1	Original Research Article
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3	Isolation and characterization of plant growth promoting
4	rhizobacteria Enterobacter hormaechei and their
5	suppression efficacy against <i>Colletotrichum falcatum</i> in
6	combination with chitosan
7	ABSTRACT
8	Aims: This study aimed to explore the suppression efficacy of plant growth promoting rhizobacteria
9	(PGPR) Enterobacter hormaechei, chitosan and its oligomers either singly or in combination on red
10	rot disease causing pathogen Colletotrichum falcatum in sugarcane.
11	Methodology: The study was conducted to isolate twenty nine bacteria from sugarcane rhizosphere
12	and investigate their potential for plant growth activities. Selected isolate PSC3 was characterized by
13	biochemical and molecular identification by 16S rRNA sequencing. The study was further preceded
14	for in vitro screening of plant growth promoting traits viz., production of Indole-3-acetic acid (IAA),
15	hydrogen cyanide (HCN) ammonia (NH $_3$) production and antifungal activity against C. falcatum.
16	Results:
17	Among twenty nine isolates strain PSC3 showed highest plant growth promoting traits viz., indole-3-
18	acetic acid, hydrogen cyanide, ammonia production and antifungal activity against C. falcatum among
19	other isolated strains. Nucleotide 16S rRNA sequence analysis using clustalW program revealed that
20	isolate PSC3 showed phylogenetic affiliation and maximum homology (99%) with E. hormaechei.
21	Antifungal activity of chitosan, chitooligosaccharides (COS) and E. hormaechei were checked by
22	inhibition of C. falcatum mycelial radial growth. Among three treatments of chitosan, COS and
23	chitosan + E. hormaechei, two treatments showed significant antifungal activity (P<0.05). Chitosan
24	treatment showed radial growth range from 2.5±0.07 to 1.9±0.03 cm against C. falcatum in
25	comparison with control (9.1±0.09cm). The significant growth inhibition 79.0% was observed in
26	chitosan at concentration 0.6% but the combination of chitosan with PGPR E. hormaechei PSC3
27	showed highest growth suppression of C. falcatum (86.8%) whereas fungal treated with only E.
28	hormaechei showed growth radial inhibition 41.3%.

Conclusion: The findings reveal that chitosan and *E. hormaechei* have significant effect on *C. falcatum*. This new antifungal combination may be help to prevent red rot disease in sugarcane.

31 *Keywards:* Enterobacter hormaechei; Colletotrichum falcatum; Sugarcane; Chitosan,
 32 Chitooligosaccharides; Antifungal activity.

33 1. INTRODUCTION

Plant growth-promoting rhizobacteria (PGPR) are plant-associated microorganisms that are known to induce plant defenses and confer beneficial effects such as increased plant growth and low susceptibility to diseases caused by pathogens¹. Therefore, their use as biofertilizers or control agents for agriculture improvement has been a focus of numerous researchers². PGPR have been proven to counteract the activities of other harmful soil borne microorganisms, thus promoting plant growth³. Some PGPRs also elicit physical or chemical changes related to plant defense, a process called "induced systemic resistance" (ISR)⁴. ISR confers plant resistance against a large variety of attackers

41 such as pathogens and herbivores⁵.

42 The red rot caused by C. falcatum Went is the most ruinous disease of sugarcane and a big 43 menace to both cane growers and sugar industry⁶. Conventional control of disease depends on the 44 use of chemical inputs and resistant varieties. Development of new variants of the fungus, health 45 hazards and environmental pollution concerned with the excessive use of agro-chemicals have 46 resulted in adopting the biological control using native strains of PGPR as a supplemental approach 47 to minimize pesticide usage⁷. Certain strains of PGPR have been used as ingenious weapon to 48 protect plants from various soil borne pathogens. These bio-antagonists adopt single or multiple mechanisms of action to suppress these pathogens which include antibiosis⁸, production of iron 49 50 chelators, secretion of hydrolytic enzymes, synthesis of hydrogen cyanide thus disease control can be 51 obtained by applying bacterial cells or their metabolic products⁹.

