ORIGINAL ARTICLE

Bioflm and Biocontrol Modulation of *Paenibacillus* **sp. CCB36 by Supplementation with Zinc Oxide Nanoparticles and Chitosan Nanoparticles**

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Abstract

Endophytic bacteria with multi-trait plant benefcial features have applications to enhance agricultural productivity by supporting the plant growth, yield, and disease resistance. In this study, *Paenibacillus* sp. CCB36 was isolated from the rhizome of *Curcuma caesia* Roxb., and its bioflm formation and antifungal properties have been evaluated in the presence of nanoparticles. Chitosan nanoparticles (CNPs) were synthesized and characterized by UV–visible spectrophotometry, Fourier transform infrared (FTIR) spectroscopy, high-resolution-transmission electron microscopic (HR-TEM) analysis, scanning electron microscopic (SEM) analysis, and dynamic light scattering (DLS). The efect of zinc oxide nanoparticles (ZnONPs) and CNPs on bioflm formation of *Paenibacillus* sp. CCB36 was evaluated by tissue culture plate assay. ZnONPs reduced its bioflm formation and was found to get modulated in the presence of CNPs as revealed by atomic force microscopy (AFM). Hence, CNPs were selected for further studies. Interestingly, biocontrol property of *Paenibacillus* sp. CCB36 against *Rhizoctonia solani* was also found to get enhanced when supplemented with chitosan nanoparticles. The results of the study indicate application of nanoparticles to improve colonization and active functioning of endophytic bacteria which can have significant application in agriculture.

Keywords Nanoparticles · Bioflm · Biocontrol · Resistance

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Introduction

Nanotechnology has the potential to positively infuence all the sectors including agriculture, food, and pharmaceutical industries and has also been utilized to reduce the harmful efects of chemical inputs on environment and human health [\[11,](#page-13-0) [21,](#page-14-0) [28\]](#page-14-1). In agriculture, the unique physicochemical properties of nanomaterials such as catalytic reactivity, high surface area, size, and shape make them to have promising applications $[23]$ $[23]$ $[23]$. Nanotechnology has been explored very much in modern agriculture to develop sustainable methods to enhance the crop yield and also to control plant diseases [[21](#page-14-0)]. Nanoparticles can have promising role in the control of phytopathogens and pests due to its enhanced solubility, specifcity, permeability, and stability [\[32\]](#page-14-3). Biofertilizers and biopesticides formulated with nanosized particles have already been described to enhance its performance in an environment friendly manner [[4](#page-13-1)].

Plant growth-promoting microorganisms have been widely used to enhance plant productivity and disease management [[5,](#page-13-2) [10](#page-13-3), [26\]](#page-14-4). Hence, nanotechnological engineering of plant benefcial microorganisms can ultimately improve the yield and resistance of plants, and the approach is comparatively novel. The plant-associated bacteria directly promote the plant growth by nitrogen fxation and phosphate solubilization and also through the synthesis of phytohormones and siderophores [\[8\]](#page-13-4). Additionally, these bacteria have diverse mechanisms to protect the plant from pathogens [\[29\]](#page-14-5). Hence, improving these functioning of endophytes through nano-supplementation will have signifcant feld applications.

Zinc oxide nanoparticles (ZnONPs) and chitosan nanoparticles (CNPs) have already been studied to promote the plant growth and disease resistance. Even though ZnONPs are known to improve the seed germination and yield, at higher concentrations, these could be detrimental [[33](#page-14-6)]. But, zinc (Zn) has a vital role in carbohydrate and protein metabolism, and it regulates the synthesis of plant growth hormones like the indole-3-acetic acid (IAA). Zn is also an essential component of dehydrogenase, proteinase, and peptide enzymes and promotes starch formation, seed maturation, and production [\[24\]](#page-14-7). Chitosan has already been recognized to have the potential to control plant diseases by suppressing the growth, sporulation, and germination of pathogens and also by disrupting pathogens while inducing diferent defense responses in plant [\[14\]](#page-14-8).

In the current study, endophytic *Paenibacillus* sp. CCB36 was isolated from the rhizome of *Curcuma caesia* Roxb. (*Zingiberaceae*), which has been previously demonstrated to have anti-oxidant [\[7\]](#page-13-5), antimicrobial [[19](#page-14-9)], and anti-ulcerogenic [\[25\]](#page-14-10) activities and hence traditionally used as medicine [\[13\]](#page-13-6). The current study has been focused on the efect of ZnONPs and CNPs on the bioflm and biocontrol properties of isolated *Paenibacillus* sp. CCB36.

