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Inheritance of aroma in aromatic rice (*Oryzasativa*L.)genotypes

Kalmeshwer Gouda Patil^{1,3}, Nagesh^{1,4}, Veeresh Gouda Patil^{1,3}, Kulkarni, R. S¹ and Shashidhar, H. E²

¹Department of Genetics and Plant Breeding, University of Agricultural Sciences, GKVK, Bangaluru, Karnataka, India

²Department of Biotechnology, University of Agricultural Sciences, GKVK, Bangaluru, Karnataka, India

³Monsanto, Bangaluru, Karnataka, India

⁴International Crop Research Institute for the Semi-Arid Tropics, Patancheru, Hyderabad, Andhra Pradesh, India

ABSTRACT

The inheritance pattern of aroma in rice (*Oryzasativa*L.) was carried out in the crosses among three non-aromatic and two aromatic varieties. All the F_1 plants of the three crosses were non-aromatic indicating that the gene controlling aroma in the parents was recessive. The segregation ratio of non-aromatic to aromatic plants was 3:1 in F_2 plants confirming the monogenic inheritance of aroma.

Keywords: rice, aroma, inheritance

INTRODUCTION

India produces some of the best quality rice in the world. Though a high-volume commodity, a class of aromatic, superfine premium rice has evolved its own market niches, making rice trade a commercial success internationally. A class of premium rice with specific grain characteristics, traditionally grown on either side of the Indus River, is popularly known as the basmati rice and these varieties are also known as “Queen of fragrance”. The superiority of basmati cultivars over other premium rice varieties is its superfine grains which have a distinct aroma, excellent elongation ability and the soft, flaky texture of the cooked rice [1]. Besides the much sought after basmati types which get high price in international markets, the country also abounds with hundreds of indigenous short grain aromatic cultivars and landraces grown in pockets of different states. Almost every state has its own collection of aromatic rice that performs well in native areas [2]. These aromatic rice lines also possess exemplary quality traits like aroma, fluffiness and taste. Even though these aromatic rice varieties fetch higher price in the market, they are susceptible to pest and diseases and are, generally low yielders. Therefore, rice breeders lay great emphasis on the development of high yielding varieties with quality grain and aroma.

From many investigations in the past, aroma in rice is known to be genetically controlled and its genetic behaviour is summarized as follows: Sood & Siddiq [3] reported, aroma is monogenic recessive to non-aromatic and Kadam & Patankar [4] reported it to be monogenic dominant. A digenic segregation of 9 non-aroma: 7 aroma Tripathi & Rao, [5], 15: 1 (Geetha, [6]), 13: 3 (Tsuzuki and Shimakawa, [7]) and trigenic ratio of 37 non-aroma: 27 aroma (Kadam and Patankar, [4] and Reddy and Sathyanarayanaiah, [8]) are also reported. Moreover, Dhulapannavar [9] showed four complementary genes indicating segregation ratio of 175 non-aroma: 81 aroma. Further, the recent inheritance studies on aroma indicated 3:1 (non-scented: scented) segregation ratio

indicating monogenic, recessive gene control (Berner and Hoff, 1986; Vivekanandan and Giridharan, 1994; Lorieux *et al.*, 1996)

There is no consensus as yet on the nature of inheritance in aromatic characters as described above. The lack of agreement among many investigators appears to be related to the differences in the aromatic varieties used and also the difficulties of the methods of evaluating aroma. The objective of the present study was to understand the nature of inheritance of aroma that may be useful in breeding and development of aromatic rice varieties.

MATERIALS AND METHODS

The experimental material for the present investigation comprised of parents, F₁ and F₂ populations of three crosses of rice (Table 1: see salient feature for parents). The rice varieties used were, IRRI 38, BI 33, Moro mutant, Jeerigesanna and Azucena. The aromatic genotype, Jeerigesanna is a local, late maturing, aromatic, low yielding variety whereas, Azucena is an exotic genotype from Philippines. The non-aromatic genotype, IRRI 38 is an exotic line while BI 33 is an advanced breeding line and Moro mutant is a natural mutant of Moroberekan cultivar. The parents and F₁ plants were grown in summer, 2007 at University of Agricultural Sciences, Bangalore, Karnataka, India and the F₂ seeds were harvested, threshed and cleaned. About 300 F₂ seeds of each cross combination were grown in *Kharif* 2007 under aerobic condition. After 60 days of seeding, determination of presence or absence of aroma was made according to the method described by Sood and Siddiq [3]. Two grams of green leaves were taken from individual plants cut into small pieces and kept in the test tubes. About 10 ml of 1.7% potassium hydroxide (KOH) solution was added to each test tube. The test tubes were covered immediately after the addition of alkali and left under room temperature for about ten minutes. The test tubes were opened one by one and the content in each was immediately evaluated by smelling. The samples were classified into two categories in the presence or absence of aroma.

