

General and specific combining abilities of local common bean varieties and cultivars

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ABSTRACT. We examined the general and specific combining abilities of common bean cultivars and varieties and tested promising hybrid combinations for the development of productive strains. Two cultivars (IPR Uirapuru and IPR Campos Gerais) and five local varieties (BGU-1, BGU-7, BGU-13, BGU-25 and BGU-36) were planted in a full diallel cross. Treatments were assessed based on a randomized block design, with 10repetitions. The following variables were assessed: mean number of days still emergence, mean number of days till blooming, mean insertion height of the first pod, mean final plant height, mean stem diameter, mean length of the pods, mean number of pods per plant, mean number of seeds per pod, mean number of seeds per plant, mean weight of 50 seeds and grain yield. A Griffing statistical analysis was performed. Cultivar IPR Uirapuru and the local variety BGU-36 were promising for most variables in the general combining ability. Combinations IPR Uirapuru x BGU-7, IPR Uirapuru x BGU-1, BGU-7 x BGU-1, BGU-25 x BGU-13 and

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BGU-13 x BGU-36 were the most promising in the specific combining ability to generate highly productive strains.

Key words: *Phaseolus vulgaris*; diallel analysis; plant breeding; genetic effects

INTRODUCTION

The common bean (*Phaseolus vulgaris*) is one of the main legumes used for human consumption (Broughton et al., 2003; Demelash, 2018), being considered a major source of proteins, carbohydrates, micronutrients, vitamins and dietary fiber (Talukder et al., 2010). In order for this legume to continue to be a key part of human diet, it is necessary that new varieties be developed, attending the demands of production verticalization (Rocha et al., 2014). Genetic enhancement is the main tool to launch new cultivars in the market.

Bean breeding programs emphasize the exploration of genetic varieties through the hybridization of superior and contrasting cultivars. However, just as it happens all around the world, genetic variability in Brazil tends to decrease due to the intense cultivation of single cultivars or of cultivars coming from few parents (Tsutsumi et al., 2015). Bitocchiet al. (2013) found that Mesoamerican-origin bean cultivars have gone through drastic genetic-diversity reduction; approximately 72% of their diversity was lost during the domestication process.

Progeny (F₁ hybrids) studies have been conducted after the hybridization process. Various methods were used to implement the diallel analysis model proposed by Griffing (1956) (Barelli et al., 2000; Gonçalves-Vidigal et al., 2008; Arunga, et al., 2010; Trindade et al., 2015; Moura et al., 2016). This analysis allows estimating parent's general combining ability (GCA) and specific combining ability (SCA) (Spragueand Tatum, 1942).

Genetic enhancement can improve creole material from a certain region, because this material presents properties and characteristics similar to that of commercial cultivars (Voysest et al.,1994). Incorporating highly productive local varieties into high-yield cultivars can help optimize production with low additional input (Fess et al., 2011). We examined the GCA and the SCA of various bean genotypes in diallel crosses, to search forthe most promising combinations for the production of productive strains.

MATERIAL AND METHODS

Seven common bean parents, including five local varieties (BGU-1, BGU-7, BGU-13, BGU-25 and BGU-36) of *Phaseolus vulgaris* from the Active Germoplasm Bank of the State University of Mato Grosso, Caceres Campus, as well as two commercial cultivars: IPR Campos Gerais and IPR Uirapuru from the Agronomic Institute of Paraná (IAPAR) were used to obtain F₁ hybrids in a full diallel scheme, without reciprocals. To obtain hybrids (F₁ generation), crosses were made with emasculation of flower buds with interlacing, requiring four successive parent plant seedings during 2016, from January through September. Crosses were made in the morning (07:00 to 10:00 h) and in the afternoon (15:00 to 18:00 h). From October to December, crosses were made in the afternoon until 18:30 due to daylight summer time, this being one hour earlier compared to earlier months. Mean temperature during the crosses ranged from 20 to 34°C.