52 Chitosan is derived from chitin, a polysaccharide found in exoskeleton of shellfish such as 53 shrimp, lobster or crabs and cell wall of fungi¹⁰. Chitosan, poly (1, 4)-2-amino-2-deoxy- β -D glucose is 54 a deacetylation product of chitin, a polysaccharide second by the prevalence in nature after 55 cellulose^{11,12}. It is a nontoxic, biodegradable biopolymer of high molecular weight. Recent studies on 56 chitosan have attracted interest for converting chitosan to oligosaccharides¹³. In this respect, chitosan 57 oligosaccharides, because of their shorter chain length, display a reduced viscosity and are soluble in 58 aqueous media at pH values close to neutrality, which increases their bioavailability and opens a wide

range of new potential applications¹⁴. Due to its properties, various studies shown that chitosan has
antifungal and antibacterial activities in different diseases ^{15, 16}.

In view of this, the focus of the work presented in this paper is directed towards isolation and identification of PGPR from sugarcane rhizosphere. Subsequently, *in vitro* screening of the potential antagonists that control red rot disease causing pathogen. Further, this research work proceeded to check antifungal activity of chitosan and their combination with *E. hormaechei*. Therefore such type of study is necessary as it advocates that use of PGPR as inoculants or biofertilizers association with chitosan is an efficient approach to replace fungicides.

67

2. MATERIALS AND METHODS

68 **2.1** Processing of soil samples for isolation of Phosphate solubilizing microrganisms

Phosphate solubilizing rhizobacteria (PSB) were isolated from sugarcane rhizospheric soil by dilution plate technique using Pikovskaya's medium¹⁷. Appropriate soil dilutions were plated on Pikovskaya's agar medium by spread plate technique and incubated at 30 ± 1 °C for 2-3 days. The colonies forming halo zone of clearance (Pikovskaya's medium) around them were counted as P-solubilizers.

73 **2.2 Morphological and Biochemical characterization**

The efficient PGPR were identified on the basis of morphological, physiological and biochemical characteristics according to the standard methods described in Bergey's manual of systematic bacteriology¹⁸ and laboratory manual of basic microbiology¹⁹.

77 2.3 Molecular characterization of efficient strains

78 Molecular characterization of most efficient bacterial isolates was done by sequencing of their 16S 79 rRNA gene. Bacteria PSC3 showed efficient plant growth promoting mechanism among the all other 80 strains. Molecular characterization of bacteria PSC3 has been completed after DNA isolation of 81 selected bacteria PSC3 followed by quantification of DNA sample; amplification of DNA by using 82 bacterial specific primers 27F (5'-AGAGTTTGATCCTGGCTCAG-3') and 1492 R (5'-83 TACGGTTACCTTGTTACGACT-3'); choosing the PCR product based on concentration and 84 processed for sequencing. Sequences have been submitted to NCBI GeneBank by Sequin.

85 **2.4 Detection of Indole-3-acetic acid (IAA) production**

Indole acetic acid production was quantitatively measured by the method given by Gordon and Weber
 (1951)²⁰. Bacterial cultures were grown in a nutrient broth amended with tryptophan (5mM) for 3-4

- days. Cultures were centrifuged at 10,000 rpm for 20 min. Two ml of supernatant was mixed with two
- 89 drops of orthophosphoric acid and 4 ml of Salkowski reagent. Pink colour indicates presence of IAA.

90 2.5 HCN production

91 All isolates were subjected for the production of hydrogen cyanide (HCN) by amending 4.4 g glycine/ I

- 92 media. Whatman No.1 filter paper was soaked in 2 % sodium carbonate and 0.5% picric acid solution
- 93 was placed in the upper lid of the plate. The plates were sealed with parafilm and incubated at 28 ±
- 94 30°C for 5 days. The formation of orange to red colour indicates the production of hydrogen cyanide²¹.
- 95 2.6 Siderophore assay
- 96 The isolates were screened for the siderophore production by adapting the universal methods
 97 explained by Schwyn and Neilands (1987)²².

98 2.7 Detection of ammonia production

- 99 Qualitative detection of ammonia production was done by the method given by Bakker and Schippers,
- 100 (1987)²³. Bacterial isolates were grown in peptone water for 2-3 days at optimum growth temperature.

101 After incubation, 1ml of Nessler's reagent was added in each tube. Tubes showing faint yellow color

- 102 indicated small amount of ammonia, and deep yellow to brownish color indicated maximum amount of
- 103 ammonia.

