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Volatile organic compounds emitted by *Trichoderma* spp. for growth promotion and management of rice diseases

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This study was conducted to evaluate the antagonistic potential of volatiles emitted from seven *Trichoderma* spp.against 4 major rice diseases and to investigate their role in enhancement of growth and vigour of rice plants. Isolates were identified based on both morphological characteristics as well as molecular assay. Their biocontrol potential was evaluated and *T. erinaceum* completely inhibited the growth of *Pyricularia oryzae* and *Bipolaris oryzae*. It also inhibited growth of *Ceratorhiza oryzae-sativae* and *Rhizoctonia solani* by 64 and 50%, respectively. Sandwich plate method showed that volatiles emitted from *T. erinaceum* could reduce the mycelial growth of all four pathogens. GC-MS analysis indicated emission of 6-Pentyl- 2H-pyran-2-one from *T. erinaceum* in large quantity. Seed treatment with *Trichoderma* enhanced vegetative and reproductive growth of rice cultivar, Sahbhagidhan. Over-expression of defense enzymes like catalase and superoxide dismutase in shoot and peroxidase in both root and shoot were observed in *Trichoderma* treated plants. Results of this study suggest that *T. erinaceum* obtained from the rice rhizosphere can be used as a potential biocontrol agent against major rice diseases and as a growth inducer. It can also be utilized as an efficient source to produce 6-Pentyl-2H-pyran-2-one for further use.

Keywords: 6-pentyl-2H-pyran-2-one, induced resistance, Sahbhagidhan, Trichoderma erinaceum, VOCs

INTRODUCTION

Rice (Oryza sativa L.) provides food for more than half of the world's population. Major challenges for sustainable rice productions are different biotic and abiotic stresses which cause severe loss to rice productivity. To prevent biotic stress, rice cultivation relies upon chemical pesticides, which further deteriorate soil health by doing harm to beneficial soil microflora and causing emergence of resistant pathogens (Bhattacharyya et al. 2015). To prevent such limitations, biocontrol agents (BCAs) could be a suitable alternative. Trichoderma is a diverse and ubiquitous genus of fungi that is commonly found in various habitats, including soil, water, and plant surfaces (Mukherjee et al. 2013, Gal-Hemedet al. 2011, Harman et al. 2004).

The genus comprises over 200 species, which can be classified into several morphological and

molecular groups (Atanasova *et al.* 2013). *Trichoderma* spp. are free-living opportunistic symbionts that are commonly found in rhizosphere of many plants such as rice (Naeimi *et al.*2010; Khalili *et al.* 2012), maize (Harman *et al.* 2004), avocado (Ruano-Rosa and Hererra, 2009), tomato (Tsahouridou and Thanassoulopoulos, 2002; Harman *et al.* 2004), *Acacia* (Mukherjee *et al.* 2013) etc. They were also isolated from tree barks (Doni *et al.* 2014; Swain *et al.* 2021).

Trichoderma spp. are among the most studied organisms in the field of biocontrol agents, and they are also intensively explored for plant growth promotion. They have been reported as effective biocontrol agents against *Bipolaris oryzae* (Khalili *et al.* 2012), *Fusarium oxysporum* (Sundaramoorthy and Balabaskar, 2013), *Rhizoctonia solani* (Kamal and Devi., 2012, Chen *et al.* 2015), *Aspergillus flavus* (Anjaiah*et al.* 2006) and many other phytopathogens. *Trichoderma* spp. have been reported recently to be biocontrol agents in

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[J.Mycopathol.Res :

rice (Swain et al. 2018, 2021). Many traits are possessed by Trichoderma that make them suitable biocontrol agents (Howell, 2003). They show mycoparasitism (Siametoet al. 2011; Gajera et al. 2012; Herath et al. 2015) and rhizosphere competence (Tsahouridou and Thanassoulopoulos, 2002). Production of cell wall degrading enzymes (Marco and Felix, 2007; Gajera et al. 2012; Agrawal and Kotasthane, 2012), antibiotics (Swain and Mukherjee, 2020) and volatile compounds (Li et al. 2018) also explains successful disease control by Trichoderma. Mechanisms such as induction of defense responses (Swain et al. 2021) and metabolism of germination stimulants (Howell, 2003) are add-on to the biocontrol potential of Trichoderma.

Several studies have reported the growthpromoting ability of *Trichoderma* in various plants (Harman *et al.* 2004;Srivastava *et al.* 2006;Vinale *et al.* 2008; Salas-Marina *et al.* 2011;Bharti *et al.* 2012;Nagaraju*et al.* 2012). The authors attributed the growth-promoting effects to the production of auxin-like compounds.*Trichoderma* spp. execute their growth promoting activity by producing phytohormones (Harman *et al.* 2004; Contreras-Cornejo *et al.* 2009; Shoresh*et al.* 2010; Harman, 2011) and mobilizing plant nutrients which in return provides higher yield (Kapriand Tewari, 2010).

