2016.

Discussion

The studied genotypes are suitable for this work, considering that DH lines obtained via maize hybridization are the most useful for research studies and for the breeding of new wheat cultivars. The probability of segregation modification and loss of desirable genotypes in these populations is significantly lower than in anther-derived populations (Adamski *et al.*, 2014). DH populations were efficiently used in other studies (Zhang *et al.*, 2009; Gegas *et al.*, 2010; Heidari *et al.*, 2011; Ganey *et al.*, 2014; Zhang *et al.*, 2014; Griffith *et al.*, 2015; Guo *et al.*, 2015; Zhang *et al.*, 2016; Liu *et al.*, 2018) for the analysis of the relationships among different yield components in wheat.

The environmental conditions of the three years had an important contribution on the genotype \times year interaction, which showed also a higher influence on TKW. The variability of plant height and ear emergence data was also affected to a similar extent by this interaction. Significant effect of GE interaction on TKW has been previously reported by other studies (Gómez-Becerra *et al.*, 2010; Tayyar, 2010; Sharma *et al.*, 2013; Khazratkulova *et al.*, 2015; Krishnappa *et al.*, 2019). The high contribution of the genotype to the variation of TKW and plant height indicates that for the studied DH lines these two traits are most stable compared with ear emergence. Van Frank *et al.* (2020), found a close stability of breeding populations and commercial varieties for plant height and TKW. Barbu *et al.* (2018) reported that the reduced plant height and its stability in wheat were not correlated with the genetic background of this trait.

Compared to the parental forms, during the study 20.83% of the DH 5 lines presented an average value of TKW higher than the parents mean, 4.16% were lower than both parents, without being registered lines with values of this trait higher than the superior parent ('G 603/86'). In the case of the 'DH 6' series, 34.34% of the lines achieved a TKW higher than the parents mean, in this respect the line 'DH 6-7' which exceeded the best parent, was highlighted. Also, 4.92% of 'DH 6' lines were lower than both parents. Significant increases of TKW compared to parents mean were also reported by other studies: 15.92 % by Bao *et al.* (2009); 5.1% by Simmonds *et al.* (2014). Considering that previously studies (Wang *et al.*, 2012; Zhang *et al.*, 2012) reported in wheat a linear correlation between TKW and favourable alleles, the DH lines with high values of TGW can be used as parents for crosses to ensure pyramiding the maximum number of favourable alleles in a valuable genetic background.

The grain yield is strongly influenced by the plant height, which can be used as a selection criterion for improving grain yield in wheat (Mohammadi *et al.*, 2012). In comparison with the parental forms, it is found that most of the lines (80.84%) of the 'DH 5' series recorded a plant height below the parents mean, or even lower than both parents in the case of 29.16% of the lines. Only the line 'DH 5-10' showed a higher value to both parents. In the case of the 'DH 6' series, 29.5% of the lines achieved a lower plant height than both parents, 70.5% had an intermediate value to parental forms while four lines were superior to both parents. A large variation of plant height in DH populations compared to parents mean was also reported by other studies (Inagaki *et al.*, 1998; Heidari *et al.*, 2012). Likewise, El-Hennawy *et al.* (2011) observed in five DH populations, several transgressive segregants for plant height in both directions. As well, Wu *et al.* (2010) reported that the plant height of wheat DH lines showed continuous variation and transgressive segregation in different growth stages and environments. Wheat DH populations are an effective material to be used for studying the molecular genetic basis of plant height (Li *et al.*, 2010; Wu *et al.*, 2010; Zhang *et al.*, 2011; Heidari *et al.*, 2012).

As against the parental forms, during the study 62.5% of the 'DH 5' lines presented a later ear emergence than the parents mean and four lines were later than both parents, without being registered earlier lines than 'F.132'. In the case of the 'DH 6' series, 49.18% of the lines showed a later ear emergence than the parents mean, in this respect the lines 'DH 6-3' and 'DH 6-20' were later than paternal parent 'G 603/86', while only the line 'DH 6-31' was earlier than maternal parent 'F.132'. These results are in agreement with those of Kuchel *et al.* (2006) who reported that more than 50% of the individuals from a wheat DH population emerge later than both parents under an average daily temperature during the duration of heading of 12.7 °C, while under an average daily temperature of an average daily temperature of 16.3 °C more than 70% of the DH population were later than the parents. In another study Lantos *et al.* (2019) reported that the DH lines showed close or a little later ear emergence in comparison with their parent. Also, an important variation of heading date up to 13 days, in DH lines compared with the parents mean was found by Heidari *et al.* (2012). A large genetic variability for total time to anthesis and duration of pre-anthesis phases in DH populations was observed by Borràs-Gelonch *et al.* (2012), Coleman *et al.* (2001) and Rebetzke *et al.* (2008). These transgressive segregations, indicates the existence of multiple, independent alleles at different loci between the parents of the DH populations (Borràs-Gelonch *et al.*, 2012).

Taking into account that grain yield can be increased by growing varieties which heading time allows to avoid different stresses during grain-filling phase (Kamran *et al.*, 2014), and considering that several QTLs for grain shape, size and yield were detected, near the loci for photoperiod sensitivity (Maphosa *et al.*, 2014), some of the studied DH lines can be used for simultaneous improvements of different target traits.

Conclusions

Based on the performed results and analyses, were highlighted the lines: 'DH 6-7' (64.53 g), 'DH 5-11' (61.72 g), 'DH 6-6' (61.02 g), 'DH 6-29' (57.48 g), 'DH 6-59' (56.49 g), 'DH 6-21' (54.78 g), 'DH 6-42' (54.37 g), which show high and stable values of TKW, associated with a plants height of approximately 85-100 cm and an ear emergence from May 11 to 17, under some climatic conditions similar to the period of study. The line 'DH 5-10' showed a specific adaptation to less favourable wheat conditions represented by a lower level of precipitation during the spring growth period in 2004, achieving a significantly higher TKW by 11-16 g compared to the values of the others years. The lines 'DH 6-31', 'DH 6-67' and 'DH 6-56', have shown a high potential to effectively exploit the favourable conditions in spring, achieving high levels of TKW amid an earlier ear emergence between April 29 and May 4. The above-mentioned doubled haploids lines can be considered as promising genotypes for using in wheat breeding programs in order to improve yield performances under temperate continental climate conditions.

Authors' Contributions

Conceptualization: SC, AG and AC; Methodology: AG; Data analysis and interpretation: SC and AC; Writing- original draft: SC, AG and AC; Writing-review and editing: SC, AG and AC. All authors read and approved the final manuscript.

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Conflict of Interests

The authors declare that there are no conflicts of interest related to this article.

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