



## Genetic analysis of kernel quality traits in rice

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**Abstract:** In rice, genetics of kernel quality traits viz. kernel length, kernel L/B ratio, kernel length after cooking, kernel elongation, and elongation index was studied in five crosses. All the quality traits were governed by additive, dominance and epistatic interactions of additive x additive, additive x dominance and dominance x dominance and duplicate type. Pedigree method of breeding followed by selection in later generations involving one or two cycles of intermating of selected segregant is suggested for improvement.

**Key words :** *Basmati rice, Kernel quality, Gene action.*

### Introduction

Studies on the nature of gene action governing the yield and its component traits in rice are available in plenty while information on the genetics of kernel quality traits such as grain size, shape and elongation on cooking are limited. Hence, an attempt was made to unravel the genetic architecture of some of the kernel quality involving basmati and non-basmati rice varieties.

### Materials and Methods

Five high yielding cosmopolitan rice varieties viz. ASD 19, ADT 38, ADT 39, CO 43 and improved White Ponni were crossed with Pusa Basmati 1 during summer 1996 at the Plant Breeding Farm, Department of Agricultural Botany, Faculty of Agriculture, Annamalai University, Annamalaiagar. Five hybrids along with the parents were raised during the Kuruvai 1996 (May-Sept). The F<sub>1</sub>s were used as female and pollen from their corresponding parents were dusted separately to obtain the two back cross progenies viz. B<sub>1</sub> (F<sub>1</sub>/P<sub>1</sub>) and B<sub>2</sub> (F<sub>1</sub>/P<sub>2</sub>). Fresh crosses were also effected to obtain F<sub>1</sub> seeds. Selfing of parents and all F<sub>1</sub>s was also done to obtain seeds of P<sub>1</sub>, P<sub>2</sub> and F<sub>2</sub> generations.

All the six generations viz. P<sub>1</sub>, P<sub>2</sub>, F<sub>1</sub>, F<sub>2</sub>, B<sub>1</sub> and B<sub>2</sub> of each of the five crosses were raised in randomised block design with three replications during late Thaladi 1997 (Oct-Feb). The plot size for each cross was 6 x

3 m with a spacing of 30 x 20 cm. Ten plants in each of P<sub>1</sub>, P<sub>2</sub> and F<sub>1</sub>; 40 plants in each of B<sub>1</sub> and B<sub>2</sub> and 70 plants in F<sub>2</sub> generation per replication were randomly selected to record observations on the following traits viz. kernel length, kernel L/B ratio, kernel length after cooking, kernel elongation and elongation index.

Seeds hulled in McGill sample sheller were used to measure kernel length in mm with Mitutoyo micrometer. The ratio of kernel length to kernel breadth was computed as kernel L/B ratio. For cooking test, four months old samples were used for analysis. The hulled seeds were milled in KETT rice polisher uniformly for 40 seconds. The milled rice was cooked as per the method of Juliano and Perez (1984). Ten randomly selected whole milled kernels before and after cooking were measured in mm with a graduated card board (Piilaiyar and Mohandoss, 1981). The ratio of the mean length of cooked rice to mean length of milled rice was computed as kernel elongation. The elongation index was derived by dividing the mean L/B ratio of cooked rice with the mean L/B ratio of milled rice (Juliano and Perez, 1984).

The adequacy of additive-dominance model was studied by scaling test. In cases where the scales A or B or C or D significantly differed from zero, a digenic interaction model was assumed. Genetic components of generation