Materials and Methods

Isolation of Endophytic Bacteria from *C. caesia* **Roxb.**

The rhizomes of *C. caesia* Roxb. collected from Indian Council of Agricultural Research (ICAR)-Indian Institute of Spices Research, Kozhikode, Kerala, India, were used for the isolation of endophytic bacteria. The sterilization of surface of rhizome and the bacterial isolation were carried out as per the previously described methods [[16](#page-14-11)]. For this,

the rhizomes were washed with running tap water to remove the soil particles, and scales were also removed. These were further washed several times with distilled water. The samples were then dipped in 70% ethanol for 1 min and washed with sterile distilled water. The rhizome pieces were then treated with 2% sodium hypochlorite for 10 min followed by washed with sterile distilled water for eight times. For the isolation of endophytic bacteria, the processed rhizome was cut into pieces of 1 cm diameter and 1–2 cm length and placed on nutrient agar plates followed by incubation for 48 h at 28 ℃.

Screening of Antiphytopathogenic Property of Endophytic Isolates

Antifungal activity of all the isolated endophytes was done by dual culture method [[27](#page-14-12)] against phytopathogenic fungi like *Pythium myriotylum* and *Rhizoctonia solani*. For this, the isolates were streaked on one side of the nutrient agar plates and incubated for 2 days at 28 ℃. Mycelial discs of phytopathogenic fungi were further inoculated on the other side of nutrient agar plates, and media with phytopathogens alone were served as the control. All the plates were then incubated at 28 \degree C for 3–5 days and were observed for fungal growth inhibition. Among these, the isolate that showed strong antiphytopathogenic activity was further evaluated for activity against *Fusarium oxysporum*, *Sclerotium rolfsii*, *Phytophthora infestans*, *Rhizoctonia solani*, and *Colletotrichum acutatum*.

Molecular Identifcation of Selected Isolate

Here, 16S rDNA sequence similarity based identifcation of the selected isolate CCB36 was carried out as it showed inhibition to all the selected phytopathogens. Genomic DNA isolated was used as the template for PCR amplifcation of 16S rDNA using the primers 16SF (5′-GAG TTT GAT CCT GGC TCA G-3′) and 16SR (5′-GAT ATT ACC GCG GCG CCT G-3′). Polymerase chain reaction (PCR) was carried out in a MycyclerTM (BIORAD, USA) with 50 μL reaction volume containing 5 μL of genomic DNA, 2 μL of both forward and reverse primers (10 pM), and 25 μ L of mastermix (Takara), and it was made up to 50 μL with Milli-Q water. PCR reactions were cycled 35 times with initial denaturation at 94 ℃ for 3 min and the denaturation at 94 ℃ for 30 s, annealing at 58 ℃ for 30 s, and extension at 72 °C for 2 min were used cyclically with a final extension at 72 °C for 7 min. Then, the amplifed product was confrmed by agarose gel electrophoresis [\[16\]](#page-14-11). The product was further sequenced at AgriGenome Labs Pvt. Ltd., Cochin, Kerala, India. The obtained sequence data was subjected to BLAST (Basic Local Alignment Search Tool) analysis.

Synthesis and Characterization of CNPs

Chitosan from shrimp shells (Cat. No: PCT0817) with molecular weight 800–20,000 Daltons and degree of deacetylation (DDA) = 75.0% was procured from the HiMedia Laboratories, Mumbai, India. For the synthesis of CNPs, 0.3 g chitosan was dissolved in 100 mL of 1% acetic acid solution and stirred at 1000 rpm for 25 min at 28 °C until the solution became clear. An opalescent suspension was further formed upon the drop by drop addition of 10 mL of 1% sodium tripolyphosphate (TPP) aqueous solution to the chitosan solution under stirring at 28 °C. Then, the CNPs were purifed by centrifuging the suspension at 10,000 rpm for 30 min at $4 °C [6, 12]$ $4 °C [6, 12]$ $4 °C [6, 12]$.

MADZU UV-2600 series), Fourier transform infrared (FTIR) (Schimadzu IR Prestige 21 FTIR) spectroscopy, High-resolution-transmission electron microscopy (HR-TEM) (JEOL, JEM-2100), Scanning electron microscopy (SEM) (JEOL JSM-7610 FPlus), and Dynamic light scattering (DLS). UV–visible spectrum of CNPs was recorded in the range of 200 to 800 nm, and FTIR spectrum was recorded in the range of 4500 to 700 cm−1. CNPs for the TEM analysis were prepared as a flm on carbon-coated copper TEM grids and were allowed to stand for 2 min. The extra solution was removed by a blotting paper, and the grid was allowed to dry before analysis.

