



Genetic Variability, Heritability and Genetic Advance for yield and its contributing characters in Rice (*Oryza sativa* L.)

Key words : Gentic Advance, Heritability and Variability

Rice is the most important staple food crop providing 43% of calorie requirement for more than 70% of Indian population. For the formulation of successful breeding programme, knowledge of the nature and magnitude of the genotypic and phenotypic variability present in any crop species plays a vital role. Therefore, present study was undertaken to find the genetic variability, heritability and genetic advance of yield and yield component traits in rice.

In the present investigation four wild abortive source CMS lines and thirteen elite *indica/indica* derivatives were crossed in line x tester design at Andhra Pradesh Rice Research Institute and Regional Agricultural Research Station, Maruteru during *khari*, 2010. The crosses along with their parents and three checks (*viz.*, MTUHR 2089-hybrid check, MTU 1075 and MTU 1010-varietal checks) were evaluated in a randomized block design with two replications during *rabi*, 2010-11.

Twenty eight days old seedlings were transplanted at a spacing of 20x15 cm and the plots were periodically top dressed with urea, watered regularly and necessary plant protection measures were taken against pests and diseases. Observations were recorded on ten randomly selected plants for plant height, number of ear bearing tillers plant⁻¹, panicle length, number of filled grains panicle⁻¹, spikelet fertility per cent and grain yield plant⁻¹ whereas data on days to 50% flowering and test weight were recorded on plot basis. The mean of replications was used for statistical analysis. Genotypic and phenotypic co-efficient of variation (Burton and Devane, 1953), heritability and genetic advance as per cent of mean (Johnson *et al.*, 1955) were estimated.

Analysis of variance indicated the existence of highly significant differences among the genotypes for all the characters studied (Table 1). The phenotypic coefficient of variance (PCV) was

slightly higher in magnitude than genotypic coefficient of variance (GCV) for all the characters studied (Table 2) indicating the influence of environment. Among the yield and yield attributing characters, PCV and GCV were observed to be highest (26.89 and 24.41) only for grain yield. Similar high PCV and GCV for grain yield was also reported by Karthikeyan *et al.* (2007), Krishna *et al.* (2008), Prasad *et al.* (2009), Mohan Lal and Chauhan (2011), Siva Parvathi *et al.* (2011) and Shiva Prasad *et al.* (2011). Value of PCV and GCV were found to be moderate for number of ear bearing tillers plant⁻¹ (14.67 and 11.78), spikelet fertility (10.99 and 10.42) and test weight (12.94 and 12.48). Similar findings were given by Saidaiah *et al.* (2010) for spikelet fertility and Satish (2000) for test weight. For the characters days to 50% flowering, plant height and panicle length PCV (9.44, 8.28 and 5.36) and GCV (9.31, 7.82 and 4.73) were low. Low PCV and GCV for days to 50% flowering was reported by Satish Chandra *et al.* (2009), Kuchanur *et al.* (2009), Saidaiah *et al.* (2010), Siva Parvathi *et al.* (2011) and Shiva Prasad *et al.* (2011), whereas for panicle length low PCV and GCV were obtained by Rita Binse *et al.* (2006), Mamta Singh *et al.* (2007), Karthikeyan *et al.* (2007), Krishna *et al.* (2008) and Prasad *et al.* (2009). The character number of filled grains panicle⁻¹ recorded high PCV (20.71) and moderate GCV (19.86). Kuchanur *et al.* (2009) reported similar results for this trait.

High estimates of heritability coupled with high genetic advance as per cent of mean was observed for number of filled grains panicle⁻¹ (92.00 and 39.22), spikelet fertility (90.00 and 20.35), test weight (93.00 and 24.79) and grain yield plant⁻¹ (82.00 and 45.63). High heritability coupled with high genetic advance as per cent of mean suggested the role of additive gene action in the inheritance of these traits and directional selection could be

Table 1. Analysis of variance for yield and yield contributing characters in rice (*Oryza sativa* L.) during *rabi*, 2010-11

Source of variations	df	Days to 50% flowering	Plant height (cm)	No. of ear bearing tillers plant ⁻¹	Panicle length (cm)	No. of filled grains panicle ⁻¹	Spikelet fertility (%)	Test weight (g)	Grain yield plant ⁻¹ (g)
MSS									
Replications	1	1.78	0.38	0.11	0.03	32.30	7.93	0.16	9.71
Genotypes	71	152.32**	181.59**	3.35**	3.84**	6358.54**	151.56**	10.91**	162.58**
Error	71	2.17	10.29	0.73	0.47	267.18	8.07	0.39	15.72

** Significant at 1 % level df = Degrees of freedom

Table 2. Estimates of genetic parameters for yield and yield component characters during *rabi*, 2010-11

S.No.	Character	Mean	Range		PCV (%)	GCV (%)	Heritability (%)	Genetic advance as % of mean (5 % GAM)
			Min	Max				
1.	Days to 50% flowering	93.10	77.50	112.50	9.44	9.31	97.00	18.90
2.	Plant height (cm)	118.34	98.10	141.00	8.28	7.82	89.00	15.22
3.	No. of ear bearing tillers per plant	9.74	7.30	14.50	14.67	11.78	64.00	19.47
4.	Panicle length (cm)	27.40	24.58	32.27	5.36	4.73	78.00	8.61
5.	No. of filled grains panicle ⁻¹	277.93	165.55	403.60	20.71	19.86	92.00	39.22
6.	Spikelet fertility (%)	81.30	42.28	91.55	10.99	10.42	90.00	20.35
7.	Test weight (g)	18.38	13.31	24.06	12.94	12.48	93.00	24.79
8.	Grain yield plant ⁻¹ (g)	35.11	11.40	54.60	26.89	24.41	82.00	45.63

PCV = Phenotypic coefficient of variation

GCV = Genotypic coefficient of variation

profitably applied on these traits in the genetically diverse material. Similar findings for grain yield plant⁻¹ and number of filled grains panicle⁻¹ were reported earlier by Karthikeyan *et al.* (2007), Prasad *et al.* (2009), Umadevi *et al.* (2010) and Siva Parvathi *et al.* (2011). For test weight, high heritability coupled with high genetic advance as per cent of mean was recorded by Vijayalakshmi *et al.* (2008), Satish Chandra *et al.* (2009), Saidaiah *et al.* (2010), Mohan Lal and Chauhan (2011) and Shiva Prasad *et al.* (2011).

High estimates of heritability coupled with moderate genetic advance as per cent of mean was observed for rest of the traits *viz.*, days to 50% flowering (97.00 and 18.90), plant height (89.00 and 15.22) and number of ear bearing tillers plant⁻¹ (64.00 and 19.47) revealing the involvement of both additive and non additive gene action in the inheritance of these traits and improvement of these traits may be possible by simultaneous exploitation of both additive and non additive components by adopting breeding procedures like biparental mating, diallel selective mating system or cyclic hybridization. High heritability (78.00) with low genetic advance as per cent of mean (8.61) was recorded for panicle length. The results were in agreement with the earlier findings of Mamta Singh *et al.* (2007), Prasad *et al.* (2009) and Shiva Prasad *et al.* (2011) indicating the role of non-additive gene action, which may be exploited through heterosis breeding. Therefore, from variability, heritability and genetic advance studies the traits *viz.*, grain yield plant⁻¹ number of filled grains panicle⁻¹ test weight and spikelet fertility may be considered as favourable attributes as they recorded high magnitude of variability parameters and high extended genetic gain obtained from selection of superior genotypes.

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