

Full Length Research Paper

Variability studies for quality traits in rice with high iron and zinc content in segregating population

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Received 1 February, 2014; Accepted 28 July, 2015

More than half of the world's population, especially women and children in the developing countries suffer from micronutrient malnutrition especially deficiency in iron and zinc. Micronutrient malnutrition problems increased the interest of researchers to increase the mineral contents (Fe and Zn) in cereals to ensure adequate attainment of dietary minerals. A lot of variability does exist for micronutrients (Fe, Zn, Vitamin A, etc.) content and bioavailability in many crops including rice. The current study was conducted to assess the variability for iron and zinc content in dehusked rice grains to identify mineral-rich families. This study was conducted with the major objectives of analysis of genetic variability for quality traits for grain iron and zinc content. Based on mean, genotypic coefficient of variability (GCV) and phenotypic coefficient of variability (PCV), heritability and genetic advance, it was understood that the progenies of ADT 37 × IR68144-3B-2-2-3 would be more useful for improving grain iron content with the desirable quality traits viz., kernel length, kernel breadth after cooking,. Similarly TRY (R) 2 × Mapillaisamba segregants could be used for improving the grain zinc content and breadth wise expansion ratio.

Key words: Genotypic coefficient of variability (GCV), phenotypic coefficient of variability (PCV), genetic advance (GA), iron and zinc content.

INTRODUCTION

Rice is the life and the prince among cereals as this unique grain helps to sustain two thirds of the world's population. Asia is the biggest rice producer, accounting for 90% of the world's production and consumption of rice (Anonymous, 2007). It is considered as the main staple food for more than 50% of the world's population. Rice provides 75% of the calories and 55% of the protein in the average daily diet of the people although it has an

incomplete amino acid profile and comprises confined amount of essential micronutrients (Bhuiyan et al., 2002). The per capita consumption of rice is very high ranging from 62 to 190 kg/year (Graham et al., 1999). To be healthy, human beings require more than 20 mineral elements and more than 40 nutrients, particularly vitamins and essential amino acids, all of which can be supplied by an appropriate diet (Philip and Martin, 2005).

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However, human diets often lack one or more of essential nutrients and cause micronutrient malnutrition. Micronutrient malnutrition is recognized as a massive and rapidly growing public health issue especially among poor people living on an unbalanced diet dominated by a single staple grain such as rice. Among the major micronutrient deficiencies common in rice consuming countries, iron and zinc deficiencies (the so called “hidden hunger”), affect over three billion people worldwide, mostly in developing countries (Welch and Graham, 2004). Plant breeding to enhance the nutrient quality of staple food crops holds promise for a low cost and sustainable approach to alleviate the problem of micronutrient malnutrition among the poorest segments of the population of developing countries. This approach has been called biofortification. Biofortification reduces malnutrition by breeding essential micronutrients into staple crops. This approach bridges the fields of human nutrition, crop science and public health to develop a set of highly sustainable nutrition interventions in a cost-effective manner. Exploiting the genetic variation in crop plants for micronutrient content is one of the most powerful tools to change the nutrient balance of a given diet on a large scale.

MATERIALS AND METHODS

Seeds of F₃ generation of four cross combinations generated from Anbil Dharmalingam, Agricultural College and Research Institute, Trichy. *viz.*, ADT 37 x IR68144-3B-2-2-3, TRY (R) 2 x Mapillaisamba were utilized as the experimental material in the present study. Among the parents *viz.*, TRY (R) 2 and ADT 37 are high yielding commercial varieties and IR68144-3B-2-2-3 is a iron donor and Mapillaisamba is a zinc donor which were used in earlier hybridization programme for introgression of high iron and zinc contributing genes. The experiment was conducted at Agricultural College and Research Institute, Madurai. The F₄ generation was raised during August to November, 2011 and F₅ generation during December 2011 to April 2012 respectively. The F₄ progenies were raised along with their parents in randomized block design with two replications. A total of five families were selected from each cross combination based on high iron and zinc content in F₃ population. For each family, 75 seedlings per replication were raised with a spacing of 20 cm between the rows and 15 cm between the plants. Each family had five rows of 15 single plants each. The recommended agronomic practices were followed throughout the crop growth period. Five single plants per family per replication were randomly selected and forwarded as single plant progeny row in F₅ generation. The mean data after computing for each character subjected to standard method of analysis of variance following Panse and Sukhatme (1961), phenotypic and genotypic coefficient of variation, heritability (Broad sense) and genetic advance as percent of mean were estimated by the formula as suggested by Burton (1952) and Johnson et al. (1955). The following quality traits are:

1. Kernel Length (KL): Ten unbroken hulled kernels with their tips intact from each genotype were measured for their length using vernier calipers. Average of length was taken in millimeters.
2. Kernel breadth (KB): Ten unbroken hulled kernels with their tips intact from each genotype were measured for their breadth using vernier callipers. Average of breadth was taken in millimetres.

