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Combining ability estimates for spike characters in F_1 hybrids developed through diallel crosses among macaroni wheat (*Triticum durum* Desf.) genotypes

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Abstract

The present study was conducted to obtain information on combining ability effects for spike related traits in durum wheat and to select suitable parents and population for designing an effective wheat breeding programme. The experimental material designed in an 8×8 diallel set to synthesize 28 hybrids excluding reciprocals. Analysis of variance (ANOVA) revealed the presence of significant variance due to General Combining Ability (GCA) and Specific Combining Ability (SCA) among the parents and F_1 hybrids for all the traits studied. Combining ability analysis revealed the involvement of both additive and non additive gene action in the inheritance of most of the traits. On the basis of GCA/SCA effects and *per se* performance, parents PDW 300 followed by RAJ 1555 and HI 8653 were found good general combiners for spike characters. Similarly, F_1 's crosses PDW 300 \times HI 8653 and PDW 300 \times RAJ 1555 were found good specific combiners for spike related traits. Therefore, these parents can be considered as best combiners and their crosses are expected to be worthwhile for subsequent utilization in genetic enhancement of macaroni wheat.

Keywords: Durum wheat, F_1 , GCA, SCA, spike characters

1. Introduction

Global demand for wheat is growing at approximately 2% year⁻¹, twice the current rate of gain in genetic yield potential (Skovmand and Reynolds, 2000) [21]. Durum wheat is the second most important wheat species, plays an important role in Indian economy being the staple food of the population. It is a naturally bi-fortified food and used for making large numbers of products including 'nutritive chapatti' and semolina, (Prasad *et al* 2014) [15]. Historically durum has received insufficient attention from plant breeders. Therefore, efforts to increase yield in durum wheat will not only support to the overall wheat production of this country but it will also help to meet the food requirements of the burgeoning population of country like India. The yield levels of durum wheat are low compared to bread wheat under both irrigated and rain fed conditions (Sharma *et al.*, 2004) [12,18]. Therefore, it is necessary to improve durum wheat with high yielding capacity. This could be only realized via intensive and effective breeding programmes. Thus, careful selection of the parents is the first important step in the process for the development of superior high yielding cultivars.

The wheat inflorescence spike has a direct bearing on grain yield. It makes an important contribution to the final grain yield through photosynthesis (Araus *et al.*, 1993) [3]. Thus any improvement of spike characteristics through selection and breeding would help improve plant⁻¹ productivity (Iqbal and Khan, 2006) [9]. For breaking the yield barrier level and make durum wheat cultivation more attractive it is now necessary to explore alternative approaches to increasing wheat production.

Combining ability analysis provides information on additive and non additive variances, which are important to decide the proper parents for hybridization and to produce superior hybrids. The present investigation was therefore an attempt to obtain information on the extent of combining ability for the spike related traits and to select suitable parents and hybrids for designing an effective wheat breeding programme.

2. Materials and Methods

The experimental material comprising of a set of 28 hybrids, developed from eight diverse parents (PDW 300, NIDW 295, AKDW-2997-16, RAJ 6560, DBP-01-11, DBP-01-12, RAJ 1555, HI 8653) of durum wheat using direct straight diallel mating system excluding reciprocals during *Rabi* 2011-12 (45 to 55 seeds). 8 parents and their 28 F_1 s hybrids were laid out in Randomized Block Design with three replications evaluated during *Rabi* 2012-13, at the

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Field Experimentation Centre of Department of Genetics and Plant Breeding Allahabad School of Agriculture, Sam Higginbottom Institute of Agriculture, Technology and Sciences, Allahabad, Uttar Pradesh, with Recommended doses of fertilizers at @ 120 kg N+60 kg P₂O₅+40 kg K₂O ha⁻¹ were applied in the experimental field along with five irrigations at all critical stage to raised a healthy crop.