The performance of parents and F_1 hybrids was assessed from February to April 2017 in a screened greenhouse at Empresa Mato-Grossense de Pesquisa, Assistência e Extensão Rural (EMPAER), Caceres County, Mato Grasso State (16°43'42" S and 57°40'51" W). According to Köppen's classification, the characteristic climate in the region is tropical, hot and humid, with a dry winter (Awa). The rainfall period is from October to March and the dry season is from April to September (Neves et al., 2011).

Cultivation of parents and of their 21 hybrids was carried out in 5.0L plastic potsfilled with the commercial substrate Topstrato HT^{\circledast} . Mineral fertilizer was added before cultivation; 20g of N, P_2O_5 and K_2O 4-14-8 was added to each vessel. Fertilization was started after 15 days by weekly applying nitrogen at a dose of 50 mg/dm³, using ammonium sulfate[(NH₄)₂SO₄]. Potassium (K_2SO_4) was applied at a dose of 75 mg/dm³ at blooming.

A randomized block design, with 26 treatments and 10 repetitions, was adopted. The experimental units were composed of two vessels with two bean plants each. The following variables were assessed: mean number of days till emergence (EMERG) mean number of days to blooming (BLOOM), mean insertion height of the first pod (INSHEI), mean final plant height (MFPR), mean stem diameter (MSD), mean length of the pods (MLLP), mean number of pods per plant (MNPPL), mean number of seeds per pod (MNSP), mean weight of 50 seeds (MWS) and grain production (GP).

The methodology by Griffing (1956) – Method 2- was used in the diallel analysis; it analyzes the p(p+1)/2 combinations, which correspond to parents and their crossing processes, without reciprocals. Parents were considered the fixed effect and only the experimental error resulted from the random effect. Conclusions in this research were limited to genotypes. The statistical analyses were conducted with the Genes software, version 2016.6.0 (Cruz, 2013).

RESULTS AND DISCUSSION

All assessed variables presented significant values at 1% probability in the F test (Table 1). This indicates genetic differences among the 21 hybrid combinations. This shows that the parents diverge from each other, which is an important feature for the success of a breeding program, because genetic variability favors the selection of promising materials (Carvalho et al., 2016).

The mean values of quadratic behaviors recorded for GCA effects were higher than the values of SCA in variables INSHEI and MLLP. This outcome indicates additive gene effects in the expression of these variables. The additive effects of INSHEI were previously reported by Barelli et al. (2000). According to the results of the aforementioned variables, the best option to enhance this population is intra populational.

The mean SCA values of quadratic components in EMERG, INSHEI, TNPP, MNSP, MNSPL, PMS and PG were higher than the GCA values; this implies non-additive type gene action. Inter populational enhancement with heterosis can, under these circumstances, provide superior genetic gains to these variables in comparison with intra populational enhancement strategies.

The assessed parents can be used either in intra or inter populational enhancement in BLOOM and MSD. Therefore, both the gene additive and non-additive effects were equally important, as can be observed by the relative proximity between the means of GCA and SCA quadratic effects (Table 1).

Table 1 – Diallel analysis based on method 2, Model 2 by Griffing (1956) applied to the 11 variables: mean number of days till emergence (EMERG), mean number of days tillblooming (BLOOM), mean insertion height of the first pod (INSHEI), mean final plant height (MFPR), mean stem diameter (MSD), mean length of the pods (MLLP), mean number of pods per plant (MNPPL), mean number of seeds per pod (MNSP), Mean weight of 50 seeds (MWS) andgrain production (GP) in common beans, by specifying the effects of the general combining ability (GCA) and of the specific combining ability (SCA).

