



Morpho-Cultural Variability Characterization of *Bipolaris oryzae* Causing Brown Leaf Spot of Rice in Jammu Sub-Tropics

Sonali Abrol ^{a*}, S. K. Singh ^a, V. B. Singh ^a, A. K. Singh ^b and Ranbir Singh ^a

^a Division of Plant Pathology, Sher-e-Kashmir University of Agricultural Sciences and Technology of Jammu, 180 009, India.

^b Division of Entomology, Sher-e-Kashmir University of Agricultural Sciences and Technology of Jammu, 180 009, India.

Authors' contributions

This work was carried out in collaboration among all authors. All authors read and approved the final manuscript.

Article Information

DOI: 10.9734/IJECC/2022/v12i111388

Open Peer Review History:

This journal follows the Advanced Open Peer Review policy. Identity of the Reviewers, Editor(s) and additional Reviewers, peer review comments, different versions of the manuscript, comments of the editors, etc are available here: <https://www.sdiarticle5.com/review-history/93194>

Original Research Article

Received 15 August 2022

Accepted 22 October 2022

Published 27 October 2022

ABSTRACT

Extensive fortnightly surveys were undertaken in major rice growing districts of Jammu division to study cultural and morphological variability amongst the isolates of brown leaf spot disease causing fungus (*Bipolaris oryzae*). Based on disease severity 24 isolates (BO₁ to BO₂₄) were divided into three groups (I, II and III). Group 'I' comprised 11 isolates with disease severity values between 51.22-67.44% and categorized under highly virulent and group 'II' comprised disease severity values between 43.17-48.17% and were categorized under moderately virulent isolates. Group 'III' included eight isolates with disease severity value between 26.11-38.17% and categorized under less virulent isolates. On the basis of colony growth pattern, margin and colour, the isolates were categorized into 4 groups (A, B, C and D). Group A comprised of six isolates had entire margin, circular form, umbonate elevation with smooth surface and black to fluffy growth. Eleven isolates were categorised under group B developed colonies with undulated margin, irregular form, umbonate elevation with grey fluffy growth and greyish colour. Group C comprised three isolates had undulated margin, irregular form, umbonate elevation with smooth surface. Group D contained four isolates developed entire margin, circular form; raised elevation with wrinkled surface, mycelium was grey which turned to black and showed grey to black

*Corresponding author: E-mail: abrol.sonali222@gmail.com;

pigmentation. The largest conidial size was recorded in isolate BO₁ (98.43-101×24.54-25.54 μm). The conidia developed mean septa of 4.5-7.4 with straight to curve shaped. The sporulation of different isolates of *B. oryzae* ranged between 6.1-7.6×10⁶ spores/ml.

Keywords: Spot disease; *Bipolaris oryzae*; leaf blade; leaf sheath; rice production; brown leaf spot.

1. INTRODUCTION

In tropical and sub-tropical countries rice production is considered as backbone of agricultural economy, including India, the world's 2nd-largest producer and consumer [1 and 2]. Rice cultivation in Jammu and Kashmir region is mostly mono-cropped, with far higher consumption and importance as a staple grain than other Indian states. Rice is highly essential in the lives of people in the state, despite the fact that the area under the crop cultivation is quite small relative to other Indian states, and hence is very significant in the state economy. In Indian sub-continent including Jammu and Kashmir region, Basmati rice is famous and distinguished all around the world for its appearance and aroma, but unfortunately rice production is reduced because of the biotic and abiotic stresses. Among the biotic constraints, fungal and bacterial diseases play a vital role in preventing normal growth of rice crops and hence reducing yield. Brown leaf spot disease caused by *Bipolaris oryzae* is one of the distinctive and most damaging diseases of rice in the world, because of the extensive distribution of numerous physiological races [3].