Table 1. Generation means and estimates of gene effects

	ASD 19/Pusa Basmati 1	ADT 38/Pusa Basmati 1	ADT 39/Pusa Basmati 1	CO 43/Pusa Basmati 1	Improved White Ponni/Pusa Basmati 1
<i>Kernel length</i>					
Generation					
	5.32 ± 0.025	6.29 ± 0.008	5.50 ± 0.012	5.47 ± 0.013	5.72 ± 0.016
	45 ± 0.019	7.45 ± 0.020	7.45 ± 0.020	7.45 ± 0.020	7.45 ± 0.020
	98 ± 0.043	6.27 ± 0.013	6.21 ± 0.013	6.17 ± 0.015	5.99 ± 0.017
	47 ± 0.006	6.34 ± 0.004	6.22 ± 0.005	6.11 ± 0.005	5.82 ± 0.005
	112 ± 0.009	5.80 ± 0.006	5.66 ± 0.007	5.73 ± 0.034	6.37 ± 0.009
	6.12 ± 0.008	5.84 ± 0.008	6.50 ± 0.07	6.04 ± 0.034	6.75 ± 0.009
Parameter					
	5.79* ± 0.04	8.96* ± 0.03	7.04* ± 0.03	7.38* ± 0.10	3.62* ± 0.03
	-1.07* ± 0.02	-0.58* ± 0.01	-0.98* ± 0.01	-0.99* ± 0.01	-0.87* ± 0.01
	-1.46 ± 0.11	-7.78* ± 0.08	-2.44* ± 0.08	-3.85* ± 0.29	6.42* ± 0.10
	0.60* ± 0.03	-2.09* ± 0.03	-0.56* ± 0.03	-0.92* ± 0.10	2.96* ± 0.03
	0.07* ± 0.02	0.54* ± 0.01	0.14* ± 0.02	0.68* ± 0.05	0.49* ± 0.02
	1.65* ± 0.11	5.09* ± 0.06	1.61* ± 0.05	2.65* ± 0.20	-4.05* ± 0.07
<i>Kernel L/B ratio</i>					
Generation					
	2.44 ± 0.014	3.33 ± 0.015	2.49 ± 0.017	2.46 ± 0.023	2.95 ± 0.019
	4.55 ± 0.035	4.55 ± 0.035	4.55 ± 0.035	4.55 ± 0.035	4.55 ± 0.035
	3.18 ± 0.023	3.41 ± 0.020	3.15 ± 0.018	3.08 ± 0.020	3.12 ± 0.030
	3.04 ± 0.010	3.43 ± 0.008	3.22 ± 0.009	2.98 ± 0.006	3.06 ± 0.009
	2.53 ± 0.008	3.01 ± 0.008	2.83 ± 0.009	2.63 ± 0.007	3.35 ± 0.015
	3.46 ± 0.014	3.11 ± 0.009	3.46 ± 0.011	3.08 ± 0.010	3.55 ± 0.012
Parameter					
	3.69* ± 0.05	5.43* ± 0.04	3.83* ± 0.05	4.02* ± 0.04	2.20* ± 0.06
	-1.06* ± 0.02	-0.61* ± 0.02	-1.03* ± 0.02	-1.04* ± 0.02	-0.80* ± 0.02
	-2.07* ± 0.14	-5.99* ± 0.11	-1.74* ± 0.13	-3.20* ± 0.11	2.52* ± 0.15
	-0.19* ± 0.05	-1.50* ± 0.04	-0.31* ± 0.04	-0.51* ± 0.04	1.55* ± 0.05
	0.12* ± 0.03	0.51* ± 0.02	0.40* ± 0.02	0.59* ± 0.02	0.59* ± 0.03
	1.57* ± 0.10	3.96* ± 0.08	1.07* ± 0.09	2.26* ± 0.08	-1.60* ± 0.11
<i>Kernel length after cooking</i>					
Generation					
	7.97 ± 0.035	8.40 ± 0.011	8.05 ± 0.017	8.18 ± 0.019	8.15 ± 0.023
	12.92 ± 0.034	12.92 ± 0.034	12.92 ± 0.034	12.92 ± 0.034	12.92 ± 0.034
	9.99 ± 0.072	10.84 ± 0.022	9.44 ± 0.308	9.63 ± 0.023	10.43 ± 0.032
	9.17 ± 0.009	10.89 ± 0.008	9.75 ± 0.070	10.75 ± 0.009	9.80 ± 0.009
	8.15 ± 0.014	9.80 ± 0.010	8.75 ± 0.068	8.95 ± 0.011	10.49 ± 0.015
	10.03 ± 0.013	9.52 ± 0.012	10.23 ± 0.011	10.66 ± 0.013	10.90 ± 0.014
Parameter					
	10.77* ± 0.06	17.02* ± 0.05	11.52* ± 0.14	14.34* ± 0.05	6.96* ± 0.06
	-2.48* ± 0.02	-2.26* ± 0.02	-2.44* ± 0.02	-2.37* ± 0.02	-2.39* ± 0.02
	-5.61* ± 0.17	-18.34* ± 0.13	-4.98* ± 0.52	-9.64* ± 0.14	7.89* ± 0.16
	-0.32* ± 0.05	-6.36* ± 0.04	-1.03* ± 0.14	-3.78* ± 0.05	3.58* ± 0.05
	0.60* ± 0.03	1.82* ± 0.03	0.96* ± 0.07	0.66* ± 0.03	1.98* ± 0.03
	4.84* ± 0.17	12.16* ± 0.09	2.90* ± 0.67	4.93* ± 0.10	-4.12* ± 0.12