104 **2.8 Peptone dextrose agar media preparation:**

- 105 The experiment was conducted 20.0 g potato, 2.0 g dextrose, 2.0 g agar were mixed in 100 ml 106 distilled water in a conical flask and was make a air tight with the cotton plug and wrapped with silver 107 foil. And it is placed in the autoclave for 1 hour at 121^oC at 15 lbs. Subsequently, it was taken out from 108 the autoclave and allowed to cool for solidify down in the laminar air flow.
- **2.9 Preparation of chitooligosaccharides**
- 110 The enzymatic method was used for preparation of COS from chitosan²⁴.

111 **2.9.1** Immobilized papain preparation

112 Chitin flakes (1.0 gm) were suspended in phosphate buffer (20 ml, 0.1 M, pH 6.5) and added 5mM 113 cysteine; 2mM ethylene diamine tetraacetate (EDTA); 17.5 mg freeze-dried papain (EC3.4.22.2). The 114 solution was kept at 5°C for 15 min in refrigerator. Then, 5% glutaraldehyde (3.1 ml) was added and 115 the suspension was kept under mild stirring at 5°C for 14 h. The chitin–papain was filtered and 116 washed with the same phosphate buffer trice, then stored in distilled water^{24,14}.

117 **2.9.2** Activity determination method of papain for chitosan

- An immobilized papain (3 gm is equals to 28.5 mg papain) was added to chitosan solution (1%, 10 ml)
- prepared by acetate buffer (0.1 M, pH 4.0). The suspension was kept under mild stirring at 45^oC for 1
- 120 h. The viscosities of the solution before and after reaction were determined by viscometer at 20°C.
- 121 Under these conditions, the papain activities decreased the viscosity of substrate chitosan.

122 2.9.3 Determination of Chitosan and chitooligosaccharides content

123 The COS contents were determined by 3, 5-DNS colorimetry 25 .

124 2.9.4 UV-Vis Spectroscopy

UV-Vis spectra of chitosan derivatives are usually recorded in aqueous acid (acetic acid) solutions in a 1.0 cm quartz cell at ambient temperature. The Diffuse Reflectance UV-Visible (DRUV) spectra of powdered or film samples are measured. Analysis in the vacuum ultraviolet through the near-infrared range has also been applied.

129 **2.10 Effect of Chitosan on mycelia radial growth**

130 Antifungal activity was determined by a radial hyphal growth of C. falcatum. Mycelium Growth 131 Inhibition in vitro was performed on growth medium treated with 0.2%, 0.4%, 0.6% chitosan, COS 132 concentration and combination of E. hormaechei. After 48 hr of incubation, agar piece of uniform size 133 (diameter, 8 mm) containing fungi were simultaneously inoculated at the centre of each petri dish containing the various concentration of chitosan followed by incubation at $25 \pm 2^{\circ}$ C for 14 days. After 134 135 incubation of fungi on culture medium containing chitosan, radial growth of fungal mycelium was 136 recorded. Radial inhibition was calculated when growth of mycelia in the control plate reached the 137 edge of the petri dish. The fungicidal effect to growth of fungi, in terms of percentage inhibition of 138 mycelial growth was calculated by using the formula % inhibition = $dc - dt/dc \times 100$ Where dc =139 Average increase in mycelial growth in control, dt = Average increase in mycelial growth in 140 treatment²⁶.

141 Statistical Analysis & Preparation of Data

All the treatment data were statistically evaluated with SPSS/16.00 software. Hypothesis testing methods included one way Analysis of Variance (ANOVA) followed by LSD's test. P<0.05 was considered to indicate statistical significance. All the results were expressed as mean \pm S.E. for the 3 replicate in each treatment.

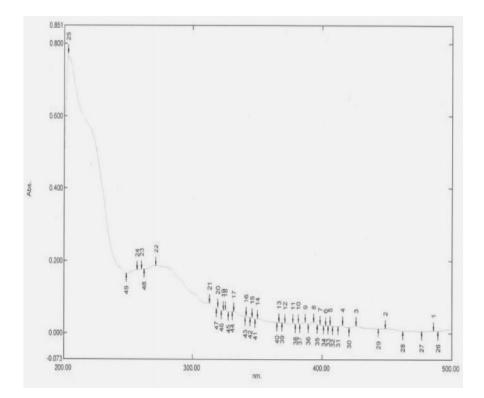
146 3. Results and Discussion

147 **3.1 Characterization of Chitooligosaccharides**

Chitosan treated with papain releases COS. COS were preliminary confirmed by 3, 5-DNS method and with formation of brown coloured complex with sugars. The results showed that the viscosity of COS decreased upto 51.47% of the beginnings chitosan solution. This was also confirmed presence of COS.

