

## Studies on Genetic Parameters for Yield and its Contributing Characters in Rice Hybrids and Their Parents

Key words : Gentic Advance, Heritability and Variability.

Rice is one of the significant cereal commodities fulfilling the nutritional requirements of more than 70% of population and source of livelihood for about 150 million rural households in India. For the formulation of successful breeding programme, genetic variability in a crop species is pre-requite and it also offers better scope for selection. Effective selection not only depends on estimation of genetic variation among the genotypes but also on the proportion of heritable variation and the expected genetic gain. Heritability coupled with genetic advance would be more useful tool in predicting the resultant effects in selection of best genotypes for yield and its contributing traits. Therefore, present study was conducted to estimate the extant of genetic variability, heritability and genetic advance as per cent of mean of yield and yield component characters in rice.

The experimental material comprised of 72 genotypes including 52 hybrids, 17 parents (4 CMS B lines and 13 restorers) and three checks (viz., MTUHR 2089, MTU 1075 and MTU 1010) were transplanted in a randomized block design in two replications with spacing of 20x15 cm during rabi 2010-11 at Regional Agricultural Research Station, Warangal. Ten competitive plants from each genotype in each replication were selected at random to record data on characters viz., plant height, number of ear bearing tillers palnt<sup>-1</sup>, panicle length, number of filled grains panicle<sup>-1</sup>, spikelet fertility per cent and grain yield plant<sup>-1</sup>, whereas data on days to 50% flowering and test weight were recorded on plot basis. The mean data was subjected to standard method of analysis of variance (Panse and Sukhatme, 1978). 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.

In the present study, analysis of variance of 72 genotypes revealed the existence of highly significant differences among the genotypes for all the characters studied (Table 1). The estimates of phenotypic coefficient of variance (PCV) were slightly higher than those of genotypic coefficient of variance (GCV) for all the characters studied indicated minimum environmental influence and greater role of genetic factors on the expression of (Table 2). High PCV and GCV values traits were exhibited by grain yield plant-1(32.03 and 29.70) and number of filled grains panicle<sup>-1</sup> (24.53) and 23.35). Similar findings were earlier reported by Prasad et al. (2009), Mohan Lal and Chauhan (2011), Siva Parvathi et al. (2011) and Shiva Prasad et al. (2011). PCV and GCV were moderate for number of ear bearing tillers palnt<sup>-1</sup> (15.73 and 13.42), spikelet fertility (10.81 and 10.03) and test weight (12.65 and 12.23). Similar findings were given by Saidaiah et al. (2010) for spikelet fertility and Satish (2000) for test weight. Low PCV and GCV values were recorded for the traits viz., days to 50% flowering (8.54 and 8.43), plant height (10.27 and 9.66) and panicle length (6.33 and 5.70). Satish Chandra et al. (2009), Kuchanur et al. (2009), Saidaiah et al. (2010), Siva Parvathi et al. (2011) and Shiva Prasad et al. (2011) were also reported low PCV and GCV for days to 50% flowering whereas, for panicle length similar results were obtained by Mamta Singh et al. (2007) and Prasad et al. (2009).

High heritability was exhibited by all the yield and yield component traits. Since high heritability does not always indicate high genetic gain, heritability with genetic advance should be used in predicting selection of superior genotypes. High estimates of heritability coupled with high genetic advance as per cent of mean was observed for number of ear bearing tillers palnt<sup>-1</sup> (72.80 and