Brown leaf spot in rice, is known to cause major qualitative and quantitative losses (up to 90%) in rice crop, especially when the leaf spotting phase reaches epiphytic dimensions, as it did during the Great Bengal Famine of 1942 [4]. The disease has spread throughout the world's rice-growing regions. It has been discovered to be pandemic in locations with high rainfall, such as the Assam, West Bengal, the Malabar Coast and Himalayas [5]. The pathogen attacks the crop from seedling to milky stage. The characteristic symptoms appear as minute spots on the coleoptile, leaf blade, leaf sheath and glume, being most prominent on leaf blades and glumes. On leaves, typical spots are brown in colour with grey or whitish centre with typical yellow halo over the spot [6]. Typically conidia are slightly curved and widest at the middle, conidia are 5-10 septate with the oldest conidium towards base. The causal fungus *B. oryzae* remains in seeds for most of its lifecycle, but can also persist on infested rice stubble and straw. It spreads from plant to plant in the field by the airborne spores.

A relative humidity of >89% at 25°C leads to successful inoculation by conidia and infection can be vigorous upon free water on leaf surface [5]. Diversity and pathogenicity of the rice brown spot pathogen were investigated earlier by many researchers using morpho-cultural characteristics as well as genetic fingerprint analysis [7,8,9,10,11]. The present paper emphasised on the morpho-cultural characterization of *B. oryzae* and identifying the most virulent isolates.

2. MATERIALS AND METHODS

2.1 Isolation of Pathogen

Naturally diseased leaves of different rice cultivars infected in varying degrees with brown leaf spot were collected from different locations of Jammu sub-tropics including Jammu, Samba, Kathua and Udhampur districts, and were brought to laboratory of the Division of Plant Pathology, SKUAST-Jammu India, for isolation of the pathogenic fungus. The collected samples were critically observed under microscope, to know the presence of pathogen in infected plant part. After confirming the presence of the pathogen under microscope, the diseased tissues from infected plant parts were subjected to isolation. The pathogen *B. oryzae* was isolated by standard hyphal tip isolation procedure and then the culture was maintained on Potato Dextrose Agar (PDA) slants and kept in a refrigerator at 5°C for further use in all the laboratory studies.

2.2 Pathogenicity Test

The pathogenicity test of *B. oryzae* (isolates BO₁ to BO₂₄) was determined on susceptible cv. Pusa Basmati-1121 under both greenhouse and field experiments. Rice seedlings were sprayed at 25 days old under greenhouse and field conditions with the spore suspension (3 x 10⁵ spore ml⁻¹) of the 24 isolates of *B. oryzae* separately. The seedlings were sprayed until runoff occurred. Three replicates were used for each isolate.

2.3 Cultural Variability

Mycelial disc (5 mm) of 10 days old culture of each isolate was transferred to the centre of

sterilized Petri plates containing PDA medium and incubated at $25\pm 2^{\circ}\text{C}$. Colony characters viz. colour and diameter were recorded after ten days of inoculation. Reverse side of cultural plate of each fungal isolate was also observed to record pigmentation on under side of the plate.

2.4 Morphological Variability

The morphological variation among different isolates of *B. oryzae* was studied on PDA under *in vitro* conditions. Mono conidial culture of each isolate was first grown on PDA medium and then semi permanent shades were prepared from 10 day old culture were stained with cotton blue in lacto phenol. The growth pattern, progressive colony growth and colony colour was recorded on PDA medium. Morphological features of the pathogen viz., type of mycelia, width of mycelia, sporulation pattern, spore size and shape were recorded.

Spores of *B. oryzae* of all the isolates from the culture were mounted on a clean glass slide. Spores were mixed with Lactophenol thoroughly in order to obtain uniform spread, on which cover slip was placed. Spores were measured under high power objective using light microscope (400X). The average size of the spores like length, width and number of septa were recorded. Microphotographs were taken to show the typical spore morphology of the pathogen.