152 3.2 UV-Vis Spectrum

153 Structure of COS was confirmed by UV-vis spectroscopy. UV-vis spectrum was recorded on Perkin 154 Elmer Lambda 3B UV-vis spectrometer. Ultraviolet protection factor (UPF) was measured using UV 155 Shimadzu 3101 PC spectrophotometer. UV-Vis spectra of chitosan derivatives are usually recorded in aqueous acid solutions in a 1.0 cm quartz cell at ambient temperature⁸. The Diffuse Reflectance UV-156 Visible (DRUV) spectra of powdered or film samples are measured²⁷. Chitosan include various ratios 157 158 of two far-UV chromophoric groups, N- acetylglucosamine (GlcNAc) and glucosamine (GlcN); as a 159 result, their extinction coefficients for wavelengths shorter than approximately 225 nm are non-zero. 160 Because GlcNAc and GlcN residues show no evidence of interacting within the chitosan chain, the 161 monomer units contribute in a simple, additive way to the total absorbance of these polymers at a particular wavelength²⁸. The UV spectra of mixtures of N-acetyl-glucosamine and glucosamine 162 163 hydrochloride are guite similar to the spectra of chitosan, and the λ max is 201 nm in 0.1 M HCl 164 solution UV-vis absorbance spectra of chitosan exhibits characteristic peak at 230 nm. After 165 preparation of chitoligosaccharides, this peak undergoes a characteristic peak at range 360-348nm is 166 which observed (Fig.1).



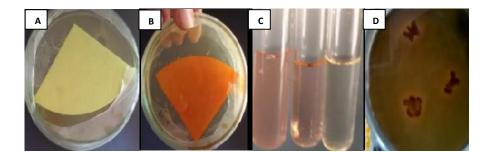
168 **Fig 1:** Characterization of chitooligosaccharides using UV- Vis spectroscopy

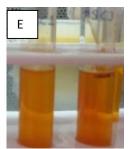
169 3.3 Isolation and Biochemical characterization of Isolates

170	The study was conducted to isolate PGPR from sugarcane rhizosphere and investigate their poter	ntial
171	for plant growth activities. Twenty nine PGPR were isolated by serial dilution in selective media fi	rom

- 172 two places of Uttar Pradesh. The study was further preceded for molecular identification of bacteria by
- 173 16S rRNA sequencing, and in vitro screening of plant growth promoting traits viz., production of
- 174 Indole-3-acetic acid (IAA), Hydrogen cyanide (HCN) Ammonia (NH₃) production and antifungal activity
- 175 against C. falcatum. Selected isolate was characterized by morphological, physiological and
- 176 biochemical method. For identification and decipher their phylogenetic affiliation with bacteria, isolate
- 177 was subjected to 16S rRNA (1492 bp long) gene sequencing. Nucleotide sequence analysis of test
- isolate using clustalW program revealed that isolate PSC3 showed maximum homology (99%) with
- 179 Enterobacter hormaechei.

- 180 *E. hormaechei* strain is gram-negative rods which are motile, catalase positive, and oxidase
- 181 negative and ferment D-glucose. The strain show negative Voges-Proskauer reactions. A detailed
- 182 biochemical profiling of the isolate is given in Table 1. Acid is produced from the compound D-sorbitol.
- 183





- 186
- 187 Fig 2: Screening of Plant growth promoting traits of E. hormaechei PSC3: (A and B) HCN production,
- 188 (C) IAA production (D) siderophore production and (E) NH₃ production.
- 189 Table 1: Biochemical characteristics of *Enterobacter hormaechei* strain isolated from sugarcane
- 190 rhizosheric soil

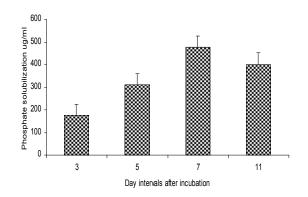
Biochemical test	E. hormaechei PSC3	Biochemical test	E. hormaechei PSC3
Colony shape	Irregular spreading	Methyl Red test	+
Colony colour	Yellow	Voges-Proskauer test	-
SIM (Motality)	+	Sucrose	-
Oxidase	-	D-Lactose	-
Catalase	+	Mannitol	-
TSI	R/R	D-Sorbitol	+
Citrate	+	Innositol	-
Nitrate	+	Maltose	-