In addition to the above mechanisms, Trichoderma species have genes that code for the generation of secondary metabolites such as volatile organic compounds (VOCs), which have biocontrol properties (Kubicek et al. 2011; Mukherjee et al. 2012). Few studies indicated the role of VOCs for the biocontrol activity of Trichoderma spp. (Contreras-Cornejo et al. 2014; Li et al. 2018). Trichoderma VOCs had shown strong inhibition of plant pathogenic fungi such as Fusarium oxysporum, Rhizoctonia solani, Sclerotium rolfsii, Sclerotinia sclerotiorum, and Alternaria brassicicola (Amin et al. 2010; Meena et al. 2017). Trichoderma spp. emit VOCs differently depending on the species, strains, cultivation environment, and substrate composition (Stoppacher et al. 2010). More than 480 different VOCs have been found in Trichoderma species, and hydrocarbons,

heterocycles, aldehydes, ketones, alcohols, phenols, thioalcohols, thioesters, and thioesters are being the most common (Siddiquee, 2014). In addition, several Trichoderma VOCs have been observed to promote plant growth directly (Hung et al. 2013; Lee et al. 2016, 2019; Nieto-Jacoboet al. 2017). Even in the absence of pathogen, Arabidopsis thaliana exposed to VOCs emitted by T. viride had enhanced fresh weight, root growth, and chlorophyll content (Hung et al. 2013). One of the major compounds detected in diverse Trichoderma species is 6-pentyl-2Hpyran-2-one (6-PP). Application of 6-PP resulted in growth promotion and a reduction in disease symptoms (Hung et al. 2013). It is also reported that all Trichoderma species did not synthesize 6-PP and yet they are able to induce plant growth promotion (Atanasova et al. 2013, Kottb et al. 2015). This suggests there may be other molecules responsible for such actions.

In this study, VOCs of seven different *Trichoderma* isolates were tested for their biocontrol and plant growth promotion activities. Among the seven isolates, two isolates were chosen for chemical characterisation of VOCs. This experiment aims to justify the role of volatile compounds produced by *Trichoderma* in growth promotion and biocontrol in rice.

MATERIALS AND METHODS

Isolation and characterisation

Seven antagonistic Trichoderma strains were isolated from rice rhizospheric soil of National Rice Research Institute, Cuttack, Odisha, India by soil dilution technique (Johnson et al. 1959). Single spore technique was used to isolate pure colonies (Khalili et al., 2012; Mukherjee et al. 2014). The isolates were stored on PDA slants in refrigerator as well as in deep freezer at -80 °C (Swain et al. 2018). The fungal pathogens used in this study were Rhizoctonia solani, CRRI-RS-8 (MTCC-12232) causing sheath blight of rice, Ceratorhiza oryzae-sativae, CRRI-RS-4 (MTCC-12231) causing aggregate sheath spot, Pyricularia oryzae, CRRI-PO-2 (KX881382.1) causing rice blast and Bipolaris oryzae, CRRI-BS-2 (ACC:KX881383.1.) causing brown-spot of rice.

The cultural and morphological characteristics of the isolated Trichoderma species were studied by growing them on Potato Dextrose Agar medium (PDA) at 27°C. Identification was based on the morphological and taxonomic keys provided by Gams and Bissett (1998). Genomic DNA was isolated from the pure cultures of Trichoderma strains (Mukherjee et.al., 1995). The ITS regions of 5.8 S r RNA gene were amplified using the primers ITS1F and ITS4R (Toju et al. 2012). The sequences were compared with those from GenBank using the BLAST program and a phylogenetic tree was constructed through http:/ /www.phylogeny.fr. (Swain et al.2018). The DNA sequences of the 6 Trichoderma isolates, and four pathogens have been submitted to NCBI GenBank (Table1). Four isolates were deposited at Microbial Type Culture Collection (MTCC),IMTech, Chandigarh.

Confrontation assay

Preliminary investigation on antagonistic potential of the *Trichoderma* isolates were carried out by confrontation assays against all the four pathogenic strains. Actively growing culture disc of the pathogen and that of antagonists were placed opposite to each other on the petri dish (diameter of 9 cm) containing PDA. Plates inoculated with pathogen alone were considered as control.

The percentage of inhibition was calculated using formula:

RI= 100* (R2-R1)/ R2.

Where RI is the percentage of ratio inhibition (Kumar et al., 2012), R1 was the growth of pathogen from the pathogen inoculums towards the inoculums of *Trichoderma* isolate and R2 was the growth of pathogen in control plates.

Effect of volatiles from Trichoderma on pathogens

Radial inhibition of pathogens by volatiles from *Trichoderma* was studied by sandwich plate method. Oatmeal agar plates were inoculated centrally with agar discs of pathogens, *Pyricularia oryzae*, *Bipolaris oryzae*. *Ceratorhiza oryzae*-sativae, *Rhizoctonia solani* and all the seven

Trichoderma were multiplied on PDA plates. The bottom petri plates containing the *Trichoderma* covered by another petri plate with pathogens were sealed together using parafilm and was incubated keeping the pathogen at top. The effect of volatile was recorded by measuring the percentage of radial growth inhibition (Anees *et al.* 2010; Li *et al.* 2018).