For the SEM analysis, mounting of dried CNPs was carried out on specimen stubs and coated with gold for 60 s using DII 29030SCTR smart coater [\[36\]](#page-14-13). The particle size of CNPs was analyzed by DLS (HORIBA SZ-100) at a sample holder temperature of 25 ℃.

Evaluation of Efect of ZnONPs and CNPs on Endophytic Isolate CCB36

ZnONPs (1.7 g/mL \pm 0.1 g/mL density at 25 °C and ≤40 nm average particle size) used were purchased from Sigma-Aldrich, USA (Cat. No.: 721077), and the CNPs were synthesized as described above. The endophytic isolate CCB36 was inoculated into 150 mL of nutrient broth and was incubated at 28 $^{\circ}$ C for 24 h with constant shaking at 200 rpm (ELTEK OS-3). After this, CCB36 was swabbed on nutrient agar plates and 2, 4, and 6 mg of ZnONPs, and CNPs were added into the respective wells made on the swabbed nutrient agar plates. The plates were further incubated at 28 $^{\circ}$ C for 24 h [\[18\]](#page-14-14).

Evaluation of Bioflm Formation of CCB36 in the Presence of ZnONPs and CNPs

The evaluation of bioflm formation was carried out by tissue culture plate assay. For this, overnight grown Luria Bertani (LB) broth (pH 7.5 ± 0.2) (Cat. No: M1245, HiMedia Laboratories, Mumbai, India) culture of isolate CCB36 was centrifuged at 5000 rpm for 5 min. The collected cells were washed with sterile phosphate-bufered saline (PBS), resuspended in LB broth, and made into the final optical density of 1 at 600 nm. Then, $200 \mu L$ bacterial suspension, bacterial suspension with 5 mg/mL CNPs, bacterial suspension with 20 μ g/ mL ZnONPs, and bacterial suspension with both 5 mg/mL CNPs and 20 μ g/mL ZnONPs were added into the wells of microtiter plate (96 wells). LB broth, LB broth with 5 mg/ mL CNPs, LB broth with 20 μ g/mL ZnONPs, and LB broth with both 5 mg/mL CNPs and 20 µg/mL ZnONPs were used as control, and the experiment was repeated for three times. After the incubation at 28 \degree C for 18–20 h, the culture suspension was removed and the wells were washed with sterile PBS twice to remove the non-adhered cells. The plates were air dried and stained with 0.1% crystal violet for 30 min at 28 °C. Excess stain was removed, and wells were washed twice with PBS. The wells were air dried, and 200 µL ethanol was added to the stained wells and kept for 10 min. Then, absorbance was recorded at 570 nm using Varioskan Flash spectral scanning multimode reader (Thermo Scientifc).

Statistical analysis

Data obtained from three replications of the experiment were statistically analyzed by using the software Origin Pro7 SRO (Northampton, MA, USA) with the Tukey's post hoc

multiple comparison tests to determine the significant difference in various groups at 5% level of signifcance [\[2](#page-13-9)].

Atomic Force Microscopy (AFM) for Bioflm Imaging

Here, overnight grown culture of CCB36 was centrifuged at 5000 rpm for 5 min at 4 $^{\circ}$ C. The collected cells were washed with sterile phosphate-bufered saline (PBS), and cells were resuspended in LB broth and diluted to the fnal optical density of 1 at 600 nm. After that, 200 µL bacterial suspension, bacterial suspension with 5 mg/mL CNPs, bacterial suspension with 20 µg/mL ZnONPs, and bacterial suspension with both 5 mg/mL CNPs and 20 µg/mL ZnONPs were added into the wells of a microtiter plate (96 wells) with control as described before. Sterile glass slide pieces $(1 \times 1 \text{ cm})$ were aseptically transferred to the bottom of each well, and after 18–20 h of incubation at 28 °C under static condition, the glass pieces were removed from wells with sterile forceps and gently washed with sterile PBS. The glass pieces were fxed with 2.5% glutaraldehyde for 1–2 h and dehydrated with 95% ethanol. Then, the fxed dried glass pieces were observed under Confocal Raman microscope with AFM (WITec Alpha300 RA, Germany) [\[15\]](#page-14-15). Here, the probe of the microscope connected to a cantilever is scanned over the surface of the glass piece, with a small repulsive force between the glass piece and the probe. The average roughness of bioflm under diferent treatments was measured by using the software "Gwyddion 2.53."

