Potential of Bacterial species as biocontrol agents of Fusarium wilt disease

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Abstract

In this work aimed to evaluate the efficiency of biological products based on Bacterial species in the control of *Fusarium* species under in vitro conditions. Six rhizospheric bacterial isolates obtained from various field of cotton crop were screened *in vitro* by dual plate assay for antagonistic activity against Fusarium wilt pathogens. Their ability to B4 was found to be the most potential antagonist with a mycelial inhibition percentage of against FO3 and FO1 mycelia followed by FO4 and FO2 respectively. For the secondary metabolites screening in liquid media, 6 antagonistic bacteria against *Fusarium* sp. were selected with broad spectrum of PGPR. The result of the ethyl acetate extracted product of the bacterial isolates namely B3 and B4, against the two pathogen F3 and F1 are as given the NIST and WILEY library based on their retention time, molecular weight, molecular formula and one chemical structure of the compounds are illustrated by GCMS analysis.

INTRODUCTION

7.4 million hectares, making it a significant commercial crop. India holds the top spot in terms of acreage, accounting for about 25% of the world's cotton crop. In India, cotton has a prominent place among all cash crops. About 65% of the raw materials used by the Indian textile industry are cotton, making it a vital raw material. With approximately 1500 mills, 4 million handlooms, 1.7 million power looms, and hundreds of garment, hosiery, and processing facilities, the Indian textile sector plays a vital role in the nation's economy and employs over 35 million people directly or indirectly (Sankaranarayanan et al., 2011). Fusarium species is a soil-borne fungal disease that affects plants through their roots at all stages of growth, causing significant economic losses by generating necrosis and wilting symptoms in many agricultural plants and having a significant overall influence on production. The disease caused by this fungus is characterized by wilted plants, yellowed leaves and root rot minimal or absent crop yield. Fusarium sp. found in its many pathogenic forms, is the most damaging species of the genus where in plants are concerned. A number of new disease reports on Fusarium have been submitted to the literature pool on agricultural research Bokhari anajat and perveen kahkashan, (2012).

India's 4 million farmers cultivate the cotton crop over an area of

It is deemed safe to employ *Bacillus amyloliquefaciens*, *Bacillus subtilis*, and *Paenibacillus polymyxa*, which are often found in soils, in the environment and with animals (Zhao *et al.*, 2015).

They generate antagonistic actions directed against many bacterial and fungal diseases (Aarab *et al.*, 2015) Thus, the primary root colonisers are Pseudomonas and Bacillus strains, which are examples of PGPR (Manikandan *et al.*, 2010) can be applied to the control of illnesses caused by dirt. In this study, we tested selected *Trichoderma* species for their biocontrol ability against the Fusarium wilt of cotton under in vitro conditions.

MATERIALS AND METHODS

Isolation of pathogenic fungi

Evaluation of infected parts of the cotton plant resulted in isolation and identification of *Fusarium* species based on the examination under microscope. Parts of plants with symptoms of Fusarium wilt infection were surface sterilised by immersion in 0.3% sodium hypochlorite for 10 minutes, and then in 70% ethanol and later rinsed thoroughly with sterile distilled water. They were transferred to potato dextrose agar (PDA) medium in petriplates and incubated at $26 \pm 20 \mathrm{C}$ for seven days (Aneja, 2001). The characteristic growth of the fungus with morphological characters of micro-conidia and macro-conidia and chlamydospores were observed by (Agrios, 2005). Pure cultures were maintained on PDA slants and stored at 4oC in the refrigerator.