3. Kernel length / breadth ratio (KLBR): The ratio between kernel length and breadth was estimated.

4. Kernel length after cooking (KLAC): Ten randomly selected whole milled rice kernels were taken in a labelled test tube and soaked in water for ten minutes. The tubes were placed in a water bath maintained at boiling temperature (100°C) for 20 min. After cooking, the cooked kernels (intact on both ends) were measured using a graph paper mounted in a glass frame and the average kernel length after cooking was expressed in millimetres.

5. Kernel breadth after cooking (KBAC): Ten randomly selected whole milled rice kernels were taken in a labelled test tube and soaked in water for ten minutes. The tubes were placed in a water bath maintained at boiling temperature (100°C) for 20 min. After cooking, the cooked kernels (intact on both ends) were measured using a graph paper mounted in a glass frame and the average kernel breadth after cooking was expressed in millimetres.

6. Linear elongation ratio (LER): The ratio of mean length of cooked rice to mean length of milled rice was computed as linear elongation ratio (Juliano and Perez, 1984):

$$\text{Linear elongation ratio} = \frac{\text{Kernel length after cooking}}{\text{Kernel length before cooking}}$$

7. Breadth wise expansion ratio (BER): The ratio of mean breadth of cooked rice to mean breadth of milled rice was computed as breadth wise expansion ratio:

$$\text{Breadth wise expansion ratio} = \frac{\text{Kernel breadth after cooking}}{\text{Kernel breadth before cooking}}$$

The zinc and iron content were determined by using atomic absorption spectrophotometer as suggested by Jackson (1973). All the statistical analysis was done by using GENRES statistical software GEN STAT 2004.

RESULTS AND DISCUSSION

In the present investigation ADT37 x IR68144-3B-2-2-3 showed high mean value for the characters *viz.*, kernel length, kernel breadth after cooking, iron content. TRY (R) 2 x Mapillaisamba had high mean value for the characters *viz.*, zinc content, breadth wise expansion ratio. The genotypic and phenotypic coefficient of variability were low in both the crosses for the most of the traits *viz.*, kernel length, kernel breadth, kernel L/B ratio, kernel length after cooking, kernel breadth after cooking, linear elongation ratio and breadth wise expansion ratio (Table 1). It indicated that all the crosses might have attained homozygosity in F₄ itself for these traits. These results were in accordance with Nandeshwar et al. (2010) who studied F₂ progenies which was derived from the crosses IR-62 x Samba mashuri and Kunti x Dudheswar and the report of Kumar et al. (2006) in rice F₂ generation for kernel length, kernel breadth recorded low genotypic and phenotypic coefficient of variability in both the generations for all the crosses. Similar findings were already have been reported by Umadevi et al. (2010) in 110 rice genotypes.

Table 1. Variability studies for quality traits in rice with high iron and zinc content in F₄ & F₅ Generation.

Crosses	ADT 37 x IR68144-3B-2-2-3								TRY (R) 2 x Mapillaisamba							
	GCV		PCV		Heritability		GA as a percent of mean		GCV		PCV		Heritability		GA as a percent of mean	
	F4	F5	F4	F5	F4	F5	F4	F5	F4	F5	F4	F5	F4	F5	F4	F5
Kernel length	1.436	2.966	1.760	3.025	66.66	96.15	2.41	5.99	1.446	0.947	1.524	1.059	90.00	80.00	2.82	1.74
Kernel breadth	3.658	3.073	3.911	3.366	87.50	83.33	7.04	5.77	2.985	3.309	3.948	3.625	57.14	83.33	4.64	6.22
Kernal L/B ratio	2.678	4.525	3.787	4.726	50.00	91.66	3.90	8.92	3.676	3.223	4.444	3.530	68.42	83.33	6.26	6.05
Kernel length after cooking	1.408	1.646	1.457	1.689	93.33	95.00	2.80	3.30	0.859	0.722	0.974	0.791	77.77	83.33	1.56	1.35
Kernel breadth after cooking	2.357	3.824	2.545	3.949	85.71	93.75	4.49	7.62	1.394	2.591	1.973	2.770	50.00	87.50	2.02	4.99
Linear elongation ratio	2.03	3.482	2.87	4.021	50.00	75.00	2.95	6.21	2.131	3.708	3.014	4.787	50.00	60.00	3.10	5.89
Breadth wise elongation ratio	4.392	6.002	5.810	6.416	57.14	87.50	6.83	11.56	3.615	2.962	4.667	3.627	60.00	66.66	5.76	4.98
Iron content	18.068	26.955	18.696	26.968	93.39	99.90	35.97	55.50	14.808	13.603	14.845	13.790	99.50	97.29	30.42	27.64
Zinc content	14.839	18.649	15.158	19.211	95.83	94.23	29.92	37.29	8.062	9.204	8.646	9.351	86.95	96.87	15.48	18.66
Single plant yield	12.81	13.69	13.77	15.32	86.51	79.89	24.54	25.22	11.82	5.20	12.38	5.50	91.19	89.48	23.25	10.14

The findings of Vanaja and Babu (2006) recorded low genotypic and phenotypic coefficient of variability for kernel length after cooking and kernel breadth after cooking, breadth wise expansion ratio in 10 quality parameters in set of 56 high yielding diverse rice genotypes which is corroborates the current investigation. For iron and zinc content moderate GCV and PCV was observed in F₅ generation. Hence, these traits need one or more generations in order to attain homozygosity. These results were in parallel with the findings of Kalaimaghal (2011) who observed moderate GCV and PCV for iron content in the in F₂ and F₃ the cross ADT37 x IR68144-3B-2-2-3.