The observations were recorded on five randomly selected competitive plants from parents as well as in F₁ hybrids for spike length, spikelet's spike⁻¹, spike weight productive tiller⁻¹, grain weight spike⁻¹, grains spike⁻¹, grains spikelet⁻¹, spike density, spike harvest index in every plot. Data on these quantitative traits in each replication were subjected to appropriate biometrical approaches. The combining ability analysis was worked out by the procedure suggested by Griffing's (1956) Method II, Model I. The mathematical model for the combining ability analysis is assumed to be:

$$Y_{ijkl} = \mu + g_i + g_j + s_{ij} + 1/bc \sum_i \sum_j \sum_k \sum_l$$

$$(ij=1, 2, 3, \dots, n);$$

$$k=1, 2, 3, \dots, b;$$

$$l=1, 2, 3, \dots, c)$$

Where, Y_{ijkl}, Mean of i^xjth genotype in kth replication; μ , the population mean; g_i, the GCA effect of ith parent; g_j, the GCA effect of jth parent; s_{ij}, the specific combining ability (SCA) effect for the cross between ith, jth parent such that s_{ij}=s_{ji}; $\sum_i \sum_j \sum_k \sum_l$, the environmental effect associated with the ijklth individual observation on ith individual in the kth block with ith as female parent and jth as male parent. The usual restrictions such as $\sum_i g_i = 0$ and $\sum_{s_{ij}} = s_{ji} = 0$ (for each i) were imposed. The analysis of variance (ANOVA) table for combining ability with expectation of mean sum of squares (M.S.) is shown in Table 1.

$$[\sum (Y_i + Y_{ii})^2 - (4/n) Y^2] S_s = \text{Sum of squares due to SCA,}$$

$$S_s = \sum < \sum Y_{ij}^2 - 1/ (n+2) [\sum (Y_i + Y_{ii})^2 + 2/ (n+1) (n+2) Y^2],$$

$$m_e = Me1/r$$

Where Y_i, Total of the array involving ith as a female parent; Y_{ii}, the value of the ith of the array; Y_{...}, the grand total; Y_{ij}, the value of i x jth cross; Me1, the error M.S. obtained from main ANOVA by Fone SCA and Patterson., 1968.

The components of variances were estimated as suggested by (Singh *et al.*, 1990) in the following ways:

$$\text{GCA expected M.S.} = \sigma^2 e + (n+2)/(n-2) \sigma^2 g_i$$

$$\text{SCA expected M.S.} = \sigma^2 e + 2/ (n-1)^2 s_{ij}$$

2.1. Estimates of various effects

The various effects were estimated as follows:

$$\text{GCA effect of } i^{\text{th}} \text{ parent} = g_i = 1/ (n+2) [(Y_i + Y_{ii}) - 2/n Y_{..}]$$

$$\text{SCA effect of } ij^{\text{th}} \text{ cross} = S_{ij} = Y_{ij} - 1/ (n+2) [Y_i + Y_j + Y_{ij}] + 2Y_{..} / (n+1) (n+2) Y_{..}]$$

Where, g_i and s_{ij}, the estimates of the GCA and SCA effects, respectively; n, Y_i, Y_{ii}, Y_{..} and Y_{ij}, the same as explained earlier; Y_j, total of the arrays involving jth parent as a male; Y_{ij}, the value of the jth parent in the array.

3. Results and Discussion

Analysis of variance for combining ability revealed that the variance due to GCA and SCA were significant for all the traits (Table 1). The extent of mean square due to GCA was higher than SCA for all the characters except spike length, which indicates existence of considerable genetic variability in the parental lines included in the present study and involvement of both additive and non-additive gene effects in the inheritance of these traits (Jaiswal *et al.*, 2013; [11] Joshi *et al.*, 2004) [12].

The ratio of GCA and SCA variances as less than one and degree of dominance more than one for all traits indicated greater role of non additive components in the inheritance of these characters.