Gelatin	+	Dextrose	-
Starch	-	Galactose	-

192 **3.4 Evaluation of isolates for their Plant growth promoting activities and physiological traits:**

The phosphorus solubilizing activity was evaluated (Fig 3). Phosphate solubilizing activity of bacterial isolates PSC3 showed the highest phosphate solubilizition efficiency 475.51 µg/ml at 7th day of intervals and lowest at 3rd day of intervals. *E. hormaechei* KU196780 was showing plant growth promoting activities like Indole-3-acetic acid production hydrogen cyanide production and ammonia production (Fig 2).

198 PGPR isolate PSC3 grew up to 250 mM and none grew at 300 mM of NaCl conditions, but 199 the concentrations of 50 to 100 mM NaCl were critical as the isolate showed discriminatory 200 performances in these NaCl concentrations. At 50 mM NaCl, isolate exhibited very luxuriant good 201 growth comparison with other concentration. At 250 mM and 300 mM NaCl, the isolates show very 202 less or no any growth respectively. PEG of 20, 40 and 60 % were found high for the growth of isolate 203 PSC3 (Table 2). The isolate PSC3 showed greatest growth at pH 7.0, 9.0. There was no any growth 204 on pH 5.0, 11.0. The finding showed that this strain might be help in drought and saline stress in 205 plants.



- 207 Fig.3: Quantification of Phosphate solubilization in g/ml by *E. hormaechei* PSC3 strains in different
- 208 day intervals (Data are expressed as mean ± SE, n=3)
- 209 **Table 2:** Plant growth promoting characteristics of *Enterobacter hormaechei* strain isolated from
- 210 sugarcane rhizospheric soil (+ Good, ++ Strong, +++ Very strong)

PGP traits	E. hormaechei	Stress tolerance	E. hormaechei PSC3

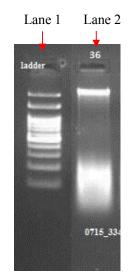
	PSC3	traits	
Phosphate solubilization	+	рН	+
NH ₃ Production	++	NaCl	+
HCN Production	+++	PEG	+
IAA Production	+++	Cu	-
Siderophore Production	+	Hg	-

212 3.5 Molecular identification of isolate PSC3

213 Molecular tools for the identification of soil bacteria were used and 16S rDNA gene analysis was 214 intensively used to understand the phylogenetic relationships. The accession numbers of the 16S 215 rDNA sequences is KU196780. Bacterial phylogenetic classification is based on sequence analysis of 216 the 16S rRNA molecule or its genes. For further identification at genus level, bacterial isolates were identified through homology search with BLAST and FASTA using partial sequence of 16S rDNA²⁹. 217 218 Sequencing data showed that the isolates belonged to genus, Enterobacter spp. being a dominant 219 species. Nucleotide sequence analysis of test isolates using clustalW program revealed that isolate 220 PSC3 showed maximum homology (99%) with Enterobacter hormaechei (Fig. 5).

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- Fig. 4 : Agarose gel electrophoresis of the 16S rDNA PCR products of bacterial isolate .Lane 1: 1kb
- 225 DNA ladder; Lane 2: bacterial isolate PSC3

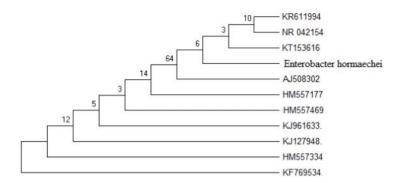


Fig 5: Neighbor-joining phylogenetic dendrogram based on a comparison of the 16S rRNA gene sequences of some of their closest phylogenetic taxa.