Effect of volatiles as plant growth promoters

One petriplate (diameter of 9 cm) containing Trichoderma culture on PDA was placed in the bottom of a closed chamber (2.5 L). In the same chamber, Karuna and Sahabagidhan seeds were placed on different petriplates containing watersoaked blotting paper. The chamber was closed with an air-tight lid and placed at 27°C for 8 days under a photoperiod of 10 h light/14 h dark. Uninoculated PDA plates were used as control. Three replications were performed for each treatment. In this study, we have used two Trichoderma isolates namely, TS-1 and TS-10 based on their maximum antagonistic activity (as studied in sandwich plate method), and fresh weight, dry weight and root and shoot lengths were measured (Tahir et al. 2017).

Characterization of volatile compounds

The fungus was cultured in 20 mL head space vial. After 48 h incubation, the vial was sealed using a crimper and was used for volatile characterization using a GC-MS (Trace 1300-TSQ9000, Thermo Scientific, USA) equipped with HS-SPME auto sampler (TriPlus RSH, ThermoScientific, USA) and a capillary column (TG-5MS; 30 m*0.25 mm, film thickness: 0.25 µm,Thermo Scientific, USA). For extraction of volatiles in headspace, the vials were incubated in the in-built head space agitator at 40 °C for 60 min with the agitator on/off for 10 seconds each. SPME fibre (1 cm, 50/30 im Divinylbenzene/ Carbon Wide Range/Polydimethylsiloxane, DVB/ CarbonWR/PDMS, Dark Gray, Thermo Scientific, USA), was used for concentration of the volatiles. The volatiles were concentrated for 30 mins on the fibre. The needle speed inside the vial was 20mm sec⁻¹. After the equilibrium time, the fibre was directly injected into split-split less injector port for 2 mins. The injector temperature

was kept at 250°C with a split flow of 5 mL min -¹and split ratio of 5:1. The fibre was conditioned at 250 °C for 1 min before the concentration and for 15 min post injection. The following oven temperature program was followed; temperature was increased from 35 to 140°C at a ramp rate of 3°C min⁻¹ and hold for 1 min, temperature was increased to 230°C at the ramp rate of 10°C min⁻¹ and hold for 3 min. Samples were ionized by the positive electron impact (EI) mode using electron energy of 70 eV. Helium (99.999 % purity) at the flow rate of 1 mL min ⁻¹ was used as the carrier gas. Both the MS transfer line temperature and ion source temperatures were set at 230 °C. The MS data was acquired for them as range of m/z of 50-500. The compounds were identified using NIST library and AMDIS software (NIST 2017).

Field Experiments

Treatment

Trichoderma strains were grown on PDA for 5 days and conidial spores were separated using 0.5 % sterile carboxy methyl cellulose (CMC). The spore concentration was adjusted to 10^7 spores mL⁻¹. Seeds of Sahbhagidhan were treated with the spore suspension and left to air dry for 2 h. The control seeds were treated with 0.5 % CMC instead of spore suspension. Seeds were sown in $1m^2$ plots with 15cmX15cm spacing in Randomized Block Design experiment with four replications.

Plant growth promotion

Plants were collected 8 weeks after sowing. Agronomical parameters such as shoot length, root length, fresh and dry weight of shoot and root was taken. Plants were harvested at maturity. Grains were threshed and dried to 15 % moisture content. Days to 50% flowering, panicle length, grain weight per hill, yield per plot, thousand grain weight and fertility percentage were noted.

Defence enzyme activity

Peroxidase (POD) activity was assayed by studying biotransformation of guaiacol spectrophotometrically (Nounjana *et al.* 2012). Catalase activity was determined by estimating the amount of enzyme required to break down H_2O_2 under the assay conditions described by Kar and Mishra (1976). The ability of superoxide dismutase to inhibit the photochemical reduction of nitroblue tetrazolium (NBT) was measured as the enzyme activity according to the method of Beauchamp and Fridovich (1971).

Statistical Analysis

Statistical analysis was done using R-software. Analysis of variance (ANOVA) was calculated and tabulated. Treatment means of all the parameters were compared through Tukey's HSD test at probability at 5%.

RESULTS AND DISCUSSION

Isolation and characterization

Pyricularia oryzae and Bipolaris oryzae were isolated from infected rice leaves at NRRI. Cuttack and cultures of Rhizoctonia solani and Ceratorhiza oryzae-sativae available in the laboratory were used for this study. The Trichoderma isolates were identified using both morphological and molecular characters (Figs.1 & 2). The PCR amplified products of ITS sequences of 5.8 S r RNA gene (Fig.3)were compared with those from GenBank using the BLAST program (Altschul et al., 1990). The DNA sequences of the 5 Trichoderma isolates were submitted to NCBI GenBank(Table1). Four isolates have been deposited at the Microbial Type Culture Collection and Gene Bank (MTCC), Chandigarh. These are CRRI-TS-1 (T. pleuroticola) as MTCC 12407, CRRI-TS-2 (T. pleurotum) as MTCC 12408, CRRI-TS-5(T. longibrachiatum)as MTCC 12409 and CRRI-TS-6 (T. harzianum) as MTCC 12410. Their phylogeny has been depicted in Fig.4.