RESULTS AND DISCUSSION

In the three crosses of IRRI 38 x Jeerigesanna, BI 33 x Jeerigesanna and Azucena x Moro mutant, the parents Jeerigesanna and Azucena were aromatic while, IRRI 38, BI 33 and Moro mutant were non-aromatic. The F₁ plants from these crosses were non-aromatic (Table 1).

The data obtained on the presence and absence of aroma of the F₂ plants in each of the crosses is presented in Table 2. In the cross IRRI 38 x Jeerigesanna, 91 out of 347 were aromatic and 256 were non-aromatic. 67 plants of 280 were aromatic in the cross BI 33 x Jeerigesanna while, 213 were non-aromatic. Similarly, 95 out of 378 were aromatic and 283 were non-aromatic in the cross Azucena x Moro mutant.

Table 1. Salient features of the parents used for the development of F₂ populations

Accession	Pedigree	Plant height	Duration	Grain type	Yield (t ha ⁻¹)	Salient features
IRRI 38	Line from IRRI, Philippines	Semidwarf	Early	Medium, slender	3.5	Drought tolerant
Jeerigesanna	Local line	Tall	Late	Short, bold	2.5	Aromatic
BI 33	Budda X IR64	Semidwarf	Medium early	Long, slender	4.5	Drought tolerant
Azucena	Exotic line	Tall	Late	Medium, bold	3.2	Aromatic
Moro mutant	Mutant from Moroberikan	Semidwarf	Medium early	Medium, bold	4.2	Drought tolerant

Table 2. Number of plants expressing aromatic and nonaromatic grain type according to KOH test

Parents	No. of plants tested	No. of plants	
		aromatic plants	nonaromatic plants
IRRI 38	10	0	10
BI 33	10	0	10
Moro mutant	10	0	10
Jeerigesanna	10	10	0
Azucena	10	10	0
F ₁ hybrids			
IRRI 38 X Jeerigesanna	10	0	10
BI 33 X Jeerigesanna	10	0	10
Azucena X Moro mutant	10	0	10

In all the aromatic x non-aromatic crosses, all F₁ hybrids were non-aromatic indicating recessive nature of the character. The F₂ population segregated into the ratio of 3 non-aromatic: 1 aromatic offspring indicating a monogenic Mendelian ratio. The calculated chi-square values were not differing from the table values fitting a monogenic Mendelian ratio at a very high level. The results showed that the aroma was controlled by a nuclear

recessive gene which agreed with those reported by Berner and Hoff [10] Vivekanandan and Giridharan [11] and Lorieux *et al.* [12].

Table 3. Inheritance pattern of aroma in all three F₂ populations of rice

Cross	No. of plants tested	No. of plants				Ratio	Chi-square	P value
		Non-aromatic plants		Aromatic plants				
		Observed	Expected	Observed	Expected			
IRRI 38 X Jeerigesanna	347	256	85.75	91	260.25	3:1	0.277	0.598
BI 33 X Jeerigesanna	280	213	70	67	210	3:1	0.171	0.678
Azucena X Moro mutant	378	283	94.50	95	283.50	3:1	0.003	0.952

REFERENCES

- [1] EA Siddiq and N Shobha Rani, 1998
- [2] N Shobha Rani and K Krishnaiah, *Science Publishers, Inc., USA*: **2001**, 49-78.
- [3] BC Sood, and EA Siddiq, *Indian J. Genet.*, **1978**, 38, 268– 271.
- [4] BS Kadam and VK Patankar, *Chron. Bot* **1938**, 4, 496-497.
- [5] RS Tripathi and MJBK Rao, *Euphytica*, **1979**, 28, 319-323.
- [6] S Geetha, *International Rice Research Notes*, **1994**, 19, 5-6.
- [7] E Tsuzuki, and E Shimakawa, *Euphytica*, **1990**, 46, 157-159.
- [8] PR Reddy and K Sathyanarayanaiah, *Indian J. Genet.*, **1980**, 40, 327-329.
- [9] CV Dhulappanavar, *Euphytica*, **1976**, 25, 659-662.
- [10] DK Berner and BJ Hoff, *Crop Science*, **1986**, 26, 876–878.
- [11] P Vivekanandan, and S Giridharan, *Oryza*, **1998**, 35, 242-245.
- [12] M Lorieux; M Petrov; N Huang; E Guiderdoni and AGhesquiere, *Theor. Appl. Genet.*, **1996**, 93, 1145-1151.