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ESTIMATION OF HERITABILITY THROUGH PARENT-OFFSPRING ANALYSIS IN SORGHUM (SORGHUM BICOLOR (L) MOENCH) CROSSES

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A study of parent progeny regression analysis and heritability in narrow sense between F₂ and F₃ population of three inter varietal crosses for six quantitative traits indicated that the correlation and regression co-efficients between F₂ mean and corresponding parental F₂ values were significant for plant height, ear length end 100 grain weight in one cross, number of leaves and ear weight in another cross and plant height and number of leaves in a third cross. This indicates that the transmission of genetic potentialities from F₂ to F₂ is more and is attributed to additive gene effects. Hence, the scheme of selection in the segregating generations will be effective for these characters of respective crosses. The correlation and regression co-efficients between F₃ mean and corresponding parental F₂ values were not significant for grain yield in all the three crosses.

Parent - offspring regression analysis enables to estimate heritability in narrow sense, resulting from additive gene effects only (Lush, 1940). The heritability estimate through parent progeny regression method is more reliable and Frey and Horner (1955) were of the opinion that this method may tend to reflect the true heritability. An evaluation of the transmission of genetic potentialities from F₂ F₃ of three inter-varietal crosses through parent-offspring regressin analysis is presented.

MATERIALS AND METHODS

The F2 s of three sorghum crosses namely, 2219 A X SPV 351 (Cross A), 2219 A x Usilampatti—2 (Cross B) and 2760 A x Co 24 (Cross C) with their parents viz, 2219 B (P1), 2760 B (p2), CO 24 (p3),

Usilampatti - 2 (local) (p4) and SPV 351 (p5) were raised during August 1982. In each cross, thirty out of 600 segregants were randomly selected and advanced to F3 generation. Each selected F2 plant from each cross was raised in one row with parents during February 83. The observations were made on all plants in F3 except the sterile segregants on plant height, number of leaves, ear length, ear weight, 100 grain weight and grain yield. The regression co-efficient 'b' was calculated following Kempthorne, 1957.

'b' =
$$\frac{\text{Sum of products of F2 and F3}}{\text{Sum of square of F3}}$$
Heritability in narrow sense between F2 and F3 = $\frac{2}{3}$ Xb

The significance of 'b' co-efficient was tested by 't' test.

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RESULTS AND DISCUSSION

The inter generation correlation and regression between selected F2 plants on their respective F3 mean values are presented in Table 1. Heritability in narrow sense was estimated through parent-progeny regression of F2 on F3 and presented in Table 2.

Table 1. Inter-Generation Correlation (r) and Regression Coefficients (b) Between F, and F, Means

Cross A		Cross	B Cross C		
770	b				
		-		***	1. 21 1.
0.25**	0.64	0.72**	1.72** -	0.52**	1.18**
0.52**	0,96**	-0,23	-0.65	0.44**	0.95**
0.21	0.66	0.61**	2.02**	-0.001	-0.02
0.61**	2,59**	-0.14	-0.87	0.14	0.59
0.13	0.25	1.00**	1.50**	0.11	0.25
0.08	0.39	0.06	0.41	0,04	C.17
	0.25** 0.52** 0.21 0.61** 0.13	0.25** 0.64 0.52** 0.96** 0.21 0.66 0.61** 2.59** 0.13 0.25	0.25** 0.64 0.72** 0.52** 0.96** -0.23 0.21 0.66 0.61** 0.61** 2.59** -0.14 0.13 0.25 1.00**	0.25** 0.64 0.72** 1.72** 0.52** 0.96** -0.23 -0.65 0.21 0.66 0.61** 2.02** 0.61** 2.59** -0.14 -0.87 0.13 0.25 1.00** 1.50**	0.25** 0.64 0.72** 1.72** 0.52** 0.52** 0.96** -0.23 -0.65 0.44** 0.21 0.66 0.61** 2.02** -0.006 0.61** 2.59** -0.14 -0.87 0.14 0.13 0.25 1.00** 1.50** 0.11

^{*} Significant at 5 per cent level

Table 2. Estimates of Heritability in Narrow Sense in F_a Generation of Sorghum Crosses

· Cross	es He	Heritability (2/3 b)		
Characters	A	В	C	
Plant height (cm)	_	1.15	0.79	
Number of leaves	0.64	**	0.54=	
Ear length (cm)	0,44	1.35		
Ear weight (g)	1.74	*#: "	0.46	
weight (g)	0.36	1.01	C.17	
Grain yield (g)	0.26	0.27	0.11	

Parent-progeny regres ion was not significant and hence heritability was not calculated

The correlation and regression co - efficient estimates between F3 means and corresponding parental F2

values were significant for number of leaves and ear weight in cross A. plant height, ear length and 100 grain weight in cross B and plant height, and number of leaves in cross C (Table 1). This indicated that there was some relationship between F2 value and its progeny mean. The grain yield was not significant in all the three crosses indicating that there was no relationship between F2 values and its progeny mean: McGinnis and Shebaski (1968) also obtained non - significant correlation between F2 and F3 and reported that on this basis alone there was no advantage in selecting F2 plants for high yield. The absence of relationship between F2 and F3 perhaps indicated that the

^{**} Significant at 1 per cent level

interference of non-additive components were very considerable, and hence, the scheme of selection should be postponed to later generations.

In the present investigation, high heritability in narrow sense obtained in cross B followed by cross C for plant height, cross A and cross C for number of leaves. cross B followed by cross A for ear. length, cross A followed by cross C for ear weight, cross B followed by cross A for 100 grain weight and grain yield (Table 2). The environment contribution to the total variability in these characters was quite low and consequently the transmission potential of the parental material for these characters was appreciably high. Panchal et al (1979) noticed high heritability in narrow sense for plant height, ear length and ear weight

For grain yield, low heritability was noticed in all the three crosses

which indicated that the environment effect on this character is high.

The negative values of parentoff spring regression in some instances showed that the F2 performance was not a good indicator of F3 progeny performance for these traits. There are several possible reasons given by Meredith and Bridge (1973) to explain why the F2 performance may not be closely related to that of the F3 progeny mean. The environment of an individual plant may differ greately from that of progeny rows A second factor may be that the genotype X environment interactions from one year to the next at a single location may differ. Dominance gene action in an F2 population i, e large non-additive gene effects may not be usable in later generations.

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