3.1. General combining ability effects

The information regarding general combining ability (GCA) effects of the parents is of prime importance because it helps in successful prediction of genetic potentiality, which would give desirable individuals in subsequent segregating populations. Estimates of GCA effects (Table 2) showed that it was not possible to pick up a good general combiner for all the spike characters because the combining ability of the parents was not consistent for all yield attributing components (Kashif *et al.*, 2008 [13]; Dhadhal and Dobariya, 2006) [6]. The GCA effects of the parents along with parental means (Table 2) indicated that there was a close and desirable relationship between parents and their GCA effects almost all of the studied characters. On the basis of overall performance, two parent PDW-300 followed by RAJ 1555 was found to be the desirable combiner for spike length, spikelets spike⁻¹, spike weight productive tiller⁻¹, grain weight spike⁻¹, grains spike⁻¹, spike density and spike harvest index. Similarly parent DBP-01-11 found to be the best combiner for more numbers of grains spikelets⁻¹. Thus, the perfect relationship could be established between *per se* performance and GCA effects of the parents. It is therefore suggested that these parental lines may be used in hybridization programme to have the superior recombinants for respective traits. Similar findings in durum wheat have been reported by Barot *et al.* (2014) [4].

3.2. Specific combining ability effects

The magnitude of specific combining ability (SCA) effects is of vital importance in selecting cross combination with higher probability of generating transgressive segregates. Significant yield performance in the specific crosses was due to the involvement of best general combiners in our study. The desirable SCA effects are presented in Table 3. The significant SCA effects with the mean performance of the genotypes indicated that among overall 28 crosses, hybrid PDW 300×HI 8653 had maximum spike length with a perfect correlation of SCA performance also. Grain weight is an essential component contributing for yield production with grains spike⁻¹ which directly determines the yield potential of a genotype. Cross PDW 300×HI 8653 performed well in all the three yield attributing traits viz., spike length, grain weight spike⁻¹ and grains spike⁻¹ with higher *per se* performance and SCA effects. Similar finding in wheat has been reported by Ismail *et al.* (2006) [10], Khan and Ali (1998) [2, 17] and Shahzad *et al.* (1998) [17]. Spike Harvest Index (SHI) is the dry weight (at anthesis) to total dry weight ratio at physiological maturity. It shows the proportion of mobilized dry matter to spike in a genotype (Donalson, 1996) [7]. A marked variation is evident for average spike harvest index among the parents and hybrids. In the present study cross RAJ 1555×HI 8653 represent the highest significant positive SCA effects for spike harvest index. Cross AKDW-2997-16×DBP-01-12 found to be common good and significant cross combination for spike density and spike harvest index. Grains spikelets⁻¹ is the most important yield determinant of wheat (Sinclair and Jamieson, 2006). The individual comparison of mean performance and SCA effects represented the significant and correlated value by the cross combination PDW 300×RAJ 1555. Spikelets spike⁻¹ and spike weight productive tiller⁻¹ is an important and effective yield

component and greater number could results in more grains spike⁻¹. The hybrid RAJ 6560×DBP-01-12 found to be best for spikelets spike⁻¹ followed by PDW 300×HI 8653 for both spike characters having significant and desirable SCA effects indicative of best yielding cross combination. Similar finding has also reported by Adel *et al.* (2013) [1], Mishra *et al.* (1996) [14] and Chowdhry *et al.* (1999) [5] in wheat.

3.3. Per se performance and SCA common superior cross combinations

The ranking of specific cross combination on the basis of their *per se* performance and SCA effects computed in Table 4. In the present study, the common good crosses on the basis of *per se* performance and SCA effects were PDW 300×HI 8653, PDW 300×RAJ 1555 and PDW 300×DBP-01-12 for spike length, RAJ 6560× DBP-01-12 and PDW 300×HI 8653 for spikelets spike⁻¹, RAJ 6560×HI 8653 AND RAJ