\*Significant at 5 per cent level.

Table 2. Generation means and estimates of gene effects

	ASD 19/Pusa Basmati 1	ADT 38/Pusa Basmati 1	ADT 39/Pusa Basmati 1	CO 43/Pusa Basmati 1	Improved White Ponni/Pusa Basmati 1
<i>Kernel elongation</i>					
Generation					
P <sub>1</sub>	1.61 ± 0.010	1.37 ± 0.002	1.47 ± 0.003	1.51 ± 0.004	1.53 ± 0.003
P <sub>2</sub>	2.10 ± 0.006	2.10 ± 0.006	2.10 ± 0.006	2.10 ± 0.006	2.10 ± 0.006
F <sub>1</sub>	1.81 ± 0.014	1.86 ± 0.004	1.75 ± 0.004	1.74 ± 0.004	1.90 ± 0.006
F <sub>2</sub>	1.80 ± 0.002	1.86 ± 0.001	1.78 ± 0.001	1.83 ± 0.002	1.82 ± 0.001
B <sub>1</sub>	1.68 ± 0.003	1.66 ± 0.002	1.66 ± 0.002	1.67 ± 0.002	1.72 ± 0.003
B <sub>2</sub>	1.75 ± 0.002	1.72 ± 0.002	1.76 ± 0.002	1.80 ± 0.002	1.78 ± 0.002
Parameter					
(m)	2.19* ± 0.01	2.42* ± 0.01	2.06* ± 0.01	2.19* ± 0.01	2.09* ± 0.01
(d)	-0.24* ± 0.01	-0.36* ± 0.003	-0.31* ± 0.003	-0.29* ± 0.004	-0.28* ± 0.003
(h)	-1.19* ± 0.03	-1.66* ± 0.02	-0.81* ± 0.02	-0.95* ± 0.02	-0.90* ± 0.02
(i)	-0.34* ± 0.01	-0.68* ± 0.01	-0.28* ± 0.01	-0.38* ± 0.01	-0.28* ± 0.01
(j)	0.17* ± 0.01	0.30* ± 0.004	0.21* ± 0.004	0.16* ± 0.005	0.22* ± 0.003
(l)	0.81* ± 0.03	1.11* ± 0.02	0.50* ± 0.02	0.47* ± 0.02	0.71* ± 0.02
<i>Elongation index</i>					
Generation					
P <sub>1</sub>	1.10 ± 0.006	0.87 ± 0.005	0.93 ± 0.006	1.15 ± 0.009	0.95 ± 0.003
P <sub>2</sub>	1.24 ± 0.010	1.24 ± 0.010	1.24 ± 0.010	1.24 ± 0.010	1.24 ± 0.010
F <sub>1</sub>	1.10 ± 0.012	1.13 ± 0.008	1.18 ± 0.009	1.19 ± 0.010	1.16 ± 0.010
F <sub>2</sub>	1.09 ± 0.009	1.14 ± 0.003	1.18 ± 0.003	1.33 ± 0.004	1.14 ± 0.003
B <sub>1</sub>	1.05 ± 0.004	1.06 ± 0.011	1.13 ± 0.006	1.21 ± 0.004	1.10 ± 0.002
B <sub>2</sub>	0.97 ± 0.004	1.03 ± 0.004	1.16 ± 0.005	1.14 ± 0.005	1.12 ± 0.002
Parameter					
(m)	1.49* ± 0.04	1.44* ± 0.03	1.21* ± 0.02	1.18* ± 0.02	1.21* ± 0.02
(d)	-1.07* ± 0.01	-0.19* ± 0.01	-0.15* ± 0.01	-0.04* ± 0.01	-0.14* ± 0.01
(h)	-1.20* ± 0.08	-0.89* ± 0.08	-0.09* ± 0.06	-0.30* ± 0.05	-0.24* ± 0.04
(i)	-1.31* ± 0.04	-0.39* ± 0.03	-0.12* ± 0.02	-0.61* ± 0.02	-0.12* ± 0.01
(j)	0.15* ± 0.01	0.21* ± 0.01	0.12* ± 0.01	0.12* ± 0.01	0.12* ± 0.01
(l)	0.82* ± 0.05	1.58* ± 0.05	0.06* ± 0.04	0.68* ± 0.04	0.19* ± 0.04