Evaluation of Bioflm Forming Property of CCB36 in the Presence of Diferent Concentrations of CNPs

Here, CCB36 cell suspension was prepared as described before. Then, 200 µL of the bacterial suspension and bacterial suspension with 1, 2.5, 5, 7.5, and 10, mg/mL of CNPs were separately added into the wells of a microtiter plate (96 well). Also LB broth and LB broth with 1, 2.5, 5, 7.5, and 10 mg/mL of CNPs were used as control. After the incubation at 28 °C for 18–20 h, the culture suspension was removed from the microtiter plate and processed as described above for analyzing bioflm [\[38\]](#page-14-16).

Evaluation of Biocontrol Property of CCB36 Against *Rhizoctonia solani*

Here, potato tuber pieces (2–3 cm in diameter) were surface sterilized by treating with 2% sodium hypochlorite solution for 2–5 min and washed thoroughly with sterile distilled water. Then, 30 mL of nutrient broth, overnight grown culture of CCB36, nutrient broth with CNPs (5 mg/mL), and CCB36 cultured with CNPs (5 mg/mL) were centrifuged at 5000 rpm for 5 min at 4 °C. The surface-sterilized potato pieces were further dipped in collected supernatant for 2–3 min and were then kept in sterilized Petri plates. Then, the pathogen *Rhizoctonia solani* was inoculated on each of the potato pieces and incubated for 3–5 days at 28 °C [[17](#page-14-17)].

Microscopic Analysis of Potato Tubers Using Lactophenol Cotton Blue (LPCB) Staining

A drop of lactophenol cotton blue stain was placed on the center of a clean slide, and tissues of potato pieces treated with *R. solani* as per described above were placed in the drop of stain and teased gently. A coverslip was kept over the stain, and the slides were observed under the microscope with $100 \times$ (oil immersion) magnification (Olympus-CX43) [[34](#page-14-18)].

Results

Plants provide vast and diverse niche for the endophytic microorganisms which could promote plant growth and disease resistance. The plant-associated microbial communities perform many vital functions for the optimal adaptation of the plant to the habitat.

Isolation of Endophytes from *C. caesia* **Roxb.**

In the study, rhizome of *Curcuma caesia* Roxb. was used for the endophytic bacterial isolation, and it resulted in the purifcation of 36 diferent bacterial isolates named as CCB1 to CCB₃₆

Antiphytopathogenic Property of Endophytic Isolates

All the bacterial isolates were screened for antiphytopathogenic property against *P. myriotylum* and *R. solani* (Table S1). Among these, CCB36 was found to have remarkable inhibitory efect against both the pathogens. Upon further analysis, CCB36 was also found to inhibit *F. oxysporum*, *S. rolfsii*, *P. infestans*, and *C. acutatum* (Fig. [1\)](#page-5-0). Hence, this was selected for further studies.

Fig. 1 Antiphytopathogenic activity of CCB36 against **a** *Fusarium oxysporum*, **b** *Sclerotium rolfsii*, **c** *Phytophthora infestans*, **d** *Rhizoctonia solani*, and **e** *Colletotrichum acutatum*

Molecular Identifcation of Endophytic Isolate CCB36

The PCR amplifcation product of 16S rDNA of CCB36 was found to have 1500 bp size which was further purifed and sequenced (Fig. S1). Sequence similarity search by BLAST analysis showed the 16S rDNA of CCB36 to have 100% identity with *Paenibacillus polymyxa*.

Synthesis and Characterization of CNPs

The clear chitosan solution was changed to an opalescent suspension upon the addition of aqueous solution of sodium tripolyphosphate (TPP), which indicated the formation of chitosan nanoparticles. This was further characterized by UV–visible spectroscopy, FTIR, TEM, SEM, and DLS analysis. The UV–visible spectrum of CNPs exhibited corresponding peak at 250 nm. Functional groups of CNPs were studied by FTIR analysis. The FTIR spectrum of the CNPs showed major absorption peaks at 3453.69, 3356.28, 3197.15, 3181.72, 1662.71, 1538.3, and 1329.98 cm−1, respectively. The spectral values of the CNPs between 3200 and 3550 cm^{-1} could be due to the -O–H bonding. The surface morphological structure was further examined by SEM and TEM analysis, which clearly showed spherical shaped chitosan nanoparticles. The particle size distribution of CNPs was measured by DLS, which showed the CNPs with average particle size of 105.7 nm (Fig. [2](#page-6-0)).

Fig. 2 Characterization of chitosan nanoparticles: **a** UV–visible spectrum, **b** Fourier transform infrared (FTIR) spectrum, **c** Transmission electron microscopic (TEM) image, **d** Scanning electron microscopic (SEM) image, and **e** Dynamic light scattering (DLS) histogram

Efect of ZnONPs and CNPs on Endophytic CCB36

Three concentrations (2, 4, and 6 mg) of ZnONPs and CNPs used were not found to have inhibitory efect on CCB36. Hence, these nanoparticles were used for further studies related to bioflm formation and biocontrol activity of CCB36.