Confrontation assays

Confrontation assays were performed using all the seven *Trichoderma* isolates against four plant pathogen strains, which exhibited varying degrees of antagonistic potential (Fig.5).Growth inhibition of the pathogens *viz.Pyricularia oryzae*, *Bipolaris oryzae*, *Ceratorhiza oryzae-sativae and Rhizoctonia solani* by all the seven *Trichoderma*

Shanti Prava Behera and others

Table 1: Rice rhizopheric Trichoderma isolates from ICAR-NRRI, Cuttack with respective NCBI accession numbers

Isolate Name	Species Name	Isolated from	Location	NCBI accession no
CRRI-TS-1	T. pleuroticola	Rice plant rhizosphere	CRRI, Cuttack	KR069050
CRRI-TS-2	T. pleurotum	Rice plant rhizosphere	CRRI, Cuttack	KR069051
CRRI-TS-3	T. koningiopsis	Rice plant rhizosphere	CRRI, Cuttack	-
CRRI-TS-5	T. longibrachiatum	Rice plant rhizosphere	CRRI, Cuttack	KR069052
CRRI-TS-6	T. harzianum	Rice plant rhizosphere	CRRI, Cuttack	KR069053
CRRI-TS-9	T. harzianum	Rice plant rhizosphere	CRRI, Cuttack	KR069054
CRRI- TS-10	T. erinaceum	Rice plant rhizosphere	CRRI, Cuttack	-

Table 2: Confrontation assay showing percentage of growth inhibition of pathogen by different Trichoderma isolates (BCA)

		Growth inl	hibition (%)		
 Trichoderma Isolates	P. oryzae	B. oryzae	C. oryzae-sativae	R. solani	
CRRI-TS-1	50.00 ^d	68.26 ^{cd}	56.14 ^{de}	73.59 ^b	
CRRI-TS-2	64.65°	62.72 ^d	50.88°	42.78 ^d	
CRRI-TS-3	100.00ª	65.67 ^d	70.24 ^b	54.03 ^{cd}	
CRRI-TS-5	83.86 ^b	60.55 ^d	65.90 ^{bc}	46.67 ^d	
CRRI-TS-6	87.66 ^b	82.21 ^b	98.25ª	86.15ª	
CRRI-TS-9	100.00ª	76.08 ^{bc}	59.54 ^{cd}	65.85 ^{bc}	
CRRI-TS-10	100.00ª	100.00ª	63.92 ^{bcd}	49.99 ^d	
CV (%)	2.89	4.40	4.22	7.31	
Tukey's HSD at 5%	6.76	9.04	7.82	12.19	

Means with same letter are not significantly different at p d" 0.05.

Table. 3: Percentage of Inhibition of Pathogens by Volatile Compounds Produced by Trichoderma spp.

 Trichoderma isolates	P. oryzae	B. oryzae		R. solani
CRRI-TS-1	9.41 ^b	46.94 ^b	0.00 ^b	0.00 ^b
CRRI-TS-2	15.38 ^b	14.11 ^c	0.00 ^b	0.00 ^b
CRRI-TS-3	17.07 ^b	1.96 ^d	0.00 ^b	0.00 ^b
CRRI-TS-5	9.76 ^b	15.77°	0.00 ^b	0.00 ^b
CRRI-TS-6	15.85 ^b	15.03°	0.00 ^b	0.00 ^b
CRRI-TS-9	15.85 ^b	2.45 ^d	0.00 ^b	0.00 ^b
CRRI-TS-10	65.88ª	59.59ª	76.08ª	20.39ª
CV (%)	9.43	8.47	2.36	9.63
Tukey's HSD at 5%	12.73	5.26	0.72	1.43

Means with same letter are not significantly different at $p \le 0.05$.

species varied from 50.00-100.00%, 60.55-100.00%, 50.88-98.25% and 42.78-86.15% respectively (Table2).Complete inhibition of *Pyricularia oryzae* (CRRI-PO-2) was observed in confrontation to CRRI-TS-3, CRRI-TS-9 and CRRI-TS-10. CRRI-TS-6 inhibited both the radial growth of C.*oryzae-sativae* and *R. solani*, and it showed more than 80 % inhibition in case of all

the four pathogens. CRRI-TS-10 was found to be most effective with 100% inhibition to foliar pathogens *viz.Pyricularia oryzae* and *Bipolaris oryzae*.

Trichoderma is a genus of fungi that has dual role in plant growth promotion and biocontrol. Strong antagonistic potential of different *Trichoderma* Characterization of VOCs from *Trichoderma* spp.

