

Determining effective yield components on seed yield by Path and Biplot Analyses in Bread Wheat (*Triticum aestivum* L.)

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Abstract: The purpose of this study was to determine the effective yield factors (plant height, spike number/m², spike height, spikelet number, seed number/spike, seed weight/spike, 1000 seed weight, test weight, harvest index and protein content) on grain yield in bread wheat genotypes by statistical methods. Results denoted that spike number/m², seed number/spike, seed weight/spike and test weight are important components; Müfitbey, Altay, Nacibey, Es 26 were high potential and stabile genotypes.

Keywords: Bread wheat, genotype, path and biplot analyses, effectiveness, yield components.

Özet: Bu çalışmanın amacı, ekmeçlik buğday genotiplerinde tane verimi üzerinde etkili olan verim faktörlerinin (bitki boyu, başak sayısı/m², başak yükseklięi, başakçık sayısı, tohum sayısı/başak, tohum aęırlığı/başak, 1000 tohum aęırlığı, tohum aęırlığı, hasat indeksi ve protein içerięi) istatistiksel yöntemlerle belirlenmesidir. Sonuçlara göre, başak sayısı/m², tohum sayısı/başak, tohum aęırlığı/başak ve tohum aęırlığı önemli bileşenlerdir; Müfitbey, Altay, Nacibey, Es 26 genotipleri yüksek potansiyelli ve stabildir.

Anahtar Kelimeler: Ekmeçlik buğday, genotip, path ve biplot analizleri, etkinlik, verim unsurları.

INTRODUCTION

Wheat is a cultivated plant of great importance in human nutrition and has the widest adaptation ability among cereals. Wheat flour, which is grown in more than a hundred countries in the world, is used as a raw material in the production of many products in many cultures and geographies due to its unique structure (Atlı,1992). Again, wheat is the most cultivated plant in the world, together with corn, in terms of planting area and production (Anon., 2019). In Türkiye, wheat ranks first among cereals with approximately 8 million hectares of acreage, approximately 20 million tons of production and 2,44 t/ha yield (Anon., 2010). Wheat, which is the basic food of approximately 35% of the world's population, provides 20% of the calories taken from food all over the world. 53% of daily calorie consumption in Turkey is provided by wheat (Kün, 1983). Wheat is the staple food of approximately 35% of the world's population due to its good nutritional raw material, wide adaptation limit, ease of production, transportation, storage and processing (Kün, 1988; Yang and Baker, 1991; Tripathi et al., 2003; Yanbeyi and Sezer, 2006).

Due to the increasing human population, the need for grain is also increasing for people and animals to be sufficiently nourished. However, due to reasons such as erosion, natural disasters, unplanned urbanization, etc., agricultural areas are gradually decreasing. Due to the increasing food need and decreasing agricultural areas, we need to increase the agricultural products obtained from a unit area for people to have sufficient and balanced nutrition. Increasing wheat production in the world is only possible by developing and introducing production genotypes that are high yielding, high quality, resistant to biotic and abiotic stresses, and stable in terms of these features. In developing genotypes in terms of these features, it is possible by carrying out the breeding studies carried out effectively and continuously. The effective execution of breeding studies can only be revealed by determining the yield components affecting high yield, and effective studies in the field of breeding and agronomy (happening literature). In many studies conducted on wheat in the world, yield components affecting yield have been determined (Ryan et al., 1991; Prystupa et al., 2003). In many studies, it has been determined that the factors such as number of spikes per square meter, number and weight of grains per spike, harvest index, test weight are quite effective on yield (Mehrotra et al., 1967; Tripathi et al., 2003). By determining the effect levels of such yield factors, more effective progress will be made in wet and agronomy programs and high quality and high yield genotypes will be developed (Prystupa et al., 2003; Tripathi et al., 2003). In this study, statistical methods were used to determine the effective yield factors (plant height, spike number/m², spike height, spikelet number, seed number/spike, seed weight/spike, 1000 seed weight, test weight, harvest index and protein content) on grain yield in bread wheat genotypes and for this purpose, effective yield factors on grain yield were determined using path and biplot analyses methods.