Median Squares of Agronomic Variables												
FV	GL	EMERG	BLOOM	INSHEI	MFPR	MSD	MLLP	TNPP	MNSP	MNPPL	MWS	GP
Genotypes	27	3.5776**	28.6031**	1247.6617**	3.3764**	3.8237**	2.9373**	107.9482**	3.5976**	3949.1582**	30.1584**	88.5817**
GCA	6	2.6735**	70.3735**	4536.5231**	10.0423**	9.8533**	10.6256**	58.8970**	9.7417**	7812.7580**	64.1581**	26.6140**
SCA	21	3.8359**	16.6688**	307.9870**	1.4719**	2.1009**	0.7407**	121.9628**	1.8421**	2845.2726**	20.4441**	106.2867**
Residue	243	0.4487	2.0849	145.9691	0.1162	0.2814	0.1946	31.3799	0.7786	685.0527	5.4727	28.4130
Median squares of the effect												
GCA		0.0046	2.0707	8480.3061	0.0417	0.4047	0.0465	3.1764	0.0394	26070.8685	1.7430	1.6197
SCA		0.1417	2.6819	1078.7568	0.2074	0.4269	0.0055	149.7706	0.3731	80963.2564	4.2271	114.2336
Residue		0.4487	2.0849	145.9691	0.1162	0.2814	0.1946	31.3799	0.7786	685.0527	5.4727	28.4130

^{**}significant effect at 1% in the F test

The GCA estimates allow breeders to obtain information about the concentration of favorable alleles; the greater the GCA magnitude, the higher the frequency of favorable alleles. Cruz et al. (2014) highlight that populations from crossing processes between high-GCA parents lead to a higher frequency of favorable alleles; thus, they increase the possibility of finding promising strains. Therefore, it is necessary to look for hybrid combinations in which at least one parent records a high GCA value in order to choose the population.

We searched for hybrid combinations presenting higher means in INSHEI, MFPR, MSD, MLLP, TNPP, MNSP, MNSPPL, MWS and GP, in which at least one parent presented positive GCA values. This was done because these are the variables demanding the lowest means; i.e., they suggest EMERG and BLOOM reduction. Interestingly, at least one of the parents recorded negative GCA values (Table 2).

EMERG showed non-additive gene effects (Table 1), and parent BGU-36 had the greatest GCA magnitude (Table 2). This outcome allows this parent to be recommended for breeding programs, because it can increment seed vigor, since it enables faster seedling emergence. Parents BGU-25, BGU-13, BGU-1 and BGU-36 recorded negative BLOOM

values: $g_{\rm I}$ -0.9714, -0.7047, -0.5381 and -0.2047, respectively (Table 2). This indicates that these parents help the development of crossing processes and of precocious individuals.

INSHEI is of fundamental importance for mechanized planting, since it facilitates harvest and prevents pods from coming in to contact with the soil; consequently, it causes less degradation in the product to be harvested (Velhoet al., 2017). According to our research, parents recommended for breeding programs to improve this variable are BGU-36 and IPR Campos Gerais, because they presented greater magnitudes: 12.2030 and 6.8252, respectively (Table 2).

Parents BGU-36 and BGU-7 presented magnitudes 0.3749 and 0.2735, respectively in MFPR; thus, they are recommended when the focus is to obtain taller plants (Table 2). The non-additive gene effects were more frequent than the additive ones, since they showed higher GCA (0.2074) than SCA (0.0417) values (Table 1). Similar and divergent values of additive and non-additive gene action in this variable were previously reported by Gonçalves et al. (2015) and Barelli et al. (2000), respectively.

Table 2 – Estimates of the general combining ability (GCA) effect on the 11 variables: mean number of days till emergence (EMERG), mean number of days till blooming (BLOOM), mean insertion height of the first pod (INSHEI), mean final plant height (MFPR), mean stem diameter (MSD), mean longitudinal length of pods (MLLP), mean number of pods per plant (MNPPL), mean number of seeds per plant (MNSP), mean number of seeds per pod (MNSP), mean weight of 50 seeds (MWS) and mean grain production (GP) in common bean. Standard deviation (SD) of the effects of two different parents.

