

Yield Ability and Yield Stability, the Effective Tools Through Selection Procedure of Classified Wheat (*Triticum aestivum*) Crosses

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Abstract

Early generation selection in wheat breeding (*Triticum aestivum*), for stable high yielding genotypes, has been attempted using various parameters. The aim of the present study was the selection of genetic parameters that could be used for the assessment of crosses in the selection procedure. The experiments were conducted at the National Agricultural Research Foundation and the research farms of the Aristotle University of Thessaloniki for four growing seasons. Four out of ten crosses were chosen, based on a set of evaluation criteria estimating productivity and stability. Positive general combining ability of the parents was a prerequisite for any cross to remain in the selection procedure. The cross Oropos x Acheloos was ranked first in F1 and exhibited significant heterobeltiosis from F2 to F4 generation, producing elite F5 lines which yielded 45.05% at the best check. The ranking of the rest of the crosses in F1 remained the same in F5 generation showing that phenotyping achieved genotyping owing to isolation environment and high selection pressure. Genetic variance per cross gave a reliable estimation of the stability of the crosses through the segregating generations with Oropos x Acheloos being the less affected cross by environmental factors. It was concluded that early generation selection can successfully produce elite F5 lines, with an appropriate methodology which estimates productivity and stability, heterotic effects and the general combining ability of the parents.

Keywords: bread wheat, heterosis, general combining ability, genetic variance cross⁻¹, early generation selection

1. Introduction

Combined improvement for stable and high yield through breeding is faced at first, with the challenge of selecting an appropriate breeding method and, second, with choosing criteria such as yield performance per se, genetic relationship matrix, heterotic patterns, and the usefulness of derived population (Lamkey, Schnicker, & Melchinger, 1995; Schnell, 1983) etc, that would help the breeder to identify among a large number of crosses those most likely to yield elite lines. The above procedure must be handled in a way that may increase the precision of phenotyping, thus, maximizing selection efficiency and saving time and effort.

It has been pointed out that on testing early generations, the adverse effects of competition obscure the evaluation data (Hinson & Hanson, 1962; Khalifa & Qualest, 1975; Shebeski & Evans, 1973). The proposed honeycomb methodology for the evaluation of quantitative traits of widely-spaced single plants under optimal field conditions minimizes the effects of competition and maximizes selection efficiency (Fasoulas, 1977; Fasoula, 2013). Therefore, the individual plant performance, as a unit of evaluation and selection, corresponds with the genotypic level of high and stable crop yield and delineates the conditions that enhance selection efficiency in plant breeding (Koutsika-Sotiriou & Traka-Maurona, 2008).

Heterotic patterns are commonly recognized as heterosis over the better parent (heterobeltiosis), or heterosis over the mid parent (relative heterosis) (Fonseca & Patterson, 1968), or as standard heterosis (Virmani, 1994) referring to the comparison of the F1 with the mean yield of broadly cultivated varieties at a region. Heterosis in F1 and F2 generation, as a criterion for the identification of promising crosses, constitutes the standard breeding practice which dominates every breeding procedure (Nass, 1979; Fasoulas, 1988; Roupakias et al., 1997; Gouli-Vardinoudi & Koutsika-Sotiriou, 1999; Singh, Sharma, & Sain, 2004; Kotzamanidis, Lithourgidis, Mauromatis, Chasioti & Roupakias, 2008). The use of heterotic patterns promoted high-yielding crosses in F1 and F2 generation (Nass, 1979). Besides, the criterion of the proposed mean yield of F1 and F2 as a prediction criterion (Roupakias et al.,

1997) was successfully applied in barley (Kotzamanidis & Roupakias, 2004) and in durum wheat (Kotzamanidis et al., 2008). At this point, it may be added that in maize breeding, the main criterion for choosing an F2 as source material from a number of single-cross hybrids is the tolerance to inbreeding depression, an indication of possessing desirable load of additive genes (Hallauer & Miranda, 1995). Another parameter, revealing the genotypic profile of a parent variety, is the general or specific combining ability, which along with heterosis aims at the discrimination of promising material (Ali Avchi, 2005; Papadopoulos, Tokatlidis, Tamoutsidis, Koutsika-Sotiriou & Koutroubas, 2007). Therefore, it could be rather conducive to the breeder's task if the performance of late generations (F5 and further) could be predicted efficiently from the early generations performance (F1 and F2), even though the choice of the appropriate evaluation criteria is yet to be decided.

The present work elaborates on presenting a framework of genetic parameters for ranking a series of crosses in order to facilitate the common plant breeder's effort to develop varieties. Particularly, the effectiveness of parameters such as: (i) heterotic patterns, (ii) productivity and stability indexes and (iii) additive/dominant effects are being tested, aiming at identifying a number of crosses from which a sample of certain stability and yield ability will enter the selection procedure, facilitating a constant turnover of the varieties.

2. Method

Six bread wheat varieties (Acheron, Yecora-E, Nestos, Orfeas, Oropos, Acheloos) were crossed to produce 10 F1's. All the varieties were developed by the Cereal Institute of Thessaloniki and were chosen on the basis of their performance and stability in the Mediterranean environment. The characteristics of the cultivars used are shown in Table 1.

Table 1. Pedigree and agronomic traits of the six cultivars used in the crosses

Cultivar	Pedigree	Agronomic characteristics	Breeder(s)
Acheron	Pedigree selection of the cross Kal-Bb x Cj 71"S"/Hork"S" (Translocation 1BL/1RS)	Tillering: Moderate to extensive 1000 grains weight: 35 ± 5 g Resistance to lodging: Very Resistant Yield: 5800 ± 600 kg / ha Type: Spring	D. Gogas and S. Stratilakis
Yecora-E	Selection within CIMMYT's cultivar Yecora-70	Tillering: Moderate 1000 grains weight: 45 ± 5 g Resistance to lodging: Resistant Yield: 5500 ± 600 kg / ha Type: Spring	Cereal Institute and S. Skorda with participation in enrichment, description and registration from D. Gogas and S. Stratilakis
Nestos	Pedigree selection for four successive generations (F5 to Fm) within the INIA66R//Hbgn/ drc material of Oregon State University	Tillering: Medium - rich 1000 grains weight: 38 ± 2 Resistance to lodging: Very Resistant Yield: 6000 ± 500 Type: Multi type	D. Gogas and S. Stratilakis
Orfeas	Selection within naturally mutated lines of the cultivar Nestos (1BL/1RS Translocation).	Tillering: Satisfactory 1000 grains weight (g): 33 ± 2 g Resistance to lodging: Very Resistant Yield: 6000 ± 500 Kg / ha Type: Spring	Cereal Institute and D. Gogas
Oropos	Pedigree selection of the cross S. Cerros' YG-3927	Medium Tillering 1000 grains weight: 35 ± 5 g Resistance to lodging: Resistant Yield: 5000 ± 500 kg / ha Type: Spring	Cereal Institute and D. Gogas
Acheloos	Pedigree selection of the cross YT-5143 x YT-3615 (Siete CERROS – T66xNIKH)	Tillering: Moderate 1000 grains weight: 36 ± 2 g Resistance to lodging: Good Yield: 6500 ± 400 kg / ha Adaptability: Special for fertile soils	Cereal Institute, D. Gogas and S. Stratilakis

The experiments were carried out for three growing seasons (F1-F3) at the farm of the Cereal Institute of Thessaloniki (latitude 40°32'N, longitude 23°00'E, elevation 15 m) and for two growing seasons (F4-F5) at the farm of Aristotle University of Thessaloniki at a distance of 1km. The selection experiments, F1 to F4 generation, were established according to the pattern of honeycomb design (Fasoulas, 1973; Fasoulas & Fasoula, 2000).

An interplant spacing of 1 m (plant density 1.16 m⁻²) between each plant and between rows was used. To ensure one plant at each position three kernels were sown per position. Five weeks after sowing, all positions were thinned to a single plant. A few days before threshing all plants were tagged for identification. Threshing occurred in the field and individual plant yield was recorded (g plant⁻¹). All experiments were subjected to growing conditions promoting high yields.

The evaluation experiment of the selected lines of the F5 generation was established under solid stand and a randomized complete block design (RCBD) consisting of three replications was used. The plots consisted of seven rows 0.60 m long, spaced 25 cm apart (a total area of 1 m² per plot) from which only the five central rows were harvested. For each plot 18 g of seed were used and appropriate agronomic practices were done timely to achieve good crop stand. Threshing occurred in the field and seeds from each plot were weighted to determine yield (g plot⁻¹).

2.1 Pre-Selection of the Classified Group of Crosses

Ten F1 crosses and their parents were evaluated for grain yield in an R-16 honeycomb design. A total of 480 single plants were evaluated with 30 replications per entry. Also, their F2 generation was evaluated in solid stand with their parents in pre-selecting manipulations (Gogas & Koutsika-Sotiriou, 2012). Three groups of evaluation criteria were used: a) Heterosis patterns, b) Productivity and stability and c) General Combining Ability (GCA) following Griffing's (1956) estimates. Four crosses were selected according to the assessment of yield ability and stability: the highest one, two with medium values and one with the lowest value. In this way from a total of 10 crosses, four crosses with classified yield ability and stability were selected: Oropos x Acheloos (OxA), Yecora x Oropos (YxO), Yecora x Acheloos (YxA) and Orfeas x Oropos (OxO). Selection intensity was 2.7% i.e. a given plant was compared with 36 plants in the grid (moving ring Figure 1). For single plant evaluation Plant prognostic equation (p-PE) was applied where the yield ability and stability were co-considered, according to Fasoula (2013). Within each of the four crosses three higher yielding F1 plants (in p-PE) were selected.

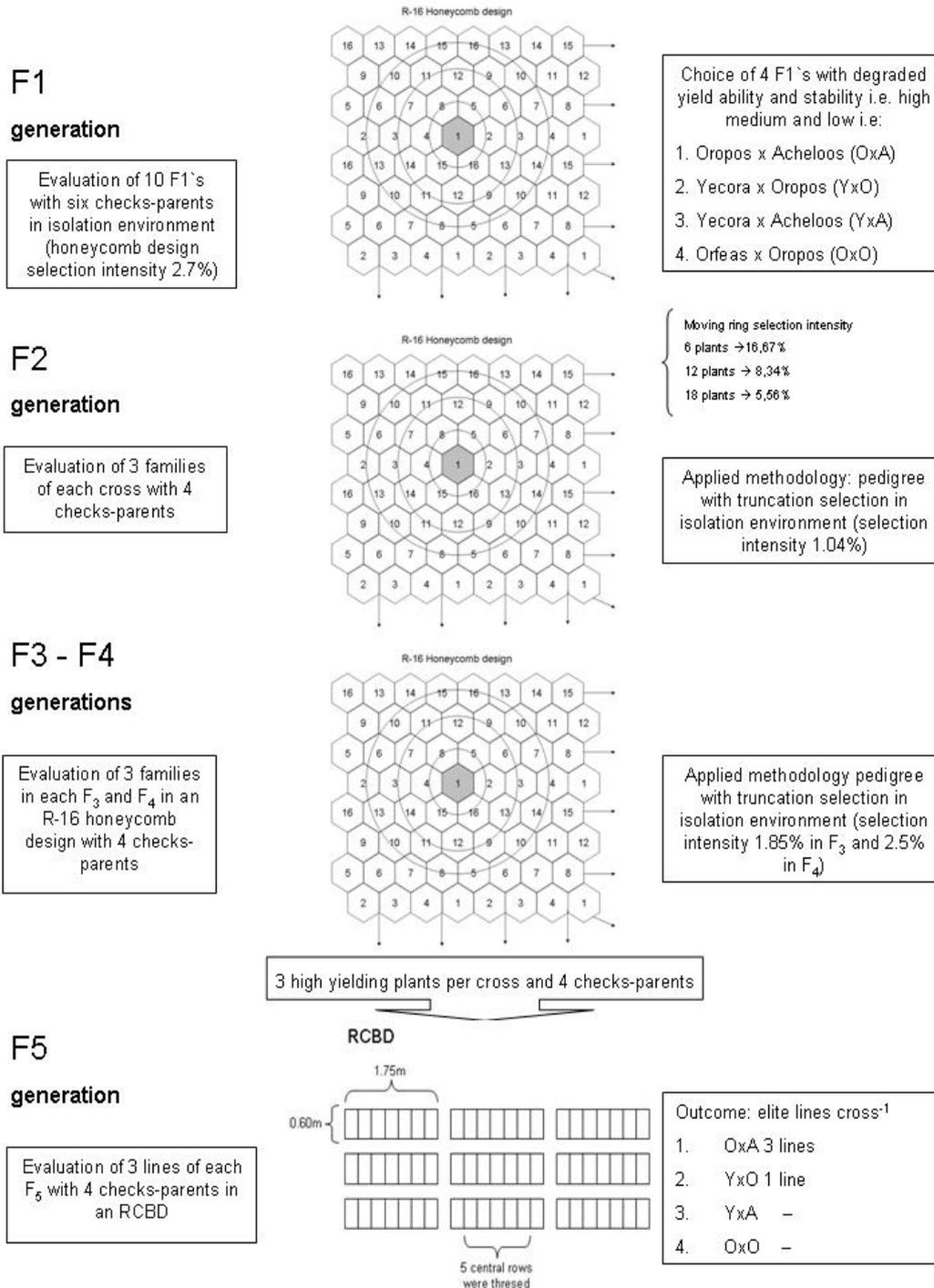


Figure 1. The followed breeding schedule

2.2 Evaluation of F2-F5 Generations

For the four classified crosses, seed from each of the three plants of F1 was sown as an independent entry in an R-16 honeycomb design (Figure 1), with the four parent-varieties used as checks (4 crosses x 3 plants per cross i.e. 12 entries, plus 4 parents a total of 16 entries). From the F2 to the F4 generation pedigree truncation selection was applied with an intensity of selection 1.04% for F2 (3 out of 288 plants per cross were selected), 1.85% for F3 (3 out of 162 plants per cross were selected) and 2.5% for F4 (3 out of 120 plants per cross were selected). Each selected plant constituted a family (i.e. 3 families per cross) in each generation (Figure 1). The evaluation from F2 to F4 generation was applied following the three groups of evaluation criteria previously mentioned, except the

GCA which was calculated through the F2 generation.

Finally, seed from the selected high yielding plants of each cross of the F4 generation was established in a RCBD in dense stand with the presence of the four parents, as checks, for the comparative trial of F5 generation. The evaluation was applied with the criteria of yield heterosis and stability of performance.

2.3 Statistical Analysis

In the experiments for mean comparisons the t-test for independent samples from populations with different standard deviations was used with application of Cochran's approximation (Snedecor & Cochran, 1967). Heterosis was estimated: (i) over the best parent (heterobeltiosis) and (ii) over mid parent (relative heterosis) (Fanseco & Peterson, 1968). At the present work the mean yield of the four parents/checks was used as the expression of standard heterosis. Standard heterosis refers to the comparison of an F1 with one or more broadly cultivated varieties at the region (Virmani, 1994). The significance of relative and standard heterosis was checked with one degree of freedom comparisons.

For the isolation environment two formulas, proposed by Fasoula (2013), were estimated: (i) the Sibling line prognostic equation (s-PE) and (ii) the Plant prognostic equation (p-PE). s-PE evaluated the performance of each entry and was specified as the product of two components: a) Coefficient of Homeostasis $CH = (\bar{x}_i/s)^2$ where \bar{x}_i is the mean of each treatment and s is the standard deviation and b) Sibling Yield Index $s - YI = (\bar{x}_i/\bar{x})^2$ where \bar{x} is the grand mean. s-PE was used for the correspondence of the isolation environment with solid stand. For the evaluation/ranking of single plants p-PE was determined as the product of two components: a) CH and b) Plant Yield Index $p - YI = (x/\bar{x}_r)^2$ where x is the yield of the plant in the center of the moving ring and \bar{x}_r is the mean yield of the plants composing the moving ring around the plant (Figure 1). (Fasoula, 2013)

The general combining ability (GCA) was estimated for the incomplete diallel according to Griffing's (1956) diallel analysis. Analysis of variance for general and specific combining abilities was performed for the F1 and F2 generation and the genetic components were estimated for grain yield.

3. Results

Table 2. Ranking of F₁'s: heterosis over better parent (BP), mid parent (MP), standard heterosis (over six parents SH), grain yield (g/plant GY), productivity (Sibling Yield Index, s-YI), stability (Coefficient of Homeostasis, CH) and yield potential (s-PE)

Genotype	BP	MP	SH	GY Mean	s-YI	CH	s-PE
Acheron x Yecora	3	5	1*	1a**	1	6	5
Nestos x Acheloos	1*	1*	4	4ab	4	7	8
Nestos x Orfeas	2	3	7	7abc	7	5	7
Nestos x Oropos	7	6	5	5abc	5	4	4
Orfeas x Acheloos	5	9	9	9abcd	9	1	2
Orfeas x Oropos	10	10	10	10bcd	10	10	10√
Oropos x Acheloos	8	4	8	8abcd	8	2	1√
Yecora x Acheloos	4	2*	2*	2ab	2	8	6√
Yecora x Orfeas	9	8	6	6abc	6	9	9
Yecora x Oropos	6	7	3*	3ab	3	3	3√

*significant positive heterosis.

** Means followed by different letters are significantly different at the 0.05 level.

√ selected crosses.

Modified data of F1 generation (Gogas & Koutsika-Sotiriou, 2012) with ten F1s were compiled in Table 2. The F1s were ranked according to three heterotic patterns and to four productivity and stability indexes. Each of the pre-mentioned parameters was ranked in a scale from 1 to 10, for higher and lower value, respectively for each F1. Accordingly, three F1s showed significant standard heterosis and were equal to best parent and mid parent values, and one surpassed significantly mid parent value. Productivity and stability indexes gave a more clear view of the

performance of the crosses classifying them into high, medium and low yielding ones.

Only F1s with, positive general combining ability (GCA) of the parents were considered for further selection. Therefore, discarding F1s with parents with negative GCA (Table 3), a selection of three classes concerning productivity and stability i.e. high, medium and low, could be entered to the experiment. According to the ranking of the F1s with the aforementioned parameters the following were selected: a) Oropos x Acheloos (OxA) having the highest s-PE value i.e. high productivity and stability and being heterotic, b) Yecora x Oropos (YxO) ranking third in productivity and stability index s-PE and exhibiting significant standard heterosis, c) Yecora x Acheloos (YxA) which ranked sixth in productivity and stability index and exhibited significant heterosis and standard heterosis and d) Orfeas x Oropos (OxO) the F1 with the lowest s-PE value and with negative heterosis over best and mid parent value.

The diallel analysis for grain yield of the F1 and F2 generation gave further insight on the genetic basis of the first generations. Mean squares of GCA and SCA were found significant for both generations, the results being in agreement with other studies (Perenzin., Corbellini, Accerbi, Vaccino, & Borghi, 1998). The mean squares for GCA were two and three times higher in the F1 and F2 generation than for SCA effects, respectively, indicating the predominance of additive gene action (Table 3). However, the ratio of $V_{GCA}:V_{SCA}$ was greater than the unity (> 1) in the F2 generation indicating the preeminent role of additive effects for grain yield (Table 3), while in the F1 generation the variance ratio of the combining abilities was lower than the unity (< 1) since in F1s heterotic effects are due to dominant gene action. Since V_{GCA} is a function of additive variance and V_{SCA} is equal to the variance component due to dominance variance, when epistasis is assumed negligible (Bernardo, 2002), heterotic patterns in the F2 generation are more likely to be inherited in the next generations than in the F1 generation.

Table 3. General combining ability (GCA) of the parents in F₁ and F₂, mean squares of GCA/SCA and the variance components

Parents	Acheron	Yecora-E	Nestos	Orfeas	Oropos	Acheloos
GCA F ₁	-26.72	13.95	-4.2	4.66	5.36	6.95
GCA F ₂	-	-0.6	-	-30.01	50.18	4.38

Source	d.f F ₁	MS F ₁	d.f F ₂	MS F ₂
GCA	5	1640.09**	3	2144.76*
SCA	9	991.89**	2	648.85**
Error	14	60.78	5	36.59

*significant at $p \leq 5\%$, **significant at $p \leq 1\%$.

Generation	F ₁	F ₂
Var GCA	205.011	536.191
Var SCA	495.946	324.425
GCA/SCA	0.41	1.65

Throughout F2 to F5 generation the initial ranking of the crosses didn't remain stable except for the first and the last classified crosses. OxA, in particular, gave significantly high BP, MP and 4P values through F2 to F5 (Tables 4 and 7) outyielding all other crosses. In terms of productivity and stability OxA outyielded the rest, scoring the highest values for s-YI and CH, with an exception in F3 generation where it was equal to YxA (Table 5). YxA performed well in F3 during the less productive year (Tables 4 and 5); however in F4 and F5 generation (dense stand) it was third in terms of productivity and stability. YxO, also classified as medium yielding cross showed an increase in s-PE from F2 to F4 generation (Table 5) exhibiting significant heterosis, heterobeltiosis and standard heterosis in F5 (Table 6). Three crosses out of four (except OxO) had $s-YI > 1$ from F1 to F4 (Tables 2 and 5) an indication of promising segregating material (Fasoula, 2008). The low yielding cross OxO, although heterotic (non significant) in F4 and F5, in terms of productivity and stability in the isolation environment, was below the 50% level of the highest yielding cross.

Table 4. Estimation of heterobeltiosis (BP), relative heterosis (MP) and standard heterosis (over four checks, SH) for F₂ to F₅ generation

Genotype/Heterosis	Generations		
	F ₂	F ₃	F ₄
Oropos x Acheloos			
BP	13.12%*	18.48%*	19.74%*
MP	17.78%*	23.56%*	35.21%*
SH	38.62%*	43.49%*	37.61%*
Yecora x Acheloos			
BP	-6.41%	46.07%*	9.91%
MP	6.64%	53.91%*	9.89%
SH	14.68%*	62.34%*	23.71%*
Yecora x Oropos			
BP	-0.50%	-2.51%	14.02%
MP	9.34%	6.89%	26.40%*
SH	12.29%	18.08%*	25.67%*
Orfeas x Oropos			
BP	-8.24%	-18.04%	16.93%
MP	12.00%	5.02%	18.08%
SH	3.55%	-0.73%	3.23%

*significant at $p \leq 5\%$.Table 5. The yield ability of successive generations (F₂ to F₄) of the degraded crosses and their parents for grain yield (g/plant GY), productivity (Sibling Yield Index, s-YI), stability (Coefficient of Homeostasis, CH) and yield potential (s-PE)

Genotypes/Generations	Mean yield			s-YI			CH			s-PE%		
	F ₂	F ₃	F ₄	F ₂	F ₃	F ₄	F ₂	F ₃	F ₄	F ₂	F ₃	F ₄
Oropos x Acheloos	66.97a*	30.6a	82.69a*	1.66	1.60	1.61	6.66	5.03	8.06	100	81.36	100.00
Yecora x Acheloos	55.4bc	34.61a*	74.34abc	1.13	2.05	1.30	5.21	4.83	7.61	53.55	100	76.26
Yecora x Oropos	54.25bc	25.17bc	75.51ab	1.09	1.08	1.35	3.85	4.17	9.03	37.94	45.76	93.40
Orfeas x Oropos	50.03c	21.16c	62.03cd	0.92	0.77	0.91	4.02	5.41	6.10	33.69	41.89	42.53
Parents/Year	2009	2010	2011	2009	2010	2011	2009	2010	2011	2009	2010	2011
Acheloos	59.2b	23.70bc	69.05bc	1.3	0.96	1.13	5.44	4.38	7.70	63.84	42.58	66.61
Orfeas	34.82e	14.48d	51.81d	0.45	0.36	0.63	4.97	8.90	5.22	33.22	32.28	25.42
Oropos	54.52bc	25.82b	53.26d	1.1	1.14	0.67	4.89	6.06	6.72	48.65	69.84	34.58
Yecora	44.71d	21.28c	66.23bc	0.74	0.77	1.04	4.97	5.96	6.24	20.17	46.69	49.60

* Means followed by different letters are significantly different at the 0.05 level.

Table 6. The genetic components of diallel analysis of F₁ and F₂ for grain yield

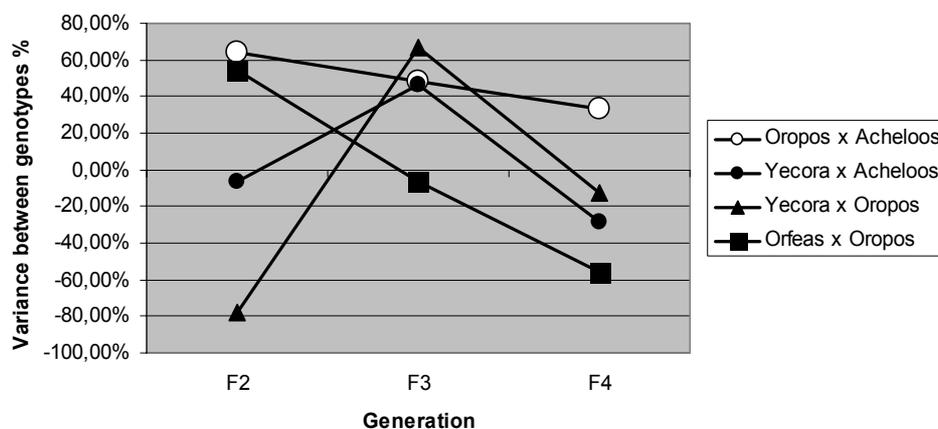
Genetic component	F ₁	F ₂
D ⁽¹⁾	25.7411	135.7337
H ₁ ⁽²⁾	1972.187	1649.376
H ₂ ⁽³⁾	1571.295	1302.23
H ₂ /4H ₁ ⁽⁴⁾	0.199	0.197
(H ₁ /D) ^{1/2} ⁽⁵⁾	8.753	3.486
F ⁽⁶⁾	92.2659	-279.683
kd/(kd+kr) ⁽⁷⁾	0.6024	0.3522
H ⁽⁸⁾	-11.5	-21
H _b ² ⁽⁹⁾	0.955	0.98
H _n ² ⁽¹⁰⁾	0.285	0.528

1) Additive variance; 2) Dominance variance1; 3) Dominance variance2; 4) Balance of positive and negative alleles; 5) Average degree of dominance; 6) Product of add. by dom. Effects; 7) Proportion of dominant genes; 8) Average direction of dominance; 9) Heritability for diallel in a broad sense; 10) Heritability for diallel in a narrow sense.

Furthermore the genetic components in both generations showed that the additive variance component (D) was higher in F₂ generation than in F₁ but relatively lower than the dominance variance components (H₁ and H₂) for both generations (Table 6). The balance of positive and negative genes (H₂/4H₁) was equal to 0.199 in F₁ generation and 0.197 in F₂ generation (lower than 0.25) suggesting unequal mean allelic frequencies at the loci with dominant and recessive genes (Table 6). The average degree of dominance (H₁/D)^{1/2} although reduced in F₂ generation it was greater than the unity suggesting the presence of overdominance.

The product of additive by dominant effects (F) in F₁ generation was positive denoting that dominant genes were more frequent than recessive ones while the opposite was found for F₂ generation (negative value for F, Table 6). In addition the proportion of dominant genes (kd/(kd+kr)) for both generations was lower than the unity reflecting an excess of recessive genes (Table 6). Average direction of dominance (H) was negative for both F₁ and F₂ generation meaning that the presence of dominant genes was an inhibitory factor for increasing grain yield (Table 6). Thus, even with the indications for the presence of overdominance, the diallel analysis pointed at the selection of recessive genes for increasing grain yield, which could be easily fixed.

Estimates of variance due to genotype and due to experimental error for generations F₂ to F₄ were made during selection procedure. At the first cross from the classified (OxA) and at the last one (OxO), a gradual decrease in σ_g^2 was observed (Figure 2) which was anticipated due to inbreeding and to an increase in homozygosity. On the other hand, the other two classified crosses YxA and YxO, gave negative values in σ_g^2 in F₂ generation which was significantly increased (in F₃) and then dropped (in F₄) (Figure 2).

Figure 2. Depicting the genetic variance cross⁻¹ through generations

The results of F5 lines clarified a high yielding line derived from the cross ranked first in the classification (i.e. OxA) with the highest yielding stability. From the beginning of the selection procedure the OxA was ranked first owing to its high yield stability (high CH Table 2) and from the results of F2 evaluation it could be selected for the development of stable and high yielding F5 lines. OxA overyielded 18.43% from the following highest yielding F5 line and 45.05% (significant superiority) from the highest yielding 'Acheloos'. Moreover, in terms of stability it accounted for only 1.46% of the experimental error while the second ranking F5 line, derived from YxO, accounted for 6.20% of the experimental error (four times higher) indicating less stable material. Besides, two more lines without significant difference in yield ability derived from the cross OxA (ranked 3rd and 4th Table 7), having also high stability performance (experimental error 2.18% and 3.22%) Thus, high and stable F5 lines could be isolated successfully from early generations after multiple criteria assesment.

Table 7. Mean separation of the F₅ lines with heterotic and stability criteria

Genotype	Heterotic criteria			Mean		Stability criteria
	BP	MP	SH			% experimental error
Oropos x Acheloos	45.05%*	69.23%*	72.93%*	264	a**	1.46%
Yecora x Oropos	54.55%*	59.90%*	41.05%*	215.33	ab	6.20%
Oropos x Acheloos	12.09%	30.77%	33.62%	204	abc	2.18%
Oropos x Acheloos	11.36%	29.91%	32.75%	202.67	abc	3.22%
Yecora x Oropos	39.23%	44.06%	27.07%	194	bcd	18.99%
Yecora x Acheloos	0.73%	14.11%	20.09%	183.33	bcde	0.80%
Acheloos				182	bcde	13.54%
Yecora x Oropos	29.19%	33.66%	17.90%	180	bcde	9.80%
Orfeas X Oropos	9.83%	20.97%	14.63%	175	bcde	7.16%
Orfeas X Oropos	6.28%	17.05%	10.92%	169.33	bcdef	15.04%
Orfeas				159.33	bcdef	6.91%
Yecora x Acheloos	-21.43%	-11.00%	-6.33%	143	cdef	1.38%
Yecora				139.33	cdef	11.32%
Oropos				130	def	0.84%
Orfeas X Oropos	-22.59%	-14.75%	-19.21%	123.33	ef	0.28%
Yecora x Acheloos	-42.86%*	-35.27%	-31.88%	104	f	0.86%

* Significant at p=5%.

**Means followed by different letters are significantly different at the 0.05 level.

4. Discussion

Early generation selection in bread wheat breeding is a challenging procedure since there is no specific methodology which can accurately discriminate promising genetic material. Heterotic patterns are typically preferred than non-heterotic ones in the selection of F₁, as they are likely to give transgressive segregants (Briggle, Cox, & Hayes, 1967; Nass, 1979; Gouli & Koutsika, 1999; Corbellini, Perenzin, Accerbi, Vaccino, & Borghi, 2002). Therefore the highly significant values of better parent and mid parent heterosis of an F₁ consists a priority to be entered to selection procedure (Singh et al., 2004). On the other hand, significant standard heterosis represents higher performance than the mean yield of well-adapted varieties and it may be considered a practical point of view as it represents its cultivation performance (Alam et al., 2004). At the present study ten out of ten F₁s exhibited positive heterotic patterns thus, when dealing with a large number of crosses heterosis alone cannot determine which of them should remain in the selection procedure. The risks of selecting crosses based only on data from F₁ generation due to dominance effects have already been mentioned (Gogas & Koutsika-Sotiriou, 2012).

Many researchers have proposed positive GCA of the parents of a cross as a selection criterion (Borghi & Perenzin,

1994; Barnard, Labuschagne, & Van Niekerk, 2002; Corbellini et al., 2002; Rebetzke, Richards, Condon, & Farquhar, 2006; Gogas & Koutsika-Sotiriou, 2012) and eventually a number of heterotic crosses can be discarded. Since high and stable yield is the aim of a wheat breeder, a classification concerning these parameters should be taken into consideration. Genes controlling productivity and stability can be approached through certain indexes (i.e. CLR, CH etc). High stability in F1/F2 can be interpreted as high heritability or as a low load of deleterious genes (Fasoula, 2008). The product of productivity and stability parameters classified the crosses to be evaluated in three classes i.e. high, medium and low.