229 3.6 Growth Inhibition

230 Antifungal activity of chitosan, COS and in combination with E. hormaechei were evaluated based on 231 the diameters of growth inhibition percentage against C. falcatum. If there is no inhibition, it is 232 assumed that there is no antifungal activity. The validation of potential antifungal activity has been 233 validated against known organisms, such as C. falcatum. In vitro prescreening showed noticeable 234 antagonistic activity of isolate PSC3 against C. falcatum with a variable range of percentage inhibition. 235 Fig.7 shows representative radial growth plates with red rot causing fungus C. falcatum after 7 day 236 incubation. Initially we found the significant result of individual effect of chitosan, COS with 0.5 %, E. 237 hormaechei and antifungal drug clotrimazole after 10 days of incubation (Fig.6). In which E. 238 hormaechei showed 41.3 % growth inhibition of fungal pathogen C. falcatum. The radial growth 239 inhibition of C. falcatum is larger in 0.6 % chitosan than other concentration 0.2%, 0.4% chitosan. 240 These findings indicated that C. falcatum is more susceptible at the dose of 0.6%. But this 0.6 % 241 chitosan is showing highest antifungal activity when combined with E. hormaechei. Fig 8 also showed 242 highest growth inhibition in chitosan combination with E. hormaechei. Chitosan with various 243 concentrations demonstrated effective inhibition on the fungi growth (P < 0.05). The average mean of 244 radial growth is 1.87 to 9.13 cm against C. falcatum. Microscopic analysis of lactophenol blue stained 245 fungal mycelia showed structural aberration in chitosan treated C. falcatum (Fig.9). These findings 246 support the fact that, virtually, all the agricultural soils possess some suppressive effect on various soil 247 borne pathogens causing diseases in plants which may be because of the antagonistic activities of 248 microbes existing in soil. This phenomenon is also known as "general suppression" or "general 249 antagonism". This may be possible due to production of HCN. Michelsen and Stougaard, 2012

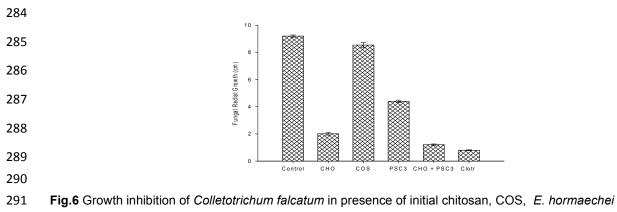
reported that HCN is a secondary metabolite produced by many antagonistic *Pseudomonas* species³⁰. He also found that production of HCN inhibited growth of hyphae of *Rhizoctonia solani* and *Pythium aphanidermatum*.

253 One way ANOVA analysis indicated significant difference among the treatments (F3, 254 11=2.298 P<0.05). The treatment T2 (Chitosan 0.2%), T3 (Chitosan 0.4%), T4 (Chitosan 0.6%), 255 showed significant result compare with T1 (Control) somewhat treatment T4 showed greatest 256 significant in this treatment. Which indicate 0.2% chitosan solution showing very efficient antifungal 257 result against C. falcatum. Chitosan at the rate of 0.6 % showed 79.0% growth inhibition of C. 258 falcatum (Fig.10). Also our study coincided with those of Meng et al., 2012 who demonstrated that 259 Chitosan and COS had stronger inhibitory effect on mycelia growth of two fungal pathogens A. 260 kikuchiana and P. piricola³¹. Numerous studies on antifungal activity of chitosan against plant pathogens have been carried out and reviewed³². Chitosan's inhibition was observed on different 261 262 development stages such as mycelial growth, sporulation, spore viability and germination, and the 263 production of fungal virulence factors. It has been commonly recognized that antifungal activity of 264 chitosan depends on its molecular weight, deacetylation degree, pH of chitosan solution and, of 265 course, the target organism. Mechanisms proposed for the antifungal activity of chitosan focused mainly on its effect on fungal cell wall³³ and cell membrane³⁴. 266

Analysis of variance was used to determine whether levels of significant with chitosan treated in *C. falcatum* fungal strain different among control. The analysis showed no significant difference among the treatment (F 3, 11= 3.89 P<0.05). The treatment T2 (COS 0.2%), T3 (COS 0.4%), T4 (COS 0.6%), showed significant result with other treatment somewhat treatment T2 showed lowest radial growth in this treatment which indicate 0.2% showing no efficient antifungal result against *C. falcatum fungal* (Fig.10).