Bioflm Formation of CCB36 in the Presence of Supplemented ZnONPs and CNPs

Microtiter plate assay was used to quantify the bioflm formed by the selected endophytic isolate. Here, CCB36 exhibited variation in bioflm forming property in the presence of supplemented ZnONPs and CNPs. Significant increase in the formation of biofilm was observed for CCB36 when supplemented with 5 mg/mL chitosan nanoparticles when compared with $ZnONPs$ (20 $\mu g/mL$) treatment and control (Fig. [3](#page-7-0)). In the presence of both ZnONPs and CNPs supplementation, CCB36 was found to form bioflm which was statistically comparable to unsupplemented and ZnONPs supplemented treatment.

Atomic Force Microscopy (AFM) Analysis on Bioflm Changes of CCB36 due to Nanoparticle Supplementation

AFM analysis showed the morphological features and 3D topography of bioflm formed by CCB36 in the presence of supplemented ZnONPs and CNPs (Fig. [4\)](#page-8-0). Here, increased thickness of bioflm of CCB36 could be observed by AFM in the presence of 5 mg/ mL CNPs supplementation when compared with other treatments. However, the bioflm

Fig. 3 Bioflm formation by *Paenibacillus* sp. CCB36 in the presence of chitosan nanoparticles and ZnO nanoparticles (LB+P, Luria Bertani broth with *Paenibacillus* sp. CCB36; LB+CNPs+P, Luria Bertani broth with chitosan nanoparticles (5 mg/mL) and *Paenibacillus* sp. CCB36; LB+ZnONPs+P, Luria Bertani broth with zinc oxide nanoparticles (20 μ g/mL) and *Paenibacillus* sp. CCB36; LB+CNPs+ZnONPs+P, Luria Bertani broth with chitosan nanoparticles (5 mg/mL), zinc oxide nanoparticles (20 µg/mL), and *Paenibacillus* sp. CCB36)

Fig. 4 AFM analysis of bioflm formed by *Paenibacillus* sp. CCB36 in the presence of zinc oxide nanoparticles and chitosan nanoparticles: **a** LB control, **b** LB with chitosan nanoparticles control, **c** LB with ZnO nanoparticles control, **d** LB with chitosan nanoparticles and ZnO nanoparticles control, **e** *Paenibacillus* sp. CCB36 inoculated LB broth, **f** *Paenibacillus* sp. CCB36 inoculated LB broth with chitosan nanoparticles, **g** *Paenibacillus* sp. CCB36 inoculated LB broth with ZnO nanoparticles, **h** *Paenibacillus* sp. CCB36 inoculated LB broth with chitosan nanoparticles and ZnO nanoparticles (LB- Luria Bertani broth and ZnO- zinc oxide)

forming property of CCB36 was decreased when 20 µg/mL of ZnONPs was added. This observation could also be refected in the evaluation of its roughness value (Fig. [5\)](#page-9-0). The roughness value of the CCB36 surface treated with CNPs showed the highest value among all the treatments used. This indicated enhanced bioflm formation by CCB36 in the presence of CNPs and was supportive to the result obtained in the microtiter plate assay*.*

Bioflm Formation of CCB36 in the Presence of CNPs

The CCB36 was observed to show diference in the thickness of bioflm produced when supplemented with diferent concentrations of chitosan nanoparticles. Specifcally, increased bioflm was found to be formed by CCB36 with the supplementation of 2.5 mg/ mL chitosan nanoparticles (Fig. 6).

Biocontrol Property of CCB36 Supplemented with CNPs Against *R. solani* **on Potato**

Surface-sterilized potato tuber pieces were used for this study. An enhanced growth of mycelia of *R*. *solani* was observed on the potato pieces which were treated with distilled water and sterile nutrient broth. Treatment with CCB36 was observed to provide protection to potato when compared with the control. Signifcantly reduced growth of *R. solani* was observed for the potato pieces treated with CCB36 cultured in the presence of CNPs. This revealed the enhanced antiphytopathogenic efect of CCB36 due to supplementation of CNPs (Fig. [7\)](#page-10-0).

Fig. 5 Surface profle of *Paenibacillus* sp. CCB36 bioflm formed under diferent treatments exhibiting variation in roughness (Rq) (LB, Luria Bertani broth; LB+CNPs, Luria Bertani broth with chitosan nanoparticles; LB+ZnONPs, Luria Bertani broth with zinc oxide nanoparticles; LB+CