

GENOTYPE SEEDLING AGE INTERACTION FOR HYBRID SEED YIELD OF RICE (*Oryza sativa* L.)

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ABSTRACT

The experiment was carried out in Randomized Completely Block Design (RCBD) during the Boro season 2006-2007 (Mid October to mid May) at Bangabandhu Sheikh Mujibur Rahman Agricultural University-Gazipur. Alok showed significant b_i but its deviation from regression (S^2_{di}) invalidated its linear prediction. Among the three CMS lines CMS-2 was stable for % OCR due to considerable mean, phenotypic index, regression coefficient near unity ($b_i \approx 1$) as well as insignificant deviation from regression ($S^2_{di} \approx 0$) and CMS-1 was highly responsive to seedling ages due to significant b_i and insignificant S^2_{di} . Although, Alok showed significant b_i but its deviation from regression (S^2_{di}) invalidated its linear prediction. Maximum out crossing rate was obtained from CMS-2 irrespective of seedling ages. On the other hand 43 days old seedling of CMS-1 gave the maximum out crossing rate. CMS-2 was highly responsive to yield over seedling ages. CMS-1 showed insignificant regression coefficient (b_i) indicating stability over seedling ages but deviation from regression (S^2_{di}) was significant which invalidated its linear prediction. Alok showed lower mean, significant regression coefficient (b_i) and insignificant deviation from regression (S^2_{di}) indicating responsive to favourable seedling ages with comparatively lower yield. 43 days old seedlings of CMS-2 gave the maximum hybrid seed production rate.

Key words: Rice (*Oryza sativa*), GxE interaction, CMS, hybrid, seed yield

INTRODUCTION

Yield stability under a wide range of environmental condition is very important for adaptation of varieties. The yield of a crop plant is a quantitative (polygenic) character and generally influenced by environmental variation. An ideal variety is one that has high mean yield but a low degree of fluctuation in performance when grown over diverse environment. Hence, testing for genotype x environment interaction has become an important task in the most of the breeding program (Prasad and Singh, 1990). Importance of genotype environment interaction is well established in the field of plant breeding. A successful evaluation of stable genotypes which could be used in future breeding program to develop stable genotypes with a high mean yield but a low degree of fluctuation in performance. It can be done through the study of genotype environment interactions. Stability for widely and specific environments have been assessed by different scientists but information on stability analysis of rice under different

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environment is little bit of scares. Therefore it is necessary to identify stable genotypes with high mean yield under wide range of environment. Thus genotype x environment interactions may play an important role and help to select widely adaptable as well as location specific varieties more efficiently. Genotype x environment interaction implies different behavior of genotypes under different environmental condition (Sarker, 2002).

MATERIALS AND METHODS

Three promising CMS lines were studied in micro-environments (three different seedling ages: 25days, 34 days and 43 days) for stability analysis. The experimental field was typically rice growing low land of 23⁰08'-24⁰09' N latitude and 90⁰26'- 90⁰82' longitudes with an elevation of 8.4-8.8 m from the mean sea level. The soil of the experimental field was Flood plain of Jamuna-Karatoya soil type under Barind Tract in (AEZ 8, 25 & 28). The experimental site is characterized by heavy rainfall during June to September and minimum rainfall during rest of the year. The temperature ranged during the experimental period 7.9⁰ C-31.8⁰ C. The maximum and minimum humidity ranged from 61%-96%. The average humidity was around 86%. Three CMS lines, Alok, CMS-01 and CMS-02 were used in the study as well as three different seedling ages i.e., seedling age of 25 days, 34 days and 34 days. The phenotypic index (Pi) is used for easy interpretation (Ram *et al.*, 1970). Stability parameters (bi and S²di) were estimated following Eberhart and Russell's model (1966).

RESULTS AND DISCUSSION

Combined analysis of variance due to environments (seedling age of 25 days, 34 days and 43 days), genotypes and genotype x seedling age interaction for grain yield and its important component of seed production (out crossing rate) are given in table 1. The mean sum of squares due to variety was significant for % OCR and grain yield that revealed the difference among genotypes and their considerable influence in out crossing rate and yield of rice. The environmental (seedling ages) mean sum of square was significant for out crossing rate and yield which revealed the variation of out crossing rate and yield of rice varieties over seedling ages. Sarker *et al.* (2002) found similar results in their study on stability for grain yield under different planting times in Bangladesh. The significant mean sum of square of genotype x environment interaction for both the traits indicated the presence of differences among the regression of the genotypes on environmental indices. This result suggests that the analysis could be extended for the stability analysis. The mean sum of squares of interaction between environment + (genotype x environment) was also significant for out crossing rate and yield which indicated very minute deviation from regression.