[J.Mycopathol.Res :

Treatments Sahbhagidhan	Shoot length (cm)	Root length (cm)	Fresh root weight (g)	Fresh shoot weight (g)	Dry root weight (g)	Dry shoot weight (g)
T-0	5.94 ^b	4.90 ^b	0.027ª	0.065 ^{ab}	0.017ª	0.034ª
CRRI-TS-1	7.62 ^ª	5.15 ^b	0.032ª	0.072 ^ª	0.019 ^a	0.039 ^a
CRRI-TS-10	6.83 ^{ab}	5.61ª	0.034ª	0.062 ^b	0.016ª	0.023ª
CV (%)	7.87	2.45	9.17	4.63	8.66	9.01
Tukey's HSD at	1.34	0.32	0.008	0.007	0.003	0.017
5%						
Karuna						
Т-0	5.85ª	3.83ª	0.023 ^b	0.049 ^b	0.014ª	0.032 ^{ab}
CRRI-TS-1	5.57ª	4.31ª	0.028ª	0.061ª	0.015ª	0.040ª
CRRI-TS-10	5.88ª	5.06ª	0.029ª	0.061ª	0.016ª	0.025 ^b
CV (%)	6.38	9.56	3.32	6.99	8.55	9.29
Tukey's HSD at 5%	0.92	1.49	0.002	0.01	0.008	0.009

Table. 4: Effect of VOCs on growth promotion of Sahbhagidhan & Karuna seedlings

Means with same letter are not significantly different at p d" 0.05.

species has been reported against *Fusarium* oxysporum and *Rhizoctonia solani* (Kamala and Devi, 2012;Sundaramoorthy and Balabaskar, 2013), *Sclerotinia* (Garcia-Nunez *et al.* 2012), *Sclerotium delphinii* (Mukherjee *et al.* 2013), *Pyricularia oryzae* (Aravindan *et al.* 2016) and *Bipolaris oryzae* (Khalili *et al.*2012). This antifungal activity is known to be exhibited through several mechanisms such as antibiosis, utilization of available space, production of cell wall degrading enzymes, mycoparasitism, induction of systemic resistance (Harman *et al.* 2004; Saba *et al.* 2012; Mukherjee *et al.* 2022).

Assay of Trichoderma volatiles

The VOCs from *Trichoderma* isolates inhibited radial growth of the test pathogens *Pyricularia oryzae and Bipolaris oryzae* (Table3). CRRI-TS-10 was the best candidate as it showed growth inhibition all the pathogens. Out of the seven isolates, only CRRI-TS-10 was able to inhibit *R. solani*. Growth of *Bipolaris oryzae* was markedly inhibited by CRRI-TS-1(Fig 6).

Assay of volatiles as plant growth promoters

In the effect of volatiles emitted from CRRI-TS-10,Sahbhagidhan seedlings showed better shoot and root growth as compared to control (Table4), whereas, CRRI-TS-1 VOCs had positive effect on shoot length.In case of Karuna, seedlings, shoot and root biomass was increased significantly with CRRI-TS-1and CRRI-TS-10 VOCs treatment.

Volatile collection and characterisation

As confirmed from the confrontation assays,CRRI-TS-10 caused maximum inhibition of pathogen growth followed by CRRI-TS-1which was most effective against *Bipolaris oryzae*. So, VOCs were collected from these two*Trichoderma* spp. and analysed using GC-MS (Figs.7 and 8). The compound, 6-pentyl- 2H-Pyran-2-one was constituted 84.10 % of the total area of VOCs produced by CRRI-TS-10.(Z)- Ocimene, 1,3-Octadiene, 1-Octen-3-ol, 3-Octanone, (Z)-Ocimene & 3-(hydroxymethyl)-2-nonanone, were detected in both the isolates(Tables 5, 6).

Production of volatile compounds is one of the mechanisms executed by *Trichoderma* species for growth promotion and antagonism. So, the isolates were tested for the effect of volatiles emitted from them.CRRI-TS-1 and CRRI-TS-10being the best pathogen inhibitors were further analysed for VOCs composition by GC-MS technology. 6-Pentyl-2H-pyran-2-one was found to be the most abundant compound (84.10%) from CRRI-TS-10 i.e., *Trichoderma erinaceum* which is already reported to have strong

 Table. 5: HS-SPME/gas chromatography-mass spectrometry (GC-MS) profile of volatile organic compounds (VOCs) produced by