3. RESULTS AND DISCUSSION

3.1 Pathogenicity Test

The pathogenicity test of *B. oryzae* (isolates BO₁ to BO₂₄) was determined on susceptible cv. Pusa Basmati-1121 (Table 1). Pathogenicity test was conducted for all the 24 isolates and control pots were maintained by spraying distilled water only. The first symptoms on the affected plant part of rice started with minute pin head shape brown flecks after 4-5 days of inoculation, later such pin head shaped flecks matured into dark brown lesions characteristics of brown spot disease. On the basis of disease severity the isolates were divided into three groups (I, II and III). Group 'I' comprised 11 isolates viz., BO₁, BO₂, BO₃, BO₄, BO₅, BO₈, BO₉, BO₁₂, BO₁₄, BO₁₉ and BO₂₀ with disease severity value between 51.22-67.44% and were categorized under highly virulent. In group 'II' comprised five isolates viz., BO₆, BO₁₁, BO₁₃, BO₁₅ and BO₁₆ with disease severity value between 43.17-48.17% and were

categorized under moderately virulent isolates. Group 'III' included eight isolates viz., BO₇, BO₁₀, BO₁₇, BO₁₈, BO₂₁, BO₂₂, BO₂₃ and BO₂₄ with disease severity value between 26.11-38.17% and were categorized under less virulent isolates (Table 2). The pathogen was re-isolated from infected leaves and its morphological characters were compared with the original culture of the pathogen and found similar in all respects i.e., morphological and microscopic. Hence, the causal agent of the brown spot disease was confirmed as *B. oryzae* after confirming of Koch's postulates. However, no such symptoms developed on uninfected leaves even after 12 days of inoculation.

The symptoms observed were in accordance with the description of the pathogen given by Sobanbabu et al. [12] who also reported variation with respect to disease incidence based on pathogenicity test. Similarly, Nazari et al. [9] also conducted pathogenicity test involving 12 isolates of *B. oryzae* under greenhouse conditions and found that isolates were pathogenic to rice seedlings in cv. Tarom. Moreover, Singh et al. [13] also reported variation in virulence among *B. oryzae* isolates based on pathogenicity test on the susceptible variety Pusa-1121.

3.2 Cultural Variability in Different Isolates of *B. oryzae* on PDA Medium

All the 24 isolates, when grown in sterilized Petri plates on sterilized PDA medium, exhibited great variability for colony diameter, growth pattern, colony margin and colour (Table 3) after 15 days of incubation at $25\pm 1^{\circ}\text{C}$. On the basis of colony growth pattern, margin and colour, the isolates were categorized into 4 groups (A, B, C and D); Group A comprised six isolates viz., BO₁, BO₆, BO₉, BO₁₁, BO₂₂ and BO₂₃ and had colony diameter of 85.67, 78.65, 81.41, 63.51, 70.34 and 64.89 mm, respectively. In addition, the isolates had entire margin, circular form, umbonate elevation with smooth surface and black to fluffy growth. Eleven isolates viz., BO₃, BO₅, BO₇, BO₈, BO₁₀, BO₁₄, BO₁₅, BO₁₇, BO₁₈, BO₁₉ and BO₂₀ were categorised under group B having colony diameter of 63.51, 70.34, 73.85, 60.32, 65.11, 81.63, 86.11, 70.43, 71.44, 71.56 and 68.24 mm, respectively. Isolates of this group developed colonies with undulated margin, irregular form, umbonate elevation with grey fluffy growth and greyish colour. Group C comprised three isolates viz., BO₄, BO₁₆ and BO₂₁ and had colony diameter of 86.96, 72.34

and 48.34 mm, respectively. The colonies of the isolates had undulated margin, irregular form, umbonate elevation with smooth surface. Initially, the mycelium was dirty white and later turned to grey in colour. Group D contained four isolates viz. BO₂, BO₁₂, BO₁₃ and BO₂₄ and had colony diameters of 48.34, 78.84, 58.54, and 71.44 mm, respectively. The colonies developed an entire margins, circular form, raised elevation with wrinkled surface, mycelium was grey which turned to black and showed grey to black pigmentation.