6560×DBP-01-12 for spike weight productive tiller⁻¹; PDW 300×HI 8653 for grain weight spike⁻¹; PDW 300×HI 8653 and RAJ 6560×DBP-01-12 for grains spike⁻¹; PDW 300×RAJ 1555 for grains spikelet⁻¹; DBP-01-11×HI 8653 for spike density and RAJ 1555×HI 8653 for spike harvest index. However some crosses showed desirable SCA effects for more than one characters. Cross combination PDW 300×HI 8653 exhibited significant and desirable SCA effects for 4 traits viz.; spikelets spike⁻¹, spike length, grain weight spike⁻¹ and grains spikelet⁻¹. Similarly cross combination RAJ 6560×DBP-01-12 recorded desirable SCA effects for 3 spike characters, spikelets spike⁻¹, spike weight productive tiller⁻¹ and grains spike⁻¹. It is thus evident that high yielding varieties involved in the crosses were predominately responsible for enhancing the yield. Similar findings in wheat have been reported by Raiyani *et al.* (2015), Adel *et al.* (2013) [1], and Ismail *et al.* (2006) [10].

Table 1: ANOVA for combining ability (GCA and SCA) for 8 characters in 8×8 diallel crosses of durum wheat

Sources of variance	Degree of freedom	Spike length	Spikelets spike ⁻¹	Spike weight productive Tillers ⁻¹	Grain weight spike ⁻¹	Grains spike ⁻¹	Grains spikelets ⁻¹	Spike density	Spike harvest index
Parents	7	0.09**	5.17**	0.34**	0.31**	71.71**	0.09**	0.03**	23.90**
Crosses	28	0.28**	1.96**	0.09**	0.10**	42.04**	0.06**	0.01**	22.10**
Error	70	0.00	0.04	0.00	0.33	0.45	0.00	0.00	2.14
GCA variance		17.79	123.06	102.35	95.62	156.54	19.05	33.3	11.14
SCA variance		55.62	46.74	29.53	31.8	91.78	13.1	18.08	10.3
GCA/SCA variance		0.03	0.26	0.3	0.3	0.17	0.14	0.18	0.1

*Significant at (p=0.05) level of significance; **, significant at (p=0.01) level of significance; GCA, general combining ability; SCA, specific combining ability

Table 2: Estimation of GCA effects of 8 parents along with their mean performance for 8 characters in F₁ hybrids of wheat

Characters	Spike length		Spikelets spike ⁻¹		Spike weight productive tillers ⁻¹		Grain weight spike ⁻¹		Grains spike ⁻¹		Grains spikelets ⁻¹		Spike density		Spike harvest index	
	GCA effect	Parent Mean	GCA effect	Parent mean	GCA effect	Parent mean	GCA effect	Parent mean	GCA effect	Parent mean	GCA effect	Parent mean	GCA effect	Parent mean	GCA effect	Parent mean
Parents	0.15**	8.27	1.23**	22.73	0.29**	3.46	0.31**	2.54	3.10**	63.13	-0.07**	2.77	0.10**	2.74	3.36**	73.31
PDW-300	-0.10**	8.07	-0.98**	19.90	-0.33**	2.29	-0.28**	1.64	-3.96**	53.47	-0.07**	2.68	-0.09**	2.44	-0.73	71.78
NIDW-295	-0.11**	8.20	-0.19**	21.60	-0.06**	3.45	-0.05**	2.55	-1.51**	60.93	-0.03	2.82	0.01	2.63	-0.29	73.98
AKDW-2997-16	0.06**	7.73	-0.19**	19.20	0.04*	2.94	0.00	1.95	-3.41**	54.33	-0.13**	2.83	-0.04**	2.48	-0.38	66.44
DBP-01-11	-0.10**	8.10	-0.55**	20.13	-0.12**	3.19	-0.14**	2.41	0.60**	59.97	0.13**	2.98	-0.02*	2.56	-1.72**	75.55
DBP-01-12	0.00	8.07	-0.31**	20.57	-0.02	3.19	-0.04*	2.29	0.70**	58.53	0.09**	2.84	-0.04**	2.55	-1.14*	71.71
RAJ-1555	0.07**	8.37	0.80**	23.47	0.17**	3.64	0.15**	2.85	2.03**	65.00	-0.03	2.60	0.06**	2.74	0.58	75.68
HI-8653	0.04	8.17	0.19**	22.03	0.04*	3.39	0.04*	2.52	2.45**	62.07	0.11**	2.81	0.01	2.69	0.33	74.70