High heritability coupled with high genetic advance was observed in ADT37 x IR68144-3B-2-2-3 only for iron and zinc content, except TRY(R)2 x Mapillaisamba which showed high heritability with moderate genetic advance. Hence, these traits were least influenced by environment and mostly governed by additive gene action. Therefore, there is scope for improvement of iron

and zinc content by exercising selection pressure on these two characters. These results were in agreement with Shanmuga sundara pandian (2007), Purusothaman (2010) and Aswini Shamak et al. (2011) in F₄ and F₅ generation. Low heritability with low genetic advance as percentage of mean was observed for kernel length, kernel breadth and kernel L/B ratio in all the crosses in both the generations. The findings of Jayasree (2007) and Umadevi et al. (2010) in genotypes observed medium to high heritability with high genetic advance as percentage of mean for kernel length after cooking, kernel breadth after cooking, linear elongation ratio and breadth wise expansion ratio which in contrast with the result of current investigation. Based on mean, GCV and PCV, heritability and genetic advance, it was understood that the progenies of ADT 37 x IR68144-3B-2-2-3 would be more useful for improving grain iron content with the desirable quality traits viz., kernel length, kernel breadth after cooking. Similarly TRY (R) 2 x Mapillaisamba

segregants could be used for improving the grain zinc content and breadth wise expansion ratio.

Conflict of Interest

The authors have not declared any conflict of interest.

REFERENCES

- Aswini Samak NR, Shailaja H, Shashidhar N, Hanumareddy B (2011). Exploratory studies on genetic variability and genetic control for protein and micronutrient content in F₄ and F₅ generation of rice. *Asian. J. Plant Sci.* 10(7):376-379.
- Anonymous (2007). *Agriculture and Industry Survey.* 17(11):18-20.
- Burton GW (1952). Quantitative inheritance in grasses. In: *Proc. 6th Inter Grassland Congr.* 1:277- 283.
- Bhuiyan NI, Paul DNR, Jabbar MA (2002). Feeding the extra millions by 2025 challenges for rice research and extension in Bangladesh. A keynote paper presented on National Workshop on Rice Research and Extension held on 29-31 January 2002 at BIRRI, Gazipur. P. 9.

- Graham RD, Senadhira D, Beebe S, Iglesias C, Monsterio I (1999). Breeding for micronutrient density in edible portions of staple food crops: Conventional approaches. *Field Crops Res.* 60:57-80.
- Jackson ML (1973). Soil and plant analysis. Prentice Hall of India Private Limited, New Delhi.
- Johnson HW, Robinson HF, Comstock RE (1955). Estimates of genetic and environmental variability in soybean. *Agron. J.* 47:314-318.
- Jayasree (2007). Characterization of landraces, cultivars and new plant types of rice for quantitative and grain quality traits. M.Sc., (Ag.) Thesis (Unpubl.), TNAU, Coimbatore.
- Kalaimaghal R (2011). Studies on genetic variability of grain iron and zinc content in F_2 , F_3 generation of rice (*Oryza sativa* L.). M.Sc. (Ag.) Thesis (Unpubl.), TNAU, Coimbatore.
- Kumar S, Gautam AS, Chandal C (2006). Estimates of genetic parameters for quality traits in rice (*Oryza sativa* L.) in mid hills of Himachal Pradesh. *Crop Res.* 32(2):206-208.
- Nandeshwar BC, Pal S, Senapati BK, De DK (2010). Genetic variability and character association among biometrical traits in F_2 generation of some Rice crosses. *Electron. J. Plant Breed.* 1(4):758-763.
- Panse VG, Sukhatme PV (1961). Statistical methods for agricultural workers. 2nd edition, Indian Council. Agric. Res. New Delhi. P. 22.
- Philip JW, Martin RB (2005). Biofortifying crops with essential mineral elements. *Trends Plant Sci.* 10:586-593.
- Purusothaman R (2010). Genetic analysis for high iron and zinc content in rice (*Oryza sativa* L.) grains. M.Sc. (Ag.) Thesis (Unpubl.), TNAU, Coimbatore.
- Shanmuga SG (2007). Studies on diversity and Genotype \times Environment interaction for micronutrient content in rice (*Oryza sativa* L.). M.Sc. (Ag.) Thesis. (Unpubl.), TNAU, Coimbatore.
- Umadevi M, Veerabhadran P, Manonmani S, Shanmugasundaram P (2010). Physico-chemical and cooking characteristics of rice genotypes. *Electron. J. Plant Breed.* 1(2):114-123.
- Welch RM, Graham RD (2004). Breeding for micronutrients in staple food crops from a human nutrition perspective. *J. Exp. Bot.* 55:353-364.