The results obtained in the present investigation showing variation in colony colour of *B. oryzae* corroborate with the findings of Kumar et al. [14] who reported that color of the isolates of *B. oryzae* showed huge variation i.e., black with

fluffy growth, grey with fluffy growth, grey with fluffy growth and white spots, grey with suppressed growth. Kumar et al. [10] grouped *B. oryzae* isolates into 5 categories on the basis of morphology and growth pattern i.e. black with fluffy growth, black with suppressed growth, grey with cottony growth, grey and white mix with cottony growth and white with cottony growth. The margins of colonies varied from circular, undulated, irregular having umbonate to raised elevation along with smooth to wrinkled surface. Nayak and Hiremath [15] also reported variations with respect to colony colour (Dark greyish black, light greyish, greyish to white, greyish black to white, black to white colour), mycelial growth (cottony and flat), margin (regular to irregular), sporulation and conidial morphology among *B. oryzae* isolates.

Table 1. Pathogenicity test of different isolates of *B. oryzae* on cv. Pusa Basmati-1121

District	Location	Isolate of <i>Bipolaris oryzae</i>	Disease severity (%)	Virulence pattern
Jammu	Bishnah	BO ₁	51.33 (45.74)	++++
		BO ₂	62.53 (52.23)	++++
	Marh	BO ₃	59.67 (50.55)	++++
		BO ₄	64.20 (53.22)	++++
		BO ₅	58.07 (49.62)	++++
		BO ₆	44.17 (41.63)	+++
		BO ₇	26.11 (30.71)	++
Samba	Vijaypur	BO ₈	54.08 (47.32)	++++
		BO ₉	51.22 (45.68)	++++
		BO ₁₀	38.00 (38.03)	++
		BO ₁₁	48.17 (43.93)	+++
	Ghagwal	BO ₁₂	53.22 (46.82)	++++
		BO ₁₃	43.17 (41.05)	+++
		BO ₁₄	52.11 (46.19)	++++
Kathua	Hiranagar	BO ₁₅	44.89 (42.04)	+++
		BO ₁₆	47.44 (43.51)	+++
	Bilawar	BO ₁₇	35.44 (36.52)	++
		BO ₁₈	38.17 (38.14)	++
	Kathua	BO ₁₉	67.44 (55.19)	++++
		BO ₂₀	52.50 (46.41)	++++
Udhampur	Udhampur	BO ₂₁	35.00 (36.25)	++
		BO ₂₂	32.55 (34.77)	++
	Tikri	BO ₂₃	34.44 (35.92)	++
		Manwal	BO ₂₄	35.17 (36.35)
	CD (p=0.05)	-	-	1.28

Table 2. Grouping of isolates of *Bipolaris oryzae* on the basis of pathogenicity test

Group	Isolate	Disease severity (%)	Virulence pattern
Group-I	BO ₁ , BO ₂ , BO ₃ , BO ₄ , BO ₅ , BO ₈ , BO ₉ , BO ₁₂ , BO ₁₄ , BO ₁₉ and BO ₂₀	>50	Highly virulent
Group-II	BO ₆ , BO ₁₁ , BO ₁₃ , BO ₁₅ and BO ₁₆	40-50	Moderately virulent
Group-III	BO ₇ , BO ₁₀ , BO ₁₇ , BO ₁₈ , BO ₂₁ , BO ₂₂ , BO ₂₃ and BO ₂₄	<40	Less virulent

Table 3. Cultural variation of isolates of *Bipolaris oryzae* on potato dextrose agar medium