·	Assessed variables										
GENOTYPES	EMERG	BLOOM	INSHEI	MFPR	MSD	MLLP	TNPP	MNSP	MNPPL	MWS	GP
IPR Uirapuru	-0.0031	1.5285	0.7907	0.1711	0.4588	0.0422	0.8198	0.4236	12.5071	-0.1300	0.5434
BGU-7	-0.0282	0.7507	-2.4891	0.2735	0.4336	-0.1988	1.3531	0.1217	12.9182	-0.8413	-0.2697
BGU-1	0.2224	-0.5381	-5.8202	-0.3561	-0.3155	0.6914	-0.9634	0.0228	-6.0817	0.3731	-0.3584
BGU-25	-0.0114	-0.9714	-5.7170	-0.2831	-0.0276	-0.0879	-0.4968	0.0695	-0.2484	-0.0583	-0.3040
BGU-13	0.1967	-0.7047	-5.7926	-0.4075	-0.1292	-0.2709	-0.0134	0.0261	-2.2595	-1.1480	-0.1846
BGU-36	-0.0928	-0.2047	12.2030	0.3749	-0.0372	-0.3029	0.3079	-0.6701	-14.2373	1.3836	0.9891
IPR Campos gerais	-0.0283	0.1396	6.8252	0.2272	-0.0469	0.1269	0.3912	0.0061	-0.5984	0.4210	-0.4156
DP (g _i - g _i)	0.0998	0.2152	1.8010	0.0508	0.0790	0.0657	0.8350	0.1315	3.9017	0.3487	0.7946

Parents IPR Uirapuru and BGU-7 presented high mean stem diameter estimates (Table 2); thus, these parents can contribute to genetic gains in this variable. Velhoet al. (2017) emphasized that this variable plays an essential role in plant support, as larger diameters avoid bedding and facilitate mechanized harvest processes. MLLP presented additive gene action prevalence; BGU-1 parents recorded 0.6914, IPR Campos Gerais, 0.1269and IPR Uirapuru, 0.042. Thus, they stood out among the others, due to their positive GCA values (Table 2). The additive gene action of this variable was previously evidenced in research conducted by Krause et al. (2012) with bean pods.

The non-additive gene effects were greater than the additive ones in TNPP, MNSP and MNSPPL. Different from results in our research, Gonçalves et al. (2015) reported additive gene effects of variables TNPP and MNSP. Barelli et al. (2000) highlighted that additive and non-additive gene effects are equally important in MNSPPL.

Among the aforementioned variables, parents IPR Uirapuru and BGU-7, in this order, were the ones presenting the greatest positive GCA magnitudes in TNPP, MNSP and MNSPPL (Table 2). Thus, when the idea is to gather parents for intercrossing to increase production components, these parents are the most recommended for inter populational enhancement with heterosis due to their non-additive gene effects.

Parent BGU-36 presented GCA values of 1.3836 in MWS and 0.9891 in GP; so it was recommended for breeding programs that focus on incrementing these variables. The non-additive action type expressed for these variables can be explained by the fact that some parents were previously selected in breeding programs focused on variables linked to production, as described by Cruz et al. (2014). Gonçalves-Vidigal et al. (2008) also found an effect of non-additive gene action on MWS and on GP in diallel crossing processes involving the following cultivars: BRSMG-Talismã, IPR Uirapuru, FT Soberano, BRS Campeiro, IAC Tybatã and IPR Juriti.

Thus, according to results recorded for the GCA effects, which set the mean performance of each parent in a series of crossing processes, besides being an estimate of the gene value, one can conclude that parents BGU-25 and BGU-36 should be favorites to be included in breeding programs focused on incrementing precocity. On the other hand, in case the aim of the breeding program is to increase yield, parents IPR Uirapuru, BGU-7 and BGU-36 are the most recommended, because they present the greatest magnitudes for variables associated with production. Either for precocity or yield, programs with such

parents may include intra- or interpopulational methods in order to obtain promising hybrids or superior segregating cultivars in future generations.