Isolation of antagonistic bacteria

Antagonistic bacteria were isolated from the rhizosphere soil sampled during the survey. One gramme of rhizosphere soil underwent repeated dilution treatment following the thorough

mixing of the soil with the root pieces (Aneja, 2001). Biochemical characterizations were performed to identify bacterial isolates using conventional methods. Cell and colony morphology and motility, Gram reaction and catalase and oxidase activities were determined by Allen et al., (2002) and Ogutcu et al., (2020). Tricalcium phosphate-containing Pikovskaya (PVK) medium was used to test for phosphate solubilization (Nautiyal, 1999). Schwyn and Neilands' (1987) description of the chromeazurol (CAS) agar assay medium was used to screen isolates for siderophore production. Using the dual culture method, the antagonistic activity of biocontrol drugs against F. oxysporum f. sp. vasinfectum was evaluated (Dennis and Webster, 1971). GC-MS was used to examine the chemical components found in nine isolates (B3) as (Verma et al., 2015).

STATISTICAL ANALYSIS

The experimental design was Completely Randomized Design (CRD) with three replicates as described by Gomez and Gomez, (1984). Test of variance was calculated using Analysis of variance (ANOVA). Differences among treatment means for each measured parameter were further separated using fishers Least Significance Difference (LSD) to determine levels of significance according to Cochran and Cox, (1992).

RESULTS AND DISCUSSION

Table - 1. Biochemical characterization of Bacterial species

The colonies formed by the bacterial isolates on the Yeast Extract Mannitol Agar (YEMA) were found to be translucent as or opaque and the surface texture were smooth, circular in shape and having entire margin. The outcomes of the biochemical tests and gramme reaction that were carried out to identify the isolates of Bacterial species shown that the findings of the gramme staining for each isolate were identical and urease test was negative, indole, catalase test and utilization of citrate was positive. Two isolates such as B2 and B5 were appeared Oxidase as negative. The detail characteristics are given table 1. Gram-positive, rod-shaped, spore-producing Bacillus mycoides bacteria develop into certain rhizoids, which produce cells that are arranged in filamentous bundles and extend end to end (Yi, 2018). According to Public Health England (2018), Bacillus species are Gram-positive rods that are frequently paired or chain cells with round or square ends. They typically have a single endospore, which is typically oval or occasionally spherical and extremely resistant to harsh environments.

The Gram-positive bacteria retain the crystal violet stain and appear purple when observed under a microscope. Meanwhile, the Gram-positive cell wall appears as a broad, 20-80 nm thick dense wall consisting of multiple interconnected layers of 60 to 90% peptidoglycan (Kaiser 2021).

S.No	Biochemical Test	В1	В2	В3	В4	В5	В6
1.	Cell Shape	Rod	Cocci	Cocci	Cocci	Rod	Cocci
2.	Gram Stain	-	-	-	-	-	-
3.	Indole test	+	+	+	+	+	+
4.	Urease	-	-	-	-	-	-
5.	Catalase	+	+	+	+	+	+
6.	Oxidase	+	-	+	+	-	+
7.	Citrate	+	+	+	+	+	+
8.	Hydrogen pro duction	+	+	+	+	+	+

The six isolates of bacterial species B1, B2, B3, B4, B5 and B6 were showed potential in biocontrol activity such as antagonistic activity against *Fusarium* wilt pathogens as well as PGPR activity (Table 2). The chitinase activities and surface area of production was more in B2, followed by B4, B1 and B5 isolate, although, B3 and B6 don't appear in chitinase activity. All isolates of bacterial species produced yellow pigmentation in blue coloured succinate medium. Further, When tetrazolium salt was added to a siderophore sample, certain bacteria species observed an immediate appearance of yellow colour, whereas B1 isolates showed a delayed emergence of yellow

colour, suggesting that the siderophore is hydroxamate type. Ghazy and El-Nahrawy, (2021) Using the CAS double-layer plate approach, two strains with strong siderophore production were tested. The strains may successfully prevent the hypha development of maize blight, as demonstrated by the results. Rouag et al., (2019) conducted a six possible the capacity of Bacillus isolates to stimulate plant growth against Foc. The results be seen that all of the isolate generated IAA, four of them also produced chitinases and cellulases, and one isolate even solubilized phosphates. Aarab et al., (2015).

Table - 2. Activities of rhizobacteria that promote plant development by the selected potential bacterial isolates.