| Source Of<br>Variations | D.F. | Days To<br>50%<br>Flowering | Plant<br>Height<br>(Cm) | No. Of ear<br>bearomg<br>tollers plant <sup>-1</sup> | Panicle<br>Length<br>(cm) | No. of filled<br>grains<br>panicle <sup>-1</sup> | Spikelet<br>fertility<br>(%) | Test<br>weight<br>(g) | Grain yield<br>plant <sup>-1</sup><br>(g) |  |  |
|-------------------------|------|-----------------------------|-------------------------|--|---------------------------|--|------------------------------|-----------------------|---|--|--|
| MSS                     |      |                             |                         |  |                           |  |                              |                       |   |  |  |
| Replications            | 1    | 0.17                        | 0.42                    | 0.05   | 0.15                      | 50.65  | 4.06                         | 0.24                  | 2.18                                      |  |  |
| Genotypes               | 71   | 157.83**                    | 232.88**                | 4.36**   | 5.44**                    | 8558.47**  | 139.85**                     | 9.85**                | 220.74**                                  |  |  |
| Error                   | 71   | 2.10                        | 14.24                   | 0.69   | 0.57                      | 421.50   | 10.45                        | 0.33                  | 16.68                                     |  |  |

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

\* Significant At 5% Level

\*\* Significant At 1% Level

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

| S. No. Character |  | Mean   | Range   |         |       |            | Heritability | Genetic<br>Advance As           |
|------------------|--|--------|---------|---------|-------|------------|--------------|---------------------------------|
|                  |  |        | Minimum | Maximum |       | GCV<br>(%) | (%)          | Per Cent<br>Of Mean<br>(5% GAM) |
| 1. I             | Days to 50% flowering                      | 104.72 | 87.50   | 124.50  | 8.54  | 8.43       | 97.40        | 17.13                           |
| 2. I             | Plant height (cm)                          | 108.24 | 85.50   | 130.00  | 10.27 | 9.66       | 88.50        | 18.72                           |
| 3. ľ             | No. of ear bearing tillers per plant       | 10.10  | 7.40    | 15.10   | 15.73 | 13.42      | 72.80        | 23.59                           |
| 4. I             | Panicle length (cm)                        | 27.37  | 24.32   | 32.02   | 6.33  | 5.70       | 81.20        | 10.59                           |
| 5. ľ             | No. of filled grains panicle <sup>-1</sup> | 273.18 | 150.00  | 409.80  | 24.53 | 23.35      | 90.60        | 45.79                           |
| 6. 5             | Spikelet fertility (%)                     | 80.16  | 39.87   | 89.90   | 10.81 | 10.03      | 86.10        | 19.18                           |
| 7. ]             | Test weight (g)                            | 17.84  | 12.85   | 23.16   | 12.65 | 12.23      | 93.50        | 24.36                           |
| 8. (             | Grain yield plant <sup>-1</sup> (g)        | 34.02  | 12.00   | 59.40   | 32.03 | 29.07      | 86.00        | 56.71                           |

PCV = Phenotypic Coefficient Of Variation

GCV = Genotypic Coefficient Of Variation

23.59), number of filled grains panicle<sup>-1</sup> (90.60 and 45.79), test weight (93.50 and 24.36) and grain yield plant<sup>-1</sup> (86.00 and 56.71). These results indicate the preponderance of additive gene action and hence direct phenotypic selection may be useful with respect to these traits. Similar findings for grain yield plant<sup>-1</sup> and number of filled grains panicle<sup>-1</sup> were reported by Umadevi *et al.* (2010) and Siva Parvathi *et al.* (2011) and for test weight, Saidaiah *et al.* (2010), Mohan Lal and Chauhan (2011) and Shiva Prasad *et al.* (2011) also observed similar findings.

High heritability with moderate genetic advance as per cent of mean was recorded for the

traits *viz.*, days to 50% flowering (97.40 and 17.13), plant height (88.50 and 18.72), panicle length (81.20 and 10.59) and spikelet fertility (86.10 and 19.18). These results were in accordance with previous reports of Mamta Singh *et al.*, 2007, Saidaih *et al.*, 2010 and Siva Parvathi *et al.*, 2011. Thus, the characters *viz.*, number of ear bearing tillers palnt<sup>-1</sup>, number of filled grains panicle<sup>-1</sup>, test weight and grain yield plant<sup>-1</sup> which recorded moderate to high PCV, GCV heritability and genetic advance as per cent of mean could be transmitted to the progeny with hybridization and phenotypic based selection.

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