 Trichoderma pleuroticola

RT	Compound Name	Calculated RI	Literature RI	RSI	Area (%)	Activity & Reference
2.39	1-Propanol, 3-amino	739.2	761	854	0.07	-
6.16	1,3-Octadiene	825	820	837	0.61	
8.08	Heptan-2-one	868.3	898	909	0.07	Metabolite in many plants & E. coli (Pubchem), <i>T. viride</i> (Carballo <i>et al.,</i> 2017)
10.71	1-Octen-3-ol	1028.3	960	907	3.24	Antimicrobial in <i>Bignonia</i> venusta(L.) (Xiong et al., 2017) & <i>T.</i> atroviride (Sharma et al. 2017)
10.92	3-Octanone	1033.2	963	894	15.83	Antibacterial and antifungal in <i>Atta</i> ant (de Lima Mendonça <i>et al.,</i> 2009),
12.24	Ocimene <(Z)-, beta- >	1063.2	1035	775	0.19	<i>T. atroviride</i> (Sharma <i>et al.</i> , 2017) Antifungal in <i>Cnidiummonnieri</i> (L.) (Sun <i>et al.</i> , 2020) Antimicrobial in Cleopatra mandarin (<i>Citrus reshni</i>) (Hamdan <i>et al.</i> , 2013)
14.18	2-Nonanone,3 - (hydroxymethyl)-	1106	1093	882	0.10	-
17.34	Dodecane	1173	1200	880	0.20	3-(Prop-2-enoyloxy)dodecane has antibiotic properties in mangrove <i>Glutamicibacter mysorens</i> (Karthik <i>et al.</i> 2023)
19.87	Cyclohexanol,2- methyl-5-(1- methylethyl)- (1g.2g.58)-	1227	1219	783	0.11	-
22.77	Tridecane <n-></n->	1289.6	1300	822	0.10	-
23.54	Isogermacrene D	1306.3	1447	918	0.87	Found in roots of <i>Leonurus sibiricus</i> L. (Sitarek <i>et al.,</i> 2017)
25.84	Alaskene <beta-></beta->	1357.2	1494	864	0.25	-
28.87	Cuprenene <gamma></gamma>	1425	1530	900	0.17	-
29.24	Anthracene, tetradecahydro-	1434	1561	871	0.38	Antimicrobial (Kim <i>et al.</i> , 2009)
30.44	4-phenyI-2-butanol, TBDMS derivatives	1462	1601	858	0.04	-

RT: Retention time RI: Retention index, RSI: Reverse similarity index

antimicrobial action in *Trichoderma koningii* (Ismaieland Ali, 2017) and *Trichoderma atroviride* (Jin *et al.*2020). Both strains, CRRI-TS-1 and CRRI-TS-10 produced 3-octanone and 1-octen-3-ol, which are fungicides reported from *T. atroviride* (de Lima Mendonça *et al.*2009;Xiong *et al.* 2017; Sharma *et al.,* 2017). (Z)- Ocimene is also a common volatile antifungal compound (Hamdan *et al.*2013, Sun *et al.* 2020) in both the species which does not have any previous report from *Trichoderma*.Tetradecahydro-anthracene, from CRRI-TS-1; Propanoic acid, Furan, 2-pentyl, Acetophenone, Benzoic acid, methyl ester,Furan <2-heptyl->&Neomethyl acetate from CRRI-TS-10 are some antimicrobial agents that were found in this study (Wang *et al.* 2011;Saharkhiz *et al.*2012; Ma *et al.* 2013; Wu *et al.* 2015; Sosa *et al.* 2016; Dias *et al.* 2017; Parra Amin *et al.* 2021). These compounds have not been earlier reported from *Trichoderma* to the best of our knowledge.

Field Experiment for evaluation of growth parameters

Trichoderma conidial suspension treatments resulted tremendous changes in both vegetative and reproductive growth of the rice cultivar Sahbhagidhan. The yield parameters like panicle length, no of panicles per square meter, 1000 grain weight and percentage fertility were also

Characterization of VOCs from Trichoderma spp.

[J.Mycopathol.Res :

Table 6: HS-SPME/gas	chromatography-mass	spectrometry	(GC-MS) p	profile of	volatile	organic	compounds	(VOCs)	produced	by
Trichoderma erinaceum										

RT	Compound Name	Calculated RI	Literature RI	RSI	Area (%)	Activity and References
2.65	Propanoic acid	745	739	856	0.02	Antimicrobial in <i>Euphorbia</i> <i>lathyrus</i> (Sosa <i>et al.</i> , 2016)
4.18 6.16 8.06	Butane, 2-cyclopropyl- 1,3-Octadiene 2-Hexanone, 5-methyl-	779 825 868	811 820 838	880 861 817	0.02 0.22 0.04	- - -
8.74	Acetate <pentyl-></pentyl->	883	915	815	0.05	-
10.69	1-Octen-3-ol	1027	960	892	1.01	Antimicrobial in <i>Bignonia</i> venusta(L.) (Xiong et al., 2017) & T. atroviride (Sharma et al., 2017)
10.92	3-Octanone	1033	963	855	0.85	antibacterial and antifungal in <i>Atta</i> ant(de Lima Mendonça <i>et al.</i> , 2009), <i>T. atroviride</i> (Sharma <i>et al.,</i> 2017)
11.07	Furan, 2-pentyl-	1036	977	912	1.64	Antifungal in <i>Streptomyces</i> <i>spp.</i> (Wu <i>et al.,</i> 2015, Dias <i>et al.</i> 2017)
11.57	Propanoic acid, pentyl ester	1048	988	843	0.12	-
12.23	Ocimene<(Z)-beta>	1062	1035	875	0.03	Antifungal in <i>Cnidium monnieri</i> (L.) (Sun <i>et al.</i> , 2020) Antimicrobial in Cleopatra mandarin (<i>Citrus reshni</i>) (Hamdan <i>et al.</i> 2013)
13.39	Acetophenone	1089	1068	855	0.02	Acetophenone derivatives are antifungal (Ma_et al., 2013)
14.17	2-Nonanone, 3 -	1106	1093	897	0.01	-
14.25	Benzoic acid, methyl ester	1108	1094	900	0.06	Benzoic acid derivative - Antifungal in <i>Piper</i> <i>cumanense</i> (Parra Amin <i>et</i> <i>al.</i> 2021)
17.14	Furan <2-heptyl->	1169	1193	891	0.06	Antiphytopathogenic in <i>Pseudomonas spp.</i> (Wang <i>et</i> <i>al.</i> 2011)
18.88	Neomethyl acetate	1206	1272	816	0.04	Antifungal in <i>Mentha piperita</i>
19.47	Acetophenone, 2-	1219	1283	884	0.11	-
23.77	Bergamotene <alpha -,<="" td=""><td>1311</td><td>1416</td><td>945</td><td>0.35</td><td>-</td></alpha>	1311	1416	945	0.35	-
24.34	2H-Pyran-2-one, 6 - pentyl-	1324	1425	925	84.10	Antimicrobial in <i>Trichoderma</i> <i>koningii</i> (Ismaiel& Ali, 2017), <i>Trichoderma atrovirid</i> e (Jin <i>et</i> <i>al.</i> , 2020)
26.02	Menth-1 <i>-</i> en <i>-</i> 9 <i>-</i> ol acetate <para></para>	1361	1426	823	1.14	-