S.No Isolates PGPR Attributes

		Colour of siderophore pigments	Surface area (mm²)	Chitinase activity (U/ml)
1	B1	Yellow	79.29	0.53
2	В2	Yellow	84.26	0.62
3	В3	Yellow	82.80	-
4	В4	Yellow	80.78	0.57
5	В5	Yellow	78.19	0.56
6	В6	Yellow	73.72	-
Values -	moan CE (n -	2)		1

Values = mean \pm SE, (n = 3)

The nitrogenase activities of all isolates were conformed to acetylene reduction assay and acetylene reduction activities as shown in Fig. 1. Isolates B4 and B1 able to converts maximum atmospheric nitrogen into ammonium under microaerophilic conditions at low nitrogen levels through the action of nitrogenase and able to grow as a pellicle when grow in Nfb semisolid medium were found to be 215.54 n moles of C_2H_4 per mg^{-1} of cell protein h^{-1} for B4 and 180.2 n moles of C_2H_4 per mg^{-1} of cell protein h^{-1} for B1. In the same context, Isolates B2 able to converts minimum atmospheric nitrogen into

ammonium under microaerophilic conditions through the action of nitrogenase activities. Around the world, isolates of the bacteria Penibacillus sp. have been found in a range of habitats, animals, plants, soils, and people. Numerous Penibacillus species are known to exhibit PGP effects, such as phosphate solubilization (Wang et al., 2012) and nitrogen fixation (Liu et al., 2019). Therefore, Rhizobiales, Sphingomonadales, Rhodospirillales, Nitrosomonadales, and Frankiales take role in the nitrogen cycle Sun et al., (2014); Dun-Dun et al., (2011).

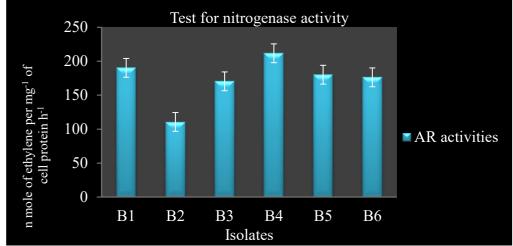


Fig. 1 Comparison of Nitrogenase activity of different isolates

Six rhizospheric bacterial isolates obtained from various field of cotton crop were screened in vitro by dual plate assay for antagonistic activity against *Fusarium* wilt pathogens for their ability to inhibit mycelial growth over control (Fig 2). Isolates, B4 and B3 was found to be the most potential antagonist with a mycelial inhibition percentage of 76.27 and 72.90 % against FO3 mycelia and 72.96 and 67.38 % against FO1 mycelia followed by FO4 (71.55 %) > FO2 (71.55 %) respectively. Among of all the bacterial isolates, isolate B2 was the least effective, reducing FO2 mycelial development by only 51.44 percent, the minimal inhibitory level. Therefore, the isolate B4 was most promising in controlling growth of all isolates of *F. oxysporum* f. sp. *vasinfectum* and B2 showed low ability to inhibit growth of the pathogen. *B. subtilis* is a somewhat effective biocontrol

formulation against club root pathogens (Zhu et al., 2019). However, reports on the antibacterial properties of Bacillus subtilis FA26 were assessed against *Clavibacter michiganesis* ssp. *Sepedonicus* (Cms), the causative agent of potato bacterial ring rot (Rajer et al., 2017).

Burkholderia cenocepacia demonstrated 44.4% inhibition in vitro and 86.1% field reduction in the incidence of FocR4T in Cavendish bananas (Ho et al., 2015). Conversely, Pseudomonas bacteria produce antibiotics, They have been identified as antagonistic agents against pathogens in suppressive soils, along with 2,4-diacetylphloroglucinol (2,4-DAPG), phenazine-1-carboxylic acid (PCA), pyrrolnitrin, rhizoxin, pyoluteorin, cyanide hydrogen (HCN) and 2-hexyl-5-propyl resorcinol (HPR) (Loper et al., 2012).