*, Significant at (p=0.05) level of significance; **, Significant at (p=0.01) level of significance

Table 3: Estimation of SCA effects and corresponding mean performance of 8 characters in 8×8 diallel cross of wheat

Characters	Spike length		Spikelets spike ⁻¹		Spike weight productive tillers ⁻¹		Grain weight spike ⁻¹		Grains spike ⁻¹		Grains spikelets ⁻¹		Spike density		Spike harvest index		
	SCA effect	Parent mean	SCA effect	Parent mean	SCA effect	Parent Mean	SCA effect	Parent Mean	SCA effect	Parent mean	SCA effect	Parent mean	SCA effect	Parent mean	SCA effect	Parent mean	
Crosses	-0.46**	8.07	-0.32	20.87	0.22**	3.41	0.02	2.35	0.2	63.20	-0.22**	2.68	0.10**	2.58	-2.69	70.67	
PDW 300×NIDW-295	-0.46**	8.07	-0.32	20.87	0.22**	3.41	0.02	2.35	0.2	63.20	-0.22**	2.68	0.10**	2.58	-2.69	70.67	
PDW 300×AKDW 2997-16	-0.72**	7.80	-1.38**	20.60	-0.26**	3.20	-0.30**	2.25	-3.41**	62.03	0.05	2.99	0.06	2.64	-3.55*	70.25	
PDW 300×RAJ 6560	-0.42**	8.27	-2.24**	19.73	-0.39**	3.17	-0.17**	2.44	-	14.68**	48.87	-0.37**	2.47	-0.15**	2.38	3.25*	76.97
PDW 300×DBP-01-11	0.03	8.57	0.28	21.90	0.02	3.42	0.11*	2.58	0.6	68.17	0.01	3.11	0.01	2.55	3.38*	75.75	
PDW 300×DBP-01-12	0.77**	9.40	1.71**	23.57	0.42**	3.92	0.40**	2.97	8.94**	76.60	0.18**	3.24	-0.03	2.50	2.67	75.61	
PDW 300×RAJ 1555	0.83**	9.53	1.16**	24.13	0.25**	3.94	0.28**	3.04	11.01**	80.00	0.38**	3.31	-0.10**	2.52	2.46	77.13	
PDW 300×HI 8653	1.00**	9.67	2.11**	24.47	0.42**	3.99	0.43**	3.08	11.19**	80.60	0.21**	3.29	-0.05	2.53	2.75*	77.17	
NIDW 295×AKDW-2997-16	0.43**	8.70	1.10**	20.87	0.35**	3.19	0.38**	2.35	1.85**	60.23	-0.05	2.88	0	2.40	3.92**	73.64	
NIDW 295×RAJ 6560	0.46**	8.90	0.1	19.87	0.18**	3.12	0.29**	2.31	-0.55	55.93	-0.02	2.81	-0.07*	2.28	4.41**	74.03	
NIDW 295×DBP-01-11	0.48**	8.77	0.46*	19.87	0.01	2.80	0.08	1.96	2.30**	62.80	0.07	3.16	-0.10**	2.26	2.27	70.55	