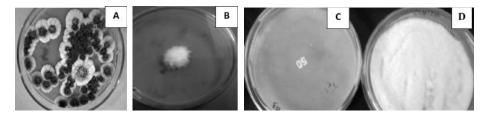
Chitosan and its derivatives offer a great potential as natural biodegradable nontoxic substances which have anti-microbial and eliciting activities. In the present study, the Chitosan was showing antifungal activity and highly effective in managing the red rot disease in sugarcane. The 0.6% of Chitosan was showing significant dose compared control. The Chitosan was showing more effect than COS. That indicate chitosan was efficient antifungal agent and highly effective in managing the complication associated with red rot disease. The 0.2% chitosan and COS was showing significant dose compared 0.2%, 0.4%., 0.6%. The study revealed that chitosan was

effective in inhibiting mycelial growth of *C. falcatum*. However, when compared to chitosan and *E. hormaechei* is relatively more effective than chitosan (Table 3). Furthermore, our results indicated that both chitosan and plant growth promoting rhizobacteria were effective in controlling diseases caused by *C. falcatum*.



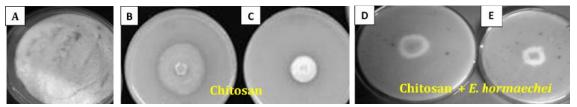
and their combination (Data are expressed as mean ± SE, n=3)

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Fig.7 Growth of *Colletotrichum falcatum* in presence of initial screening of different concentration of chitosan after 7 days. (A) 0.1% Chitosan (B) 0.5% Chitosan(C) 1.0% Chitosan (D) without chitosan control



298 299

Fig.8 Growth of *Colletotrichum falcatum* in presence of chitosan and chitosan with *E. hormaechei* and
 their combination after 7 days of inoculation (A) Control, (B) 0.2% Chitosan, (C) 0.6% Chitosan, (D)
 0.2% Chitosan with *E. hormaechei*, (E) 0.6% Chitosan with *E. hormaechei*

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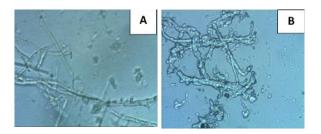
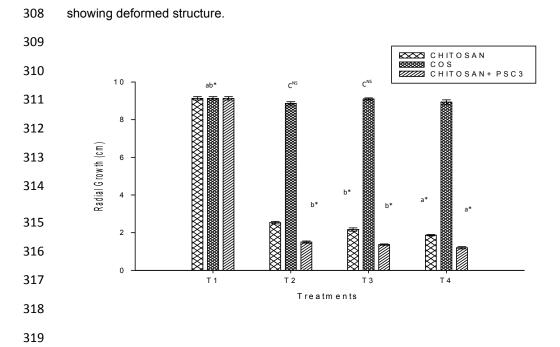


Fig.9 Microscopic analysis of lactophenol blue stain mycelia (a) Control (b) Chitosan treated mycelia



- 320 Fig.10 Antifungal Activity of different concentration of chitosan against Colletotrichum falcatum (Data
- 321 are expressed as mean ± SE, n=3).

- 322 Values are statistically significant at *p<.05. Significance determined by ANOVA was compared within
- the treatments as follows: a Control vs 0.6% chitosan and chitosan+ E. hormaechei; b 0.2%, 0.4%,
- 324 0.6% Chitosan vs. Control and ^{CNS} Not significant

325 **Table 3.** Percentage of radial growth inhibition of *Colletotrichum falcatum*

Concentration of /	Growth Inhibition	Growth Inhibition	Growth Inhibition % by
COS Chitosan	% by Chitosan	% by COS	Chitosan + <i>E.</i>
			hormaechei
0.2 % Chitosan/COS	71.3%	2.7%	83.5%
0.4% Chitosan/COS	76.1%	1.0%	84.9%
0.6% Chitosan/COS	79.0%	3.0%	86.8%

327 4. Conclusion and Future Prospects

This present research findings proved that this study is helpful for developing new biocontrol combination from chitosan and plant growth promoting rhizobacteria for managing fungal diseases and associated complications. The study reveals that chitosan solution and their concentration have significant effect of antifungal activity but their combination with plant growth promoting rhizobacteria *E. hormaechei* showed greatest growth inhibition of *C. falcatum* (86.8%). The chitosan and *E. hormaechei* seems promising for the development of a new formulation for fungal infection in plants.

A further investigation of the best antifungal result of chitosan and *E. hormaechei* like time of application, concentration, combination with other components, physiological changes in plants and molecular mechanism are needed and provide future line of work for controlling red rot disease of sugarcane for sustainable agriculture.

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