RT: Retention time RI: Retention index, RSI: Reverse similarity index

increased considerably. In comparison to untreated ones. CRRI-TS-9 treated plants yielded highest with 836.64g/m² followed by CRRI-TS-6 (757.5 g/m², CRRI-TS-3 (744.04 g/m²) and CRRI-TS-10 (738.48 g/m² (Table7).

Plant growth promotion activity of this genus has been reported in many plants, such as rice (Swain *et al.*2018), sunflower (Nagaraju *et al.* 2012), sugarcane (Srivastava *et al.* 2006), tomato (Bharti *et al.* 2012) etc. Various physiological activities contribute towards growth promotion. These are production of phytohormones, such as auxin (Contreras-Cornejo *et al.* 2009), modification of factors affecting nutrient uptake, such as mineral solubilization (Kapriand Tewari, 2010) and protection of plants against phytopathogens (Khalili *et al.*2012; Kamala and Devi, 2012; Garcia-Nunez *et al.*2012). The *Trichoderma* isolates which have been characterized in our

Table 7: Effect of Trichoderma treatments on vegetative growth of rice cultivar Sahbhagidhan

Shanti Prava Behera and others

	Shoot	Root	Fresh	Dry	Fresh root	Drv root		No of	Panicle	Grain	Grain	1000	
Treatmant	anoth	lanoth	shoot	shoot	waicht	wainht		aloidad	lanath	wainht	wainht nar	grain	Fertility
ון פמווום וו			weight	weight			- 5		India			weight	%
	(cm)	(cm)	2	: D	(a)	(a)		per m ²	(cm)	Per hill (g)	plot (g)	- E	2
	,		(6)	(6)								(B)	
Т-0	57.30℃	13.44 °	12.73₫	3.87 ^e	3.77 d	0.73℃	75.75ª	416.50 ^a	18.98	12.96 ₫	330.01 °	21.96ª	52.09ª
CRRI -TS-1	77.98 ^{ab}	19.57ª	37.14ª	11.59 ^b	5.38 °	1.47ª	72.00ª	575.75 ^a	23.69 ª	20.04 bc	699.81 ^b	23.14ª	58.73ª
CRRI -TS-2	4 6 2.07	15.83 ^{bc}	21.83°	7.34 ^d	6.97 ^{ab}	1.45 ^{ab}	72.50ª	553.70 ª	26.07 ^b	23.11 ^{ab}	709.50 ^{ab}	22.78ª	65.20 ^ª
CRRI -TS-3	72.88 ^{ab}	16.64 ^{abc}	20.23°	5.23 ^e	5.43°	1.37 ^{ab}	74.25ª	485.10 ^a	26.02 ^b	18.33 °	744.04 ^{ab}	22.84ª	61.23ª
CRRI -TS-5	76.38 ^{ab}	19.00 ^{ab}	30.89 ^b	6.75 ^d	4.95 cd	0.79°	73.50ª	619.85 ^a	24.75°	19.61 ^{bc}	695.84 ^b	22.00ª	61.20ª
CRRI -TS-6	79.84ª	17.33 ^{ab}	32.22 ^{ab}	13.15 ^a	8.11 ^a	1.57ª	74.75ª	600.25 ^a	23.98 ^{cd}	22.97 ^{ab}	757.50 ^{ab}	23.80ª	59.01 ª
CRRI -TS-9	76.63 ^{ab}	18.76 ^{ab}	35.34 ^{ab}	8.98°	6.20 ^{bc}	1.25 ^{ab}	71.00ª	600.25 ^a	27.35ª	23.77 ª	836.64 ª	21.46ª	59.23 ª
CRRI -TS-10	72.66 ^{ab}	16.93 ^{ab}	32.48 ^{ab}	7.88 ^{cd}	6.04 ^{bc}	1.04 ^{bc}	73.50ª	612.50 ^ª	24.77°	22.03 ^{ab}	738.48 ^{ab}	23.70 ª	60.52 ^a
CV (%)	4.60	8.17	8.22	7.93	9.52	9.72	3.70	9.35	1.64	7.75	8.31	6.045	9.68
Tukey's HSD at	7.86	3.28	5.35	1.50	1.4	0.41	6.36	252.84	0.94	3.69	134.0	3.21	14.92
5%													