NIDW 295×DBP-01-12	0.29**	8.67	-0.55**	19.10	0.06	2.94	-0.07	1.91	-1.50*	59.10	0.03	3.09	-0.14**	2.20	-3.56*	65.30
NIDW 295×RAJ 1555	-0.42**	8.03	-1.60**	19.17	-0.16**	2.91	-0.30**	1.87	1.21	63.13	0.36**	3.29	-0.06	2.38	-4.31**	66.27
NIDW 295×HI 8653	-0.35**	8.07	-1.05**	19.10	-0.09	2.86	-0.20**	1.87	1.45*	63.80	0.26**	3.33	-0.04	2.35	-5.07**	65.26
AKDW 2997-16×RAJ 6560	0.44**	8.87	1.54**	22.10	0.25**	3.46	0.31**	2.55	9.84**	68.77	0.24**	3.11	0.05	2.49	3.62*	73.69
AKDW 2997-16×DBP-01-11	-0.14*	8.13	-0.67**	19.53	-0.09	2.97	-0.22**	1.88	-0.48	62.47	0.07	3.19	-0.06*	2.40	-5.36**	63.36
AKDW 2997-16×DBP-01-12	-0.23**	8.13	0.55**	21.00	-0.1	3.05	0.05	2.25	-3.14**	59.90	-0.24**	2.85	0.13**	2.58	3.99**	73.29
AKDW 2997-16×RAJ 1555	-0.1	8.33	-1.19**	20.37	-0.42**	2.92	-0.45**	1.94	0.6	64.97	0.22**	3.19	-0.10**	2.44	-4.40**	66.63
AKDW 2997-16×HI 8653	0.43**	8.84	-2.01**	18.93	-0.41**	2.81	-0.46**	1.82	-5.46**	59.33	0.02	3.13	-0.35**	2.14	-5.88**	64.89
RAJ 6560×DBP-01-11	-0.08	8.37	-1.20**	19.00	-0.26**	2.89	-0.44**	1.72	-0.98	60.07	0.13*	3.16	-0.14**	2.27	-9.31**	59.33
RAJ 6560×DBP-01-12	0.56**	9.10	2.22**	22.67	0.42**	3.67	0.41**	2.67	11.03**	72.17	0.18**	3.18	0.09**	2.49	3.57*	72.78
RAJ 6560×RAJ 1555	0.12	8.73	0.38	21.93	-0.01	3.43	-0.03	2.42	-7.27**	55.20	-0.36**	2.51	0.02	2.51	-0.48	70.45
RAJ 6560×HI 8653	0.65**	9.23	1.92**	22.87	0.55**	3.87	0.34**	2.68	8.00**	70.90	0.09	3.10	0.03	2.47	2.01	72.69
DBP-01-11×DBP-01-12	0.15	8.53	-0.89**	19.20	-0.27**	2.83	-0.32**	1.79	3.48**	68.63	0.32**	3.57	-0.16**	2.25	-4.51**	63.37
DBP-01-11×RAJ 1555	0.61**	9.07	1.43**	22.63	0.21**	3.50	0.07	2.38	4.18**	70.67	-0.01	3.12	-0.01	2.49	-1.62	67.97
DBP-01-11×HI 8653	-0.66**	7.77	0.01	20.60	-0.02	3.14	-0.07	2.13	1.09	68.00	0.03	3.29	0.19**	2.65	-1.34	68.00
DBP-01-12×RAJ 1555	-0.48**	8.07	-1.57**	19.87	0.02	3.40	-0.17**	2.24	-3.21**	63.37	0.08	3.17	-0.03	2.46	-4.42**	65.75
DBP-01-12×HI 8653	-0.22**	8.30	-1.96**	18.87	-0.56**	2.70	-0.45**	1.85	-2.17**	64.83	0.20**	3.43	-0.17**	2.27	-4.28**	65.64
RAJ 1555×HI 8653	-0.06	8.53	-0.44*	21.50	-0.04	3.41	0.13*	2.62	-0.7	67.63	0.06	3.16	-0.02	2.51	5.19**	76.83

*, Significant at ($p=0.05$) level of significance; **, Significant at ($p=0.01$) level of significance

Table 4: Ranking of good cross combination on the basis of *per se* performance and their SCA effect in a 8×8 diallel cross of wheat