MATERIALS AND METHODS

This research was carried out in 2020-2021 and 2021-2022 in the experimental area of Eskişehir Osmangazi University Faculty of Agriculture under arid conditions. In the 2020-2021 year when the experiment was carried out, a total of 363,7 mm of precipitation and the average temperature was 10,4 ° C. In 2021-2022, a total of 385,7 mm of precipitation and the average temperature was 9,4 ° C. This amount of precipitation was determined to be slightly higher than the long-term average (361,45 mm). Again, the average temperature in both years was close to the long-term average (10.1 ° C). The experimental site soils are salt-free, contain 1,7% organic matter and 5,24% lime, and are clayey loamy and slightly alkaline in structure.

The experiment was carried out according to a randomized block design under arid conditions. Sowing was done on 15 October with a plot drill and the sowing density was applied as 0,22 t/ha. In the experiment, 0,06 t/ha N and 0,06 t/ha P₂O₅ were applied to the wheat genotypes as base fertilization together with sowing. Müfitbey, Altay, Nacibey, Es 26, Reis, Bayraktar, Bezostaja, Sönmez and Alpu genotypes were used as bread wheat

genotypes in the experiment. Plant height, spike number/m², spike height, spikelet number, seed number/spike seed weight/spike, 1000 seed weight, test weight, harvest index, protein content and seed yield. Path and biplot analyses of the average data obtained from the genotypes in the experiment were performed using Minitab 17, Tarist and Lisrel software programs.

RESULTS AND DISCUSSION

In wheat breeding, it is very important to evaluate the genetic material in terms of yield and quality components as well as yield in order to improve high yielding and high quality, stable in terms of quality and yield, and resistant to diseases. In this context, ten wheat genotypes used in the study; plant height, spike number/m², spike height, spikelet number, seed number/spike, seed weight/spike, 1000 seed weight, test weight, harvest index and protein content were evaluated and effective yield components on grain yield were determined. The mean, maximum and minimum values of yield components examined on ten wheat genotypes tested in the study are given in Table 1.

Table 1. Mean, maximum and minimum values of the yield components.

Variable	Mean	Minimum	Maximum	Variable	Mean	Minimum	Maximum
Plant Height	85,68±3,79	67,84	103,81	1000 Seed Weight	37,41±1,46	24,43	40,13
Spike Number/m ²	349,60±19,62	281,66	462,85	Test Weight	79,23±0,851	71,76	80,99
Spike Height	7,90±0,11	8,41	9,51	Harvest Index	41,54±1,36	29,71	46,54
Spikelet Number	13,64±0,31	11,37	14,04	Protein Content	12,34±0,39	10,26	14,09
Seed Number/Spike	31,46±1,31	24,43	36,76	Seed Yield	359,72±20,56	230,68	428,75
Seed Weight/Spike	2,03±0,05	1,89	2,35	±			

Correlation analysis between yield components is given in Table 1, path and biplot analysis revealing the components effective on yield are given in Table 1 and Figure 1a and b. Spike number/m², seed number/spike, seed weight/spike, test weight and harvest index had important relationship at %/1% with seed yield (Table 2). Path analysis revealed that spike number/m², spikelet number and seed number/spike via plant height; seed number/spike, seed weight/spike, test weight and harvest index via spike number/m²; spike number/m², seed number/spike, seed weight/spike, harvest index via spike height; seed number/spike, harvest index, protein content

Table 2. Correlation analysis between yield elements and path and biplot analyses revealing the components that are effective on yield.