Table 1. Pooled analysis of variance for out crossing rate (% OCR) and grain yield of three CMS lines

Source of variation	df	Mean sum of squares	
		% OCR	Yield
Genotype	2	1.41*	0.16*
Environment (Seedling ages)	2	16.58*	0.15*
Genotype x Environment	4	3.67**	0.05*
Environment + (genotype x Environment)	6	28.83*	0.18*
Environment (linear)	1	31.89**	0.81**
Genotype x Environment (linear)	2	2.35ns	0.89**
Pooled deviation	3	32.64	0.08
Pooled error	12	1.65	0.02

* and ** significant at 5 % and 1 % level of significance, and ns = insignificant

The linear portion of environment mean sum of square was highly significant for out crossing rate and yield (Mannan, 1991). The linear portion of genotype x environment mean sum of square was insignificant for out crossing rate but it was highly significant for yield. Significant linear components indicating the presence of differences among the regression values of the genotypes on the environmental indices. Insignificant non linear components (pooled deviation) for the traits suggested the stability of the genotypes with the changes of seedling age. It also explains that the production based on regression analysis for the genotypes differed considerably in respect to their stability performance. Das *et al.* (1991) also found significant linear and non-linear component of the interaction in %OCR and yield of rice. According to Eberhart and Russell (1966), a stable genotype is characterized by a slope not differ from unity ($b_i = 1$) and the deviation from regression close to zero ($S^2di = 0$). The stability parameters mean phenotypic index (Pi), regression coefficient (b_i) and deviation from regression (S^2di) for yield and out crossing rate are presented in table 2 and 3 (Manonmani *et al.*, 1991).

Table 2. Environment wise mean out crossing rate (% OCR) and estimated stability of three CMS lines

Genotype	Environments (seedling age)			Mean	Phenotypic index (PI)	Regression coefficient (b_i)	deviation from regression (S^2di)
	25 days old	34 days old	43 days old				
Alok	39.99	42.7	42.60	41.77	-3.15	2.58*	5.26*
CMS-1	39.04	42.93	50.32	44.09	-0.83	3.91*	0.72 ^{ns}
CMS-2	42.12	49.83	54.75	48.90	3.98	1.01 ^{ns}	0.16 ^{ns}
Env mean	40.38	45.15	49.23	(GM)			
Env index	-4.54	0.23	4.31	44.92			
SE	0.24						

* Significant at 5 % and ns = insignificant

The linear portions of G x E interaction was significant and non linear portions was insignificant for both out crossing rate and grain yield (t/ha) but in case of individual test, Alok exhibited significant deviation from regression (S^2di) for % OCR and CMS-1 exhibited significant deviation from regression (S^2di) for yield, thus stability appears to be the property of individual buffering rather than population buffering for these characters. From the environmental index it was clear that 43 days old seedlings were suitable for favorable environments for % OCR (Rashid *et al.*, 1991). Among the three CMS lines CMS-2 was stable for % OCR due to considerable mean, phenotypic index, regression coefficient near unity ($b_i \approx 1$) as well as insignificant deviation from regression ($S^2di \approx 0$) and CMS-1 was highly responsive to seedling ages due to significant b_i and insignificant S^2di . Although, Alok showed significant b_i but its deviation from regression (S^2di) invalidated its linear prediction (table 2). Maximum out crossing rate was obtained from CMS-2 irrespective of seedling ages (Rao *et al.*, 1991). On the other hand 43 days old seedling of CMS-1 gave the maximum out crossing rate. CMS-2 was highly responsive to yield over seedling ages as judge through high mean, phenotypic index, significant regression coefficient (b_i) and insignificant (S^2di). CMS-1 showed insignificant regression coefficient (b_i) indicating stability over seedling ages but deviation from regression (S^2di) was significant which invalidated its linear prediction (Table 3).

Table 3. Environment wise mean yield and estimated stability of three CMS lines

Genotype	Environments (seedling age)			Mean	Phenotypic index (PI)	Regression coefficient (bi)	deviation from regression (S ² di)
	25 days old	34 days old	43 days old				
Alok	3.10	3.08	3.36	3.18	-0.37	0.83*	-1.25 ^{ns}
CMS-1	3.32	3.42	3.62	3.44	-0.11	0.36 ^{ns}	1.11*
CMS-2	4.04	3.84	4.21	4.03	0.48	0.93*	-0.21 ^{ns}
Env mean	3.48	3.44	3.73	(GM)			
Env index	-0.07	-0.11	0.18	3.55			
SE	0.03						

* Significant at 5 % and ns = insignificant

Alok showed lower mean, significant regression coefficient (bi) and insignificant deviation from regression (S²di) indicating responsive to favourable seedling ages with comparatively lower yield. 43 days old seedlings of CMS-2 gave the maximum hybrid seed production rate.

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