Characters	Parents with higher <i>Per se</i> Performance	Good cross combination	Superior Common Cross Combinations
Spike length	[1] PDW 300×HI 8653	PDW 300×HI 8653 PDW 300×RAJ 1555 PDW 300×DBP-01-12	PDW 300×HI 8653 PDW 300×RAJ 1555
	[2] PDW 300×RAJ 1555		
	[3] PDW 300×DBP-01-12		
	[4] RAJ 6560×DBP-01-12		
	[5] RAJ 6560×HI 8653		
Spikelets spike ⁻¹	[1] PDW 300×HI 8653	RAJ 6560×DBP-01-12 PDW 300×HI 8653 RAJ 6560×HI 8653 AKDW 2997-16×NIDW 295	RAJ 6560×DBP-01-12 PDW 300×HI 8653 RAJ 6560×HI 8653
	[2] PDW 300×RAJ 1555		
	[3] PDW 300×DBP-01-12		
	[4] RAJ 6560×HI 8653		
	[5] RAJ 6560×DBP-01-12		
Spike weight productive tillers ⁻¹	[1] PDW 300×HI 8653	RAJ 6560×HI 8653 RAJ 6560×DBP-01-12	RAJ 6560×HI 8653 RAJ 6560×DBP-01-12
	[2] PDW 300×RAJ 1555		
	[3] PDW 300×DBP-01-12		
	[4] RAJ 6560×HI 8653		
	[5] RAJ 6560×DBP-01-12		
Grain weight spike ⁻¹	[1] PDW 300×HI 8653	PDW 300 x HI 8653 RAJ 6560×DBP-01-12	PDW 300 x HI 8653
	[2] PDW 300×RAJ 1555		
	[3] PDW 300×DBP-01-12		
	[4] RAJ 6560×HI 8653		
	[5] RAJ 6560×DBP-01-12		
Grains spike ⁻¹	[1] PDW 300×HI 8653	PDW 300×HI 8653 RAJ 6560×DBP-01-12	PDW 300×HI 8653 RAJ 6560×DBP-01-12
	[2] PDW 300×RAJ 1555		
	[3] PDW 300×DBP-01-12		
	[4] RAJ 6560×DBP-01-12		
	[5] RAJ 6560×HI 8653		
Grains spikelets ⁻¹	[1] DBP-01-11×DBP-01-12	PDW 300×RAJ 1555 NIDW 295×RAJ 1555	PDW 300×RAJ 1555
	[2] DBP-01-12×HI 8653		
	[3] PDW 300×HI 8653		
	[4] PDW 300×RAJ 1555		
Spike density	[1] DBP-01-11×HI 8653	DBP-01-11×HI 8653 AKDW 2997-16×DBP-01-12	DBP-01-11×HI 8653
	[2] PDW 300×AKDW 2995-16		
	[3] PDW 300×DBP-01-11		
	[4] PDW 300×HI 8653		
Spike harvest index	[1] PDW 300×HI 8653	RAJ 1555×HI 8653 NIDW 295×RAJ 6560	RAJ 1555×HI 8653
	[2] PDW 300×RAJ 1555		
	[3] PDW 300× RAJ 6560		
	[4] RAJ 1555×HI 8653		

4. Conclusion

The GCA and SCA variances were significant for almost all the characters, indicating the importance of both additive and non additive gene effects. On the basis of GCA effects it was observed that PDW 300 was best general combiner for almost all the spike characters, followed by RAJ 1555 and HI 8653. The best specific combiners PDW 300×HI 8653 was considerable for spike length, spikelets spike⁻¹, grain weight spike⁻¹, grains spike⁻¹, the cross PDW 300×RAJ 1555 was found desirable for grains spikelet⁻¹ and spike length, and cross DBP-01-11×HI 8653 for spike density and cross RAJ 1555×HI 8653 for spike harvest index. The presence of parents having significant GCA effects, indicating additive×additive gene action played an important role for the expression of all the spike traits.

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