	Seed Yield	Plant Height	Spike Number/m ²	Spike Height	Spikelet Number	Seed Number/Spike
Plant Height	0.572ns					
Spike Number/m ²	0.778**	0.486ns				
Spike Height	-0.383ns	0.289ns	-0.579ns			
Spikelet Number	0.554ns	0.707*	0.444ns	0.100ns		
Seed Number/Spike	0.770**	0.324ns	0.661*	-0.559ns	0.661*	
Seed Weight/Spike	0.763*	0.158ns	0.638*	-0.639*	0.293ns	0.566ns
1000 Seed Weight	0.032ns	0.245ns	0.406ns	-0.375ns	0.328ns	0.192ns
Test Weight	0.761*	0.616ns	0.477ns	-0.128ns	0.634*	0.569ns
Harvest Index	0.638*	-0.038ns	0.610ns	-0.732*	0.347ns	0.839**
Protein Content	0.290ns	0.572ns	0.612ns	-0.158ns	0.612ns	0.438ns
	Seed Weight/Spike	1000 Seed Weight	Test Weight	Harvest Index		
1000 Seed Weight	0.293ns					
Test Weight	0.701*	0.169ns				
Harvest Index	0.649*	0.052ns	0.384ns			

Protein Content		0.071ns		0.698*		0.193ns		0.072ns	
Dependent Variable: Seed Yield									
For Plant Height					For Spike Number/m ²				
Variables	Path Coef.	%	Variables	Path Coef.	%				
Direct Effect	0.3393	19.6554	Direct Effect	0.5422	19.2812				
Indirect Effects		Indirect Effects		Indirect Effects					
Via	Path Coef.	%	Via	Path Coef.	%				
Spike Number/m ²	0.2634	15.2602	Plant Height	0.1648	5.8612				
Spike Height	-0.0600	3.4787	Spike Height	0.1204	4.2812				
Spikelet Number	0.1799	10.4244	Spikelet Number	0.1131	4.0200				
Seed Number/Spike	0.2537	14.6988	Seed Number/Spike	0.5181	18.4229				
Seed Weight/Spike	0.0832	4.8216	Seed Weight/Spike	0.3360	11.9494				
1000 Seed Weight	-0.0786	4.5543	1000 Seed Weight	-0.1304	4.6386				
Test Weight	-0.1671	9.6787	Test Weight	-0.1293	4.5981				
Harvest Index	0.0294	1.7045	Harvest Index	-0.4673	16.6165				
Protein Content	-0.2714	15.7234	Protein Content	-0.2905	10.3309				

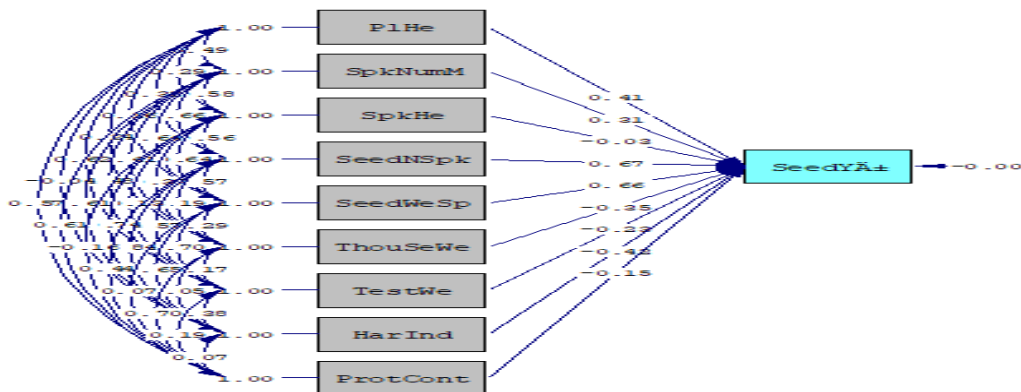
For Spike Height					For Spikelet Number				
Variables	Path Coef.	%	Variables	Path Coef.	%				
Direct Effect	-0.2078	9.3962	Direct Effect	0.2546	11.2583				
Indirect Effects		Indirect Effects		Indirect Effects					
Via	Path Coef.	%	Via	Path Coef.	%				
Plant Height	0.0981	4.4351	Plant Height	0.2398	10.6010				
Spike Number/m ²	-0.3142	14.2108	Spike Number/m ²	0.2407	10.6437				
Spikelet Number	0.0254	1.1489	Spikelet Number	-0.0207	0.9165				
Seed Number/Spike	-0.4381	19.8124	Seed Number/Spike	0.5184	22.9196				
Seed Weight/Spike	-0.3365	15.2197	Seed Weight/Spike	0.1541	6.8153				
1000 Seed Weight	0.1205	5.4480	1000 Seed Weight	-0.1052	4.6501				
Test Weight	0.0347	1.5710	Test Weight	-0.1718	7.5978				
Harvest Index	0.5608	25.3622	Harvest Index	-0.2657	11.7459				
Protein Content	0.0751	3.3956	Protein Content	-0.2907	12.8519				