\* Significant at 5 per cent level.

mean were estimated following six parameter model (Mather and Jinks, 1982).

## Results and Discussion

Six varieties with diverse kernel characters were crossed to generate six generations involving five cross combinations. Means of each generation were calculated and scaling test was applied to test the adequacy of additive-dominance model. Based on scaling tests, a digenic epistatic model was assumed for all the traits studied. The mean value of kernel traits for six generations

and estimates of gene effects are given in Table 1 and 2. The observed value of F<sub>1</sub> for all the kernel characters was intermediate between the two parental means.

Additive (d) and dominance (h) effects were negative and significant for kernel length. Murai and Kinoshita (1986) and Kato (1989) opined that grain size was governed by additive-dominance model. The additive x additive interaction effect was positive and significant in two crosses indicating enhancing effect. The additive x dominance and dominance x dominance



Effects were also significant and positive. The 'I' effects recorded opposite signs for the kernel characters revealing the presence of duplicate interaction. Additive, dominance and epistatic interactions of all the three and duplicate type appear to govern kernel length.

The additive and dominance effects were positive and significant for kernel L/B ratio and kernel length after cooking. Singh and Richharia (1977), Somrith *et al.* (1979), and Singh and Singh (1985) reported that this was controlled by both additive and non-additive gene actions, while Reddy and Nerkar (1991) reported the absence of epistasis for this trait. Sarawgi *et al.* (1991) observed predominance of non-additive component for kernel L/B ratio. Additive x additive interaction effect was significant and negative revealing a diminishing effect. Both additive x dominance and dominance x dominance effects were positive and significant. Hence, additive, dominance and epistatic interactions of all the three and duplicate type appear to influence kernel L/B ratio and kernel length after cooking.

The additive and dominance effects were positive and significant for kernel elongation and elongation index. Siddiq (1980) reported that kernel elongation was predominantly governed by non-additive gene action. The additive x additive interaction effect was negative and significant while the additive x dominance and dominance x dominance effect was positive and significant. Hence, additive, dominance and epistatic gene actions of all the three and duplicate interaction control kernel elongation and elongation index. Tomar (1987) reported that additive and epistatic reaction mainly of duplicate effects influenced kernel elongation.

The grain quality traits viz. kernel length, kernel L/B ratio kernel length after cooking, kernel elongation and elongation index were under the control of additive, dominance and all the three kinds of epistatic interactions along with duplicate type.

As additivity was present for all the traits, pedigree method of breeding may be followed for improvement of these traits. Since considerable amount of dominance effect was also present,

selection has to be postponed to later generations until homozygosity is achieved. Since epistatic interactions of various kinds were also observed, one or two cycles of intermating of selected segregants will result in the improvement of kernel traits.

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