Isolate	Radial growth (mm)	Group	Colony characteristics	Pigmentation
BO ₁	85.67	A	Entire margin, circular form, umbonate elevation with smooth surface, black with fluffy growth	Black
BO ₆	78.65			
BO ₉	81.41			
BO ₁₁	63.51			
BO ₂₂	70.34			
BO ₂₃	64.89			
BO ₃	63.51	B	Undulated margin irregular, umbonate elevation with grey fluffy growth	Greyish
BO ₅	70.34			
BO ₇	73.85			
BO ₈	60.32			
BO ₁₀	65.11			
BO ₁₄	81.63			
BO ₁₅	86.11			
BO ₁₇	70.43			
BO ₁₈	71.44			
BO ₁₉	71.56			
BO ₂₀	68.24			
BO ₄	86.96	C	Undulated margin, irregular form, umbonate elevation, smooth surface, mycelium initially dirty white and later turning to grey	Mixture of grey
BO ₁₆	72.34			
BO ₂₁	48.34			
BO ₂	48.34	D	Entire margin, circular form, raised elevation with wrinkled surface, mycelium grey turning to black.	Grey to black
BO ₁₂	78.84			
BO ₁₃	58.54			
BO ₂₄	71.44			

Table 4. Morphological variation of conidia of different isolates of *Bipolaris oryzae*

Isolate	Mean		Size of conidia (µm) Range	Number of septa	Number of conidia/ml (1×10 ⁶)	Shape of Spore
	Length (µm)	Breadth (µm)				
BO ₁	99.71	25.04	55.34-57.32 x 12.22-15.23	7.1	7.4	Curved
BO ₂	56.33	13.72	59.08-60.43 x 10.25-11.23	6.6	6.9	Straight
BO ₃	59.75	10.74	98.54-99.43 x 23.44-26.43	4.5	6.6	Curved
BO ₄	98.98	24.93	82.56-84.34 x 15.63-33.21	5.6	7.2	Curved
BO ₅	83.45	24.42	57.21-58.44 x 12.54-14.33	6.9	7.6	Straight
BO ₆	57.82	13.43	89.43-90.11 x 14.32-15.43	5.9	7.2	Curved
BO ₇	86.77	14.87	38.87-39.54 x 9.54-10.43	6.4	6.3	Curved
BO ₈	39.20	9.98	95.87-99.32x18.45-21.32	4.9	6.5	Curved
BO ₉	97.59	19.88	51.39-55.34x14.32-16.43	6.1	7.2	Curved
BO ₁₀	53.36	15.37	80.32-85.39x19.20-20.23	6.3	6.1	Curved
BO ₁₁	82.85	19.71	89.21-93.33x23.54-24.32	5.2	7.2	Curved
BO ₁₂	91.27	23.93	59.23-60.12x 12.32-14.34	6.9	6.4	Curved
BO ₁₃	59.67	13.33	89.33-92.12x12.56-14.32	5.0	7.4	Curved
BO ₁₄	90.72	13.44	69.32-71.65x12.23-15.77	4.8	7.6	Curved
BO ₁₅	70.48	14.00	83.34-87.33x11.34-13.34	5.6	6.9	Straight
BO ₁₆	85.33	12.34	78.44-89.34x9.54-10.55	4.9	6.2	Straight
BO ₁₇	83.89	10.04	80.34-81.44 x12.33-14.32	5.0	6.9	Curved
BO ₁₈	80.89	13.32	39.33-42.56x12.33-14.33	7.4	7.5	Curved
BO ₁₉	40.94	13.33	59.23-60.12x12.32-14.34	7.3	6.5	Curved
BO ₂₀	59.67	13.33	89.33-92.12x12.56-14.32	6.7	7.2	Straight
BO ₂₁	90.72	13.44	69.32-71.65x12.23-15.77	7.1	7.2	Curved
BO ₂₂	70.48	14.05	89.21-93.33x23.54-24.32	5.8	7.2	Curved
BO ₂₃	91.27	23.93	98.43-101 x 24.54-25.54	5.4	7.2	Curved
BO ₂₄	59.67	13.33	55.34-57.32 x 12.22-15.23	6.4	6.5	Curved

3.3 Morphological Variability in Different Isolates of *B. oryzae*

The isolates of *B. oryzae*, grown on PDA culture medium, developed conidial length ranged between 39.20-99.71 μm and breadth between 9.98-25.04 μm . The largest conidial size was recorded in isolate BO₁ (98.43-101 \times 24.54-25.54 μm), followed by BO₄ (98.54-99.43 \times 23.44-26.43 μm) and it was minimum in isolate BO₈ (38.87-39.54 \times 9.54-10.43 μm). The conidia developed mean septa of 4.5-7.4 with straight to curved shaped (Table 4). The sporulation of different isolates of *B. oryzae* ranged between 6.1-7.6 \times 10⁶ spores/ml.