The estimate of combining ability effects (Table 3) aids in the selection of the best combinations. Selection should prioritize the most favorable SCA estimates involving at least one of the parents, which must have shown favorable GCA effects (Griffing, 1956). The best hybrid contributions in EMERG originated from BGU-1 x BGU-36 and BGU-13 x BGU-36, which recorded SCA values: -1.1435 and -1.0928, respectively. The hybrid combinations IPR Uirapuru x BGU-36 and BGU-1 xBGU-36 in BLOOM presented the greatest negative magnitudes: -1.667 and -1.600, respectively. Parent BGU-36 presented the best GCA result (Table 2). These results indicate that these specific combinations help diminish the number of days till germination and blooming.

The best combinations for INSHEI were BGU-7 x BGU-36 (9.8504), BGU-1 x IPR Campos Gerais (8.1293), IPR Uirapuru x BGU-36 (7.6395) and BGU-13 x BGU-36 (5.1229) (Table 3). Among the combinations mentioned above, the ones with parent 6 (BGU-36) presented more potential to be used in breeding programs, since this parent had the highest positive GCA estimate: 12.2030 (Table 2).

Fourteen hybrid combinations presented positive GCA values in MFPR, with emphasis on combinations BGU-13 x BGU-36 and BGU-1 x IPR Campos Gerais, which recorded the greatest magnitudes (Table 3). Cabralet al. (2011) reported that cultivars with the greatest plant height are desirable, since plant height has a positive and significant relation with grain yield.

If the aim of the breeding program is to develop greater stem diameter cultivars in MSD, combinations BGU-25 x BGU-13 (1.3689), IPR Uirapuru x BGU-7 (0.9081) and IPR Uirapuru x BGU-13 (0.3209) are the most viable options, since they present the highest positive SCA values (Table 3). Among all these combinations, the ones including parent IPR Uirapuru should be given priority due to its higher GCA estimates (Table 2).

In MLLP, combinations BGU-1 x BGU-25 (0.3408) and IPR Uirapuru x BGU-13 (0.3270) recorded the greatest SCA positive effects (Table 3). Thus, these combinations should be favored if the intension is to develop plants that have longer bean pods.

Combinations showing higher SCA in TNPP were BGU-25 x BGU-13, IPR Uirapuru x BGU-7, IPR Uirapuru x BGU-1, BGU-7 x BGU-1, BGU-13 x BGU-36, IPR Uirapuru x IPR Campos Gerais, BGU-1 x BGU-36 and BGU-36 x IPR Campos Gerais (Table 3). Among all these parents, IPR Uirapuru and BGU-7 presented the highest GCA estimates (Table 2) since they participated in 50% of the combinations recording the highest positive estimates and the second combination presenting the highest SCA estimate.

Eleven combinations presented positive SCA values in MNSPPL, and the greatest magnitudes among them were shown by combinations BGU-25 x BGU-36, BGU-1 x BGU-13, IPR Uirapuru x BGU-1 and BGU-1 x IPR Campos Gerais (Table 3). However, only parents IPR Uirapuru and BGU-7 presented positive GCA values (Table 2). This made the best combination, composed of these two parents; this combination recorded an SCA value 35.4013 (Table 3). Griffing (1956) emphasizes that the best combination must be the one presenting the highest SCA, whose parents presented high GCA scores.

Parent BGU-36 showed the highest positive GCA magnitude in mean weight of 50 seeds (Table 2). This result was expressed in SCA, in which combinations between BGU-7 and BGU-13, and BGU-36 presented the greatest SCA magnitudes: 2.7246, 2.1782, respectively (Table 3).