For Seed Number/Spike					For Seed Weight/Spike				
Variables	Path Coef.	%	Variables	Path Coef.	%				
Direct Effect	0.7843	27.0315	Direct Effect	0.5269	22.0151				
Indirect Effects		Indirect Effects		Indirect Effects					
Via	Path Coef.	%	Via	Path Coef.	%				
Plant Height	0.1098	3.7832	Plant Height	0.0536	2.2398				
Spike Number/m ²	0.3582	12.3454	Spike Number/m ²	0.3459	14.4519				
Spikelet Number	0.1160	3.9997	Spikelet Number	0.1327	5.5454				
Spikelet Number	0.1683	5.8008	Spikelet Number	0.0745	3.1131				
Seed Weight/Spike	0.2984	10.2859	Seed Number/Spike	0.4443	18.5642				
1000 Seed Weight	-0.0618	2.1293	1000 Seed Weight	-0.0940	3.9282				
Test Weight	-0.1542	5.3134	Test Weight	-0.1901	7.9422				
Harvest Index	-0.6424	22.1418	Harvest Index	-0.4974	20.7825				
Protein Content	-0.2080	7.1691	Protein Content	-0.0339	1.4177				

For 1000 Seed Weight					For Test Weight				
Variables	Path Coef.	%	Variables	Path Coef.	%				
Direct Effect	-0.3211	21.2920	Direct Effect	-0.2710	12.4168				
Indirect Effects		Indirect Effects		Indirect Effects					
Via	Path Coef.	%	Via	Path Coef.	%				
Plant Height	0.0831	5.5092	Plant Height	0.2092	9.5824				
Spike Number/m ²	0.2203	14.6092	Spike Number/m ²	0.2587	11.8524				
Spikelet Number	0.0780	5.1693	Spikelet Number	0.0266	1.2200				
Spikelet Number	0.0834	5.5314	Spikelet Number	0.1615	7.3968				
Seed Number/Spike	0.1509	10.0078	Seed Number/Spike	0.4461	20.4385				
Seed Weight/Spike	0.1543	10.2295	Seed Weight/Spike	0.3695	16.9272				
Test Weight	-0.0458	3.0390	1000 Seed Weight	-0.0543	2.4872				
Harvest Index	-0.0399	2.6467	Harvest Index	-0.2941	13.4739				
Protein Content	-0.3312	21.9658	Protein Content	-0.0918	4.2050				

For Harvest Index					For Protein Content				
Variables	Path Coef.	%	Variables	Path Coef.	%				
Direct Effect	-0.7661	30.5859	Direct Effect	-0.4749	24.9694				
Indirect Effects		Indirect Effects		Indirect Effects					
Via	Path Coef.	%	Via	Path Coef.	%				
Plant Height	-0.0130	0.5202	Plant Height	0.1939	10.1978				
Spike Number/m ²	0.3307	13.2044	Spike Number/m ²	0.3318	17.4449				
Spikelet Number	0.1521	6.0717	Spikelet Number	0.0329	1.7274				
Spikelet Number	0.0883	3.5253	Spikelet Number	0.1559	8.1965				
Seed Number/Spike	0.6577	26.2569	Seed Number/Spike	0.3436	18.0653				
Seed Weight/Spike	0.3420	13.6551	Seed Weight/Spike	0.0376	1.9794				
1000 Seed Weight	-0.0167	0.6678	1000 Seed Weight	-0.2240	11.7769				
Test Weight	-0.1040	4.1538	Test Weight	-0.0524	2.7547				
Protein Content	-0.0340	1.3589	Harvest Index	-0.0549	2.8877				