Existence of variability among the isolates of *B. oryzae* with respect to conidial size was well documented in earlier studies. Ou [16] reported that size of conidia of *B. oryzae* isolates varied from 45-106 \times 14-17 μm in India. Jaiganesh and Kannan [17] reported that the colour of the conidia was brown to light brown and it was slightly curved with a bulge in the middle and tapering towards the ends. They also reported that the size of the conidia varied from 29.3-33.2 μm length and 13.5-14.8 μm width.

4. CONCLUSION AND RECOMMENDATIONS

The pathogenicity test of *B. oryzae* isolates was determined on susceptible rice cultivar Pusa Basmati-1121. The disease severity ranged between 26.11-67.44 per cent, 11 isolates of *B. oryzae*, with higher disease severity (51.22-67.44%) were categorized as highly virulent, whereas five and 8 isolates with disease severity between 43.17-48.17 and 26.11-38.17% were moderately and less virulent, respectively.

All the 24 test isolates exhibited great variability when cultured on PDA medium with colony colour varying from black, grey and whitish grey. The margins of colonies were circular, irregular and undulated and the colonies showed umbonate and raised elevation along with smooth and wrinkled surface. On the basis of colony growth pattern, margin and colour, the isolates were categorized into 4 groups (A, B, C and D). Group A comprised six isolates viz., BO₁, BO₆, BO₉, BO₁₁, BO₂₂ and BO₂₃ whereas, 11 (BO₃, BO₅, BO₇, BO₈, BO₁₀, BO₁₂, BO₁₄, BO₁₅, BO₁₇, BO₁₉ and BO₂₀), 3 (BO₄, BO₁₆ and BO₂₁) and 4 (BO₂, BO₁₃, BO₂₃ and BO₂₄) isolates were categorized under group B, C and D, respectively.

The isolates of *B. oryzae*, grown on PDA culture medium, developed conidial lengths ranging between 39.20-99.71 μm and breadth between 9.98-25.04 μm . The conidia developed mean septa of 4.5-7.4 with straight to curve shaped. The sporulation of different isolates of *B. oryzae* ranged between 6.1-7.6 \times 10⁶ spores/ml.

Investigating the response of the varieties to the isolates will enable us to appreciate the level of resistance of those varieties. To well understand the level of resistance of these varieties to brown leaf spot disease, it would be important to test them in field conditions and to understand the effects of abiotic factors on their behavior. Selecting genotypes with horizontal resistance could help us in taking part in achieving the set goals of global food security.

ACKNOWLEDGEMENT

The author acknowledges the support and help of all the members of Division of Plant Pathology, SKUAST-Jammu India.

COMPETING INTERESTS

Authors have declared that no competing interests exist.

REFERENCES

1. Nazir A, Bhat MA, Bhat TA, Fayaz S, Mir MS, Basu U, Ahanger SA, Altaf S, Jan B, Lone BA, Mushtaq M. Comparative analysis of rice and weeds and their nutrient partitioning under various establishment methods and weed management practices in temperate environment. *Agronomy*. 2022;12(4):816.
2. Bhat TA, Kanth RH, Jan B, Nazir A, Ahanger SA, Mir MS, Naikoo NB, Fayaz S, Dar KA, Gul A, Mansoor T. Real-time nitrogen application of rice varieties based on leaf colour chart under system of rice intensification in temperate climate. *Agronomy*. 2022;12(9):2229.
3. Arshad HM, Khan JA, Jamil FF. Screening of rice germplasm against blast and brown spot diseases. *Pakistan Journal of Phytopathology (Pakistan)*. 2008;25:84-90.
4. Ghoze RL, Ghatge MB, Subramamanyan V. *Rice in India* (revised edition). New Delhi, India, Indian Council of Agricultural Research. 1960;474.
5. Abrol S, Singh S, Singh V, Basu U, Singh R, Singh A, Ahanger Sa, Mehta A, Singh