Means with same letter are not significantly different at $p \le 0.05$.



Fig.1: Morphological growth patterns of isolates(5 days after inoculation on PDA)



Fig. 2: Conidiophore structure bearing conidia (indicated by arrow mark) as observed under light microscope

L	CRRI-TS1	CRRI-TS2	CRRI-TS3	CRRI-TS5	CRRI-TS6	CRRI-TS9	CRRI-TS10
-			-	-			-
100	_	_	_				

Fig. 3: ITS amplification of genomic DNA of Trichoderma spp.

Shanti Prava Behera and others

Treatment Name	Catalas unit/r	e activity (in nin/g)	Peroxida (in unit	ase activity :/min/g)	Superoxide Dist unit/n	mutase activity (in nin/g)
	Root	Shoot	Root	Shoot	Root	Shoot
T-0	4.00°	5.00 ^h	0.27 ^d	0.28 ^d	0.82 ^f	1.91 ^f
CRRI-TS-1	6.00ª	18.00 ^b	0.48 ^b	0.65ª	7.52ª	4.60 ^d
CRRI-TS-2	6.00ª	12.00 ^e	0.33 ^{cd}	0.36 ^{cd}	2.83 ^d	3.98°
CRRI-TS-3	5.00 ^b	6.00 ^g	0.32 ^{cd}	0.39 ^{bc}	1.83 ^e	5.03°
CRRI-TS-5	5.00 ^b	10.00 ^f	0.40 ^{bc}	0.30 ^d	4.82 ^b	10.03ª
CRRI-TS-6	6.00ª	13.00 ^d	0.35 ^{cd}	0.47 ^b	2.67 ^d	4.72 ^{cd}
CRRI-TS-9	5.00 ^b	15.00°	0.25 ^d	0.42 ^{bc}	2.87 ^d	4.94 ^{cd}
CRRI-TS-10	6.00ª	33.00ª	0.77ª	0.65ª	3.72°	9.02 ^b
CV (%)	1.02	3.12	8.74	6.95	5.46	2.65
Tukey's HSD at	0.55	0.23	0.097	0.08	0.52	0.41
5%						

Table. 8: Expression of stress related enzyme activities in the rice cultivar Sahbhagidhan

Means with same letter are not significantly different at $p \le 0.05$.





study exhibited growth promotion in rice in terms of enhanced biomass and yield. Positive effect on rice plant growth by *Trichoderma* species has been reported by Doni *et al.* (2014), Swain *et al.* (2018), Woo *et al.*(2023) and many others.

Defence enzyme activity

Experimental results revealed that the *Trichoderma* seed dressing could activate defence enzymes in rice plants. Catalase activity in Sahbhagidhan shoots was maximum increased by seed treatment with CRRI-TS-10 followed by CRRI-TS-1. Similarly, Both CRRI-TS-1 & CRRI-TS-10 were best peroxidase activity enhancers in shoot and root. In case of SOD activity in Sahbhagidhan roots, CRRI-TS-1 showed highest potential followed by CRRI-TS-5 and CRRI-TS-



Fig. 5: Confrontation assay showing inhibition of pathogen by Trichoderma spp.



Fig.6 : Inhibition of Pathogens by Volatile Compounds Produced by Trichoderma spp.



Fig. 7: Gas chromatography-mass spectroscopy profiling of volatile compounds from Trichoderma pleuroticola



Fig. 8: Gas chromatography-mass spectroscopy profiling of volatile compounds from Trichoderma erinaceum

10. But in shoots, CRRI-TS-5 & CRRI-TS-10 performed significantly well (Table 8).

CONCLUSION

All the seven Trichoderma isolates were effective against four plant pathogens. However, the VOCs emitted by these Trichoderma isolates were not equally effective against plant pathogens. Among the isolates, the VOCs emitted by Trichoderma erinaceum had antagonistic potential against all four pathogens, whereas VOCs released by other isolates were not effective. The compound, 6-Pentyl-2H-pyran-2-one released by Trichoderma erinaceum is already reported to have both biocontrol and growth promotion action. This isolate improved the growth of rice plants under field conditions. Trichoderma erinaceum could be promoted for use in rice disease management as an effective, environment-friendly benign tool. The VOCs emitted by the effective Trichoderma spp. can be utilized for the proper management of paddy health.

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