According to this logic four classified crosses were chosen: the first and the last one were in accordance to Nass (1979) suggestions that high yielding F1s give better F4s than the low yielding ones. For medium yielding crosses i.e. YxA and YxO, one may clarify the following: The YxA significantly outyielded four parents value, from F2 to F4 generation; however in F5 generation, its performance was equal to the lowest yielding cross, so it may be characterized as poor material. The YxO outyielded four parents value in F3 and F4 generation while in F5 generation its performance exhibited a significant heterobeltiosis, relative and standard heterosis. Searching for help in GCA data of their parents, 'Yecora' and 'Orfeas' had negative values while 'Oropos' and 'Acheloos' had positive values. Thus the highest yielding F5 line derived from varieties with positive GCA in F2 and the lowest yielding F5 line derived from varieties with negative GCA in F2. From the medium yielding crosses YxO had as parents 'Yecora' (with slightly negative GCA) and 'Oropos' (with the highest GCA in F2), which eventually led to a high yielding F5. On the other hand, YxA with 'Yecora' and 'Acheloos', the latter having positive GCA but approximately 8.7% the GCA of 'Oropos', ended up eventually with a non promising F5. According to Soriano Viana (2000) such a variation in GCA values indicates strong differences in allele frequencies and genetic divergence among the diallel parents for the trait under study.

The diallel analysis showed an anticipated lower frequency of dominant genes in F2 generation while overdominance was assumed for both F1 and F2. Overdominance hypothesis states that heterosis is due to inter allelic interactions and it isn't fixable (Singh et al., 2004). In the case that single gene overdominance is important in heterosis then it is impossible to obtain homozygotes as vigorous as heterozygotes. However, the fixation of dominant genes is controversial in the sense that disadvantageous overdominant genes have a low probability of fixation whereas advantageous overdominant genes are more probable to be fixed (Masatoshi & Roychoudhury, 1973). Furthermore, there is substantial evidence that heterosis is not controlled by a single locus alone, even if that locus behaves in a dominant or overdominant way (Yu., Hiatt, Chan, Sweeney, & Dawe, 1997; Monforte & Tanksley, 2000). Moreover, overdominance can be in fact pseudo-overdominance, as linkage and epistasis among multiple loci may contribute in the expression of heterosis (Budak, 2002). At the present study, besides the presence of overdominance, an excess of recessive alleles were also found. Since the direction of dominance was negative, selection for recessive genes may increase grain yield, a situation that can be easily fixed with the appropriate methodology applied, which discriminated those genotypes. Additionally, the extremely low selection intensity that was applied (ranging from 2.7% and 1.04% for F1 and F2 up to 1.85% and 2.5% for F3 and F4 respectively) favored them as homozygosity was increased.

The main issue emerged from the present study was the significance of agronomic stability in classified crosses. Their catalytic role overshadowed all other multiple criteria. Control could be achieved by a plethora of recessive alleles that contributed to the late generations for elite lines. In terms of agronomic stability the cross OxA maintained positive values of σ^2_g through inbreeding from F2 to F4 (Figure 2). The increase of σ^2_e (residual percentage of σ^2_g) in F3 and F4 generations is due to the increase of homozygosity rather than environmental effects thus rendering OxA both stable and productive. Indeed, the exploitation of a positive GE-interaction-based package was the core of the success of the Green Revolution (Simmonds, 1979; Simmonds, 1991). All the other crosses showed negative values of σ^2_g in at least two of the three generations of selection.

Numerous scholars dealt with the issue of the negative variance components. Some, on one hand, supported the exact reporting without changing. Others, on the other hand, kept a different line, insisting on negative estimates being tagged as zero (Littell, Stroup, & Freund, 2002) since variances cannot be negative by definition (Brown & Mosteller, 1991). On the other hand a negative variance component might be a useful diagnostic tool (Hocking, 1983; Searle, Casella, & McCulloch, 1992) especially in the interpretation of the biological significance of this value. The results of the present study indicated negative values of σ^2_g for three crosses, namely YxA, YxO and OxO. From a genetic point of view this could be interpreted as a "masking" effect of the environment over the genotype, which means significant genotype by environment interaction as the generations were evolving, exhibiting negative values of σ^2_g . Finally, the fluctuation, occurred in σ^2_g value in the two of them (YxA and YxO), possibly showed that these crosses were less affected from the genotype by environment interaction in a stressful year.

Studying a couple of genetic parameters that are used in the evaluation of segregating generations in wheat breeding, one may conclude the following: The isolation environment of selection clarified the whole procedure owing to the fact that phenotyping achieve genotyping in the differentiation of the crosses (Newton et al., 2010), discriminated and predicted successfully the elite F5 lines. The low yielding, non heterotic crosses in F1 and F2 generation fail to succeed promising lines in late generations and may be ignored in a breeding program with a lot of crosses. The standard check heterosis suggested more steadily crosses with high yield potential, under the prerequisite of the general combining ability in F1 and F2 generation; the genetic components provide further insight for the genetic background of the parents and could help in the choice of source material. Therefore, as an epilogue, an appropriate methodology which estimates productivity and stability, heterotic effects and the general combining ability of the parents, may be a prosperous leader for a successful outcome.

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