- S, Kakraliya Dp, Bhagat S. Effect of agromet conditions on the progression of brown leaf spot disease in Basmati-370 rice. Asian Jr. of Microbiol. Biotech. Env. Sc. 2022;24(2):335-340.
6. Abrol S, Singh SK, Mehta A, Ahanger SA, Basu U, Vaid A, Singh R, Singh AK, Singh VB. Distribution pattern and prevalence of brown spot of rice (*Bipolaris oryzae*) in Jammu region. The Parma Innovation. 2022;11(2):732-736.
 7. Ouedraogo I, Correll JC, Boza EJ, Cartwright RD, Lee FN, Sankara P. Pathogenic, molecular, and genetic diversity among *Bipolaris*, *Drechslera*, and *Exserohilum* species on rice. BR Wells Rice Research Studies. Fayetteville, AR, USA, AAES Research Series. 2004;529: 111-9.
 8. Kamal MM, Mia MA. Diversity and pathogenicity of the rice brown spot pathogen, *Bipolaris oryzae* (Breda de Haan) Shoem. in Bangladesh assessed by genetic fingerprint analysis. Bangladesh Journal of Botany. 2009;38(2):119-25.
 9. Nazari S, Javan-Nikkhah M, Fotouhifar KB, Khosravi V, Alizadeh A. *Bipolaris* species associated with rice plant: Pathogenicity and genetic diversity of *Bipolaris oryzae* using rep-PCR in Mazandaran province of Iran. Journal of Crop Protection. 2015;4(4): 497-508.
 10. Kumar A, Solanki IS, Akhtar J, Gupta V. Morpho-molecular diversity of *Bipolaris oryzae* causing brown spot of paddy. Indian Journal of Agricultural Sciences. 2016;86(5):615-20.
 11. Boka A, Bouet A, Tiendrebeogo A, Kassankogno AI, Ouedraogo I, Nda GN, Denezon OD, Adiko A. Pathogenic variability of *Bipolaris oryzae* causing leaf spot disease of rice in West Africa. International Journal of Phytopathology. 2018;7(3):103-10.
 12. Sobanbabu G, Sabarinathan KG, Parthiban VK, Ramamoorthy V. Isolation, screening and identification of virulent Isolates of *Bipolaris oryzae* causing rice brown spot and *Sarocladium oryzae* causing sheath rot disease. Int. J. Curr. Microbiol. App. Sci. 2018;7(9):930-9.
 13. Singh HP, Raigar OP, Basandrai D, Sharma N, Upmanyu S, Basandrai AK. Screening for blast resistance in aromatic and basmati rice under north-western himalayas. Plant Disease Research. 2021;36(1):62-8.
 14. Kumar P, Anshu V, Kumar S. Morpho-pathological and molecular characterization of *Bipolaris oryzae* in Rice (*Oryzae sativa*). Journal of Phytopathology. 2011;159(1):51-6.
 15. Nayak MS, Hiremath SV. Cultural, morphological and molecular characterization of *Bipolaris oryzae* causing brown leaf spot of rice in Northern Karnataka. Journal of Pharmacognosy and Phytochemistry. 2019;8(2):1235-9.
 16. Ou SH. Rice diseases. IRRI; 1985.
 17. Jaiganesh V, Kannan C. Studies on the cultural characters and pathogenicity studies of brown leaf spot of rice caused by *Helminthosporium oryzae* (Syn: *Bipolaris oryzae*). Plant Archives. 2019;19: 585-7.

© 2022 Abrol et al.; This is an Open Access article distributed under the terms of the Creative Commons Attribution License (<http://creativecommons.org/licenses/by/4.0>), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

Peer-review history:

The peer review history for this paper can be accessed here:
<https://www.sdiarticle5.com/review-history/93194>