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Table 3 – Estimates of specific combining ability effects (S_{ii} and S_{ji}) on the 11 variables: mean number of days to emergence (EMERG), mean number of days to blooming (BLOOM), mean insertion height of the first pod (INSHEI), mean final plant height (MFPR), mean stem diameter (MSD), mean length of the pods (MLLP), mean number of pods per plant (MNPP), mean number of seeds per plant (MNSPL), mean number of seeds per pod (MNSP), mean weight of 50 seeds (MWS) and mean grain production (GP) in the common bean. Standard deviations (SD) of the effects of two F_1 with, and without, common parent between two parents resulting from diallel crossing between seven parents.

Hybrid combinations	Assessed variables											
(s _{ii} ands _{ij})	EMERG	BLOOM	INSHEI	MFPR	MSD	MLLP	TNPP	MNSP	MNPPL	MWS	GP	
1x1	-0.2077	-0.4000	-5.4481	-0.4487	-0.2407	-0.3983	-4.7986	-0.1410	-25.6375	-1.3071	-4.6485	
1x2	-0.3325	0.2777	-2.2182	-0.1291	0.9081	0.1028	7.7681	0.0109	35.4013	-0.5608	5.5026	
1x3	0.3168	1.9666	-4.0371	0.3976	0.1707	0.2706	4.1347	0.5557	11.7013	2.0357	4.1888	
1x4	0.3257	-0.2000	3.2596	0.3446	-0.5277	-0.0230	-0.3319	-0.0649	6.3180	-0.2568	-0.6217	
1x5	0.1754	0.5333	4.9852	0.3870	0.3209	0.3270	-0.6653	0.2225	5.9291	1.3819	1.1136	
1x6	-0.0429	-1.6667	7.6395	0.0435	-0.3319	0.1620	-2.0208	-0.5083	-20.5431	0.7482	-0.4083	
1x7	-0.0271	-0.1112	1.2673	-0.1158	-0.0589	-0.0429	0.7125	0.0665	12.4680	-0.7342	-0.4724	
2x2	-0.0075	0.3555	-4.6884	-0.2586	-0.0152	-0.2590	-3.3653	0.0757	-18.1098	-1.3984	-3.7792	
2x3	0.1579	-0.1556	-1.1572	0.2882	-0.1326	0.1657	2.9514	-0.2294	14.3402	1.2220	2.4045	
2x4	-0.1162	0.0777	4.6395	0.3542	-0.2935	0.2251	-0.3653	0.0920	1.3569	-0.1694	-0.1269	
2x5	0.7505	0.9111	1.0151	0.2425	-0.3233	0.0841	-0.1486	-0.0236	3.4680	0.7772	-1.7263	
2x6	0.0072	-0.4889	9.8504	0.1331	-0.3077	-0.0769	-3.1042	-0.2304	-17.9542	2.7246	1.5614	
2x7	-0.4520	-1.3334	-2.7528	-0.3713	0.1793	0.0172	-0.3708	0.2304	-0.3931	-1.1969	-0.0568	
3x3	0.7232	0.5333	-1.0761	-0.3540	0.2306	-0.6095	-1.1819	-0.7235	-13.6098	-0.6246	-2.0004	
3x4	-0.7749	-0.8334	2.3706	-0.2230	-0.1873	0.3408	-2.6486	0.2569	-4.7931	-0.5620	-0.9248	
3x5	-0.0831	-0.1000	-2.6538	-0.3747	-0.1842	-0.0222	-1.8319	0.8203	-15.0319	-0.6803	-2.9226	
3x6	-1.1435	-1.6000	-0.4995	0.2119	0.0670	0.2698	0.6125	-0.3505	15.6958	-0.7780	2.8961	
3x7	0.0803	-0.3445	8.1293	0.4385	-0.1950	0.1940	-0.8542	0.3943	5.3069	0.1160	-1.6011	
4x4	0.5590	1.3000	-0.3826	-0.4600	-0.0628	-0.2828	-2.3153	-0.2868	-16.0764	-0.3164	-2.3866	
4x5	0.4507	1.0333	-3.9070	-0.2467	1.3689	-0.0988	9.1513	-0.4244	32.0347	-0.5188	5.7365	
4x6	-0.3846	-1.5667	-8.3777	0.2809	0.1451	0.1852	-1.1042	0.8339	0.5625	0.1836	0.0542	
4x7	-0.6188	-1.1112	2.7801	0.4105	-0.3799	-0.0637	-0.0708	-0.1194	-3.3264	1.9561	0.6615	
5x5	-0.1745	-0.6334	-4.1315	-0.4334	-0.3799	-0.2058	-3.2319	-0.2289	-13.5042	-1.8411	-4.4773	
5x6	-1.0928	-1.0334	5.1229	0.5002	0.1173	-0.1088	0.9625	-0.3397	-0.7764	2.1782	3.5769	
5x7	0.1480	-0.0778	3.7006	0.3589	-0.3063	0.2303	-1.0042	0.2031	1.3847	0.5438	3.2167	
6x6	1.3719	3.2666	-7.7728	-0.4473	0.2659	-0.3018	2.1069	0.1905	8.8013	-1.6174	-4.1815	
6x7	-0.0873	-0.1778	1.8100	-0.2746	0.0129	0.1723	0.4403	0.2125	5.4125	-1.8219	0.6829	
7x7	0.4784	1.5777	-7.4672	-0.2229	0.3739	-0.2536	0.5736	-0.4949	-10.4264	0.6207	-1.2153	
SD (sii-sij)	0.2232	0.4813	4.0272	0.1136	0.1768	0.1470	1.8672	0.2941	8.7245	0.7798	1.7767	
SD (sij-sik)	0.2824	0.6088	5.0941	0.1437	0.2236	0.1860	2.3619	0.3720	11.0357	0.9863	2.2474	
SD (sij-skl)	0.2642	0.5694	4.7651	0.1344	0.2092	0.1739	2.2093	0.3480	10.3229	0.9226	2.1023	