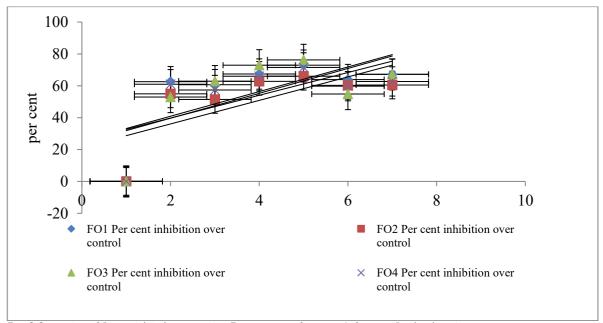


Fig. 2 Screening of Bacterial isolates against *F. oxysporum* f. sp. *vasinfectum* (Dual culture) Two antagonistic bacteria against *Fusarium oxysporum* with a metabolites screening broad spectrum of PGPR were chosen for the secondary (ethyl acetate) was used.

metabolites screening in liquid medium. An organic solvent (ethyl acetate) was used to extract the extracellular bioactive

compounds that the bacterial isolates in the culture filtrate generated. The outcome of the bacterial isolates B3 - Bacillus subtilis ethyl acetate extracted product against the two pathogens F3 and F1 are as given below (Table 3).

Thirty chemicals were found to be present in the ethyl acetate extract of *Bacillus subtilis* according to GC-MS analysis. By comparing the retention time, molecular formula, and molecular weight with the Wiley spectrum library search software, the identity of the chemical ingredients was verified. Fig. 2 display the active principles together with their retention time (RT), molecular weight, and molecular formula. Benzothiazole and 4-ditert-butylthiophenol were identified by Gao *et al.*, (2018) as the primary inhibitory volatile organic compounds generated by

B. subtilis CF-3. The postharvest fungus Colle and Monilinia fructicola were significantly inhibited in their growth. Furthermore, Three-methyl-2-pentanone, 2-heptanone, 2-octanone, 2-decanone, 5-methyl-2-hexanone, 2-nonanone, 2-dodecanone, 2-undecanone, and 5-methyl-2-heptanone and 2-pentanone were among the eight Bacillus strains that produced ketones, which are abundant sources of bioactive volatile organic compounds and were found to be effective in preventing the growth of Fusarium solani (Li et al., 2015) and Mycelial growth and spore germination were strongly inhibited by 2-methylenecyclohexanol, 2-nonanone, 2-nonanol, 2-(2-methylpropyl)-3-(1-methylethyl)pyrazine, and 2-dodecanone (Goswami and Kistler, 2004).

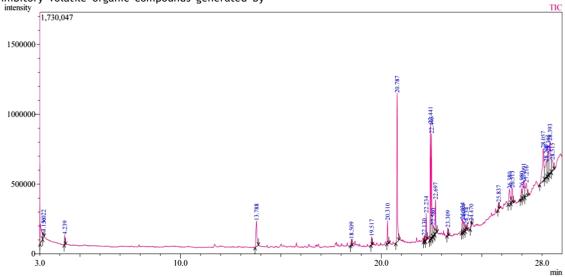


Fig. 2 GC-MS chromatogram of Bacillus subtilis

Table 3. Chemical compounds detected through GCMS

S. No.	R.Time	Area%	Height%	Name of the compound	Structure
1.	3.022	3.83	2.51	ethyl acetate	
2.	13.788	0.70	0.83	Pyrrolidine	
3.	18.509	0.58	1.25	Dibutyl ester	

CONCLUSION

most promising antagonists were determined to be isolates B4 and B3, with a mycelial inhibition percentage of 76.27 and 72.90 % against FO3 mycelia and 72.96 and 67.38 % against FO1 mycelia followed by FO4 (71.55 %) > FO2 (71.55 %) respectively, and their antagonists were the most effective in reducing the occurrence and severity of Fusarium oxysporum, an incidental pathogen that causes cotton wilt. There is strong evidence that microbial inoculants based formulations may be highly effective in suppressing soil borne diseases and enhancing crop productivity.

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