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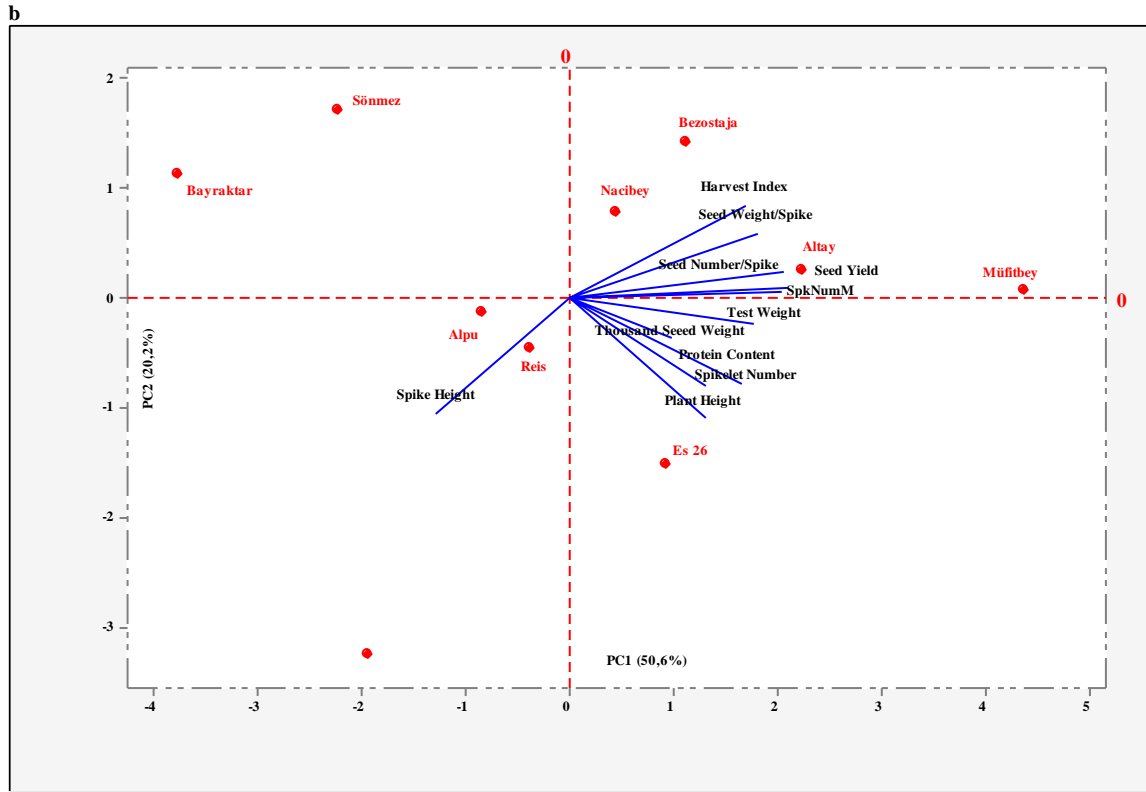


Figure 1. Path and biplot analyses showing the effect of yield components on seed yield.

via spikelet number; spike number/m², seed weight/spike, harvest index via seed number/spike; spike number/m², seed number/spike, harvest index via seed weight/spike; spike number/m², seed number/spike, seed weight/spike, protein content via 1000 seed weight; spike number/m², seed number/spike, seed weight/spike via harvest index; plant height, spike number/m², seed number/spike, 1000 seed weight via protein content were determined as important components on seed yield (Table 1 and Figure 1a). Correlation and path analyses revealed that spike number/m², seed number/spike, seed weight/spike and harvest index had important effect on seed yield in bread wheat. Biplot analysis revealed that spike number/m², seed number/spike, seed weight/spike and test weight were found as important factors on seed yield (Figure 1b). Besides, Müfitbey, Altay, Nacibey, Es 26 were high potential and stable genotypes. Results denoted that spike number/m², seed number/spike, seed weight/spike and test weight are important components; Müfitbey, Altay, Nacibey, Es 26 were high potential and stable genotypes.

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