*Parents: 1-IPR Uirapuru, 2-BGU-7, 3-BGU-1, 4-BGU-25, 5-BGU-13, 6-BGU-36 e 7-IPR Campos Gerais.

Twelve hybrid combinations showed positive SCA values in GP (Table 3). Among these combinations, the following have one parent with positive GCA values: IPR Uirapuru x BGU-7, IPR Uirapuru x BGU-1, IPR Uirapuru x BGU-13, BGU-7 x BGU-36, BGU-1 x BGU-36, BGU-25 x BGU-36, BGU-13 x BGU-36 and BGU-36 x IPR Campos Gerais. All these combinations tend to reinforce the selection of promising strains in order to produce high grain weight. However, the combinationBGU-25 x BGU-13 presented the highest SCA, even though none of the parents presented positive GCA values. The SCA value in this combination can be explained by a high genetic complementation degree between parents (Idahosa and Alika, 2013).

Based on the set of assessed variables, we conclude that the hybrid combinations IPR Uirapuru x BGU-7, IPR Uirapuru x BGU-1, BGU-7 x BGU-1, BGU-25 x BGU-13 and BGU-13 x BGU-36 are the most favorable for the generation of segregating populations for the production of highly productive strains.

CONCLUSIONS

Additive gene effects prevailed in the expression of INSHEI and MLLP. Non-additive gene effects prevailed in the expression of EMERGE, MFPR, TNPP, MNSP, MNSPPL, MWS and GP.

Local varieties presented good general and specific combinations.

Parents IPR Uirapuru and BGU-36 presented the best combining ability in most of the variables, and hybrid combinations IPR Uirapuru x BGU-7, IPR Uirapuru x BGU-1, BGU-7 x BGU-1, BGU-25 x BGU-13 and BGU-13 x BGU-36 were the most promising for the development of highly productive strains.

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CONFLICT OF INTEREST

All authors declare they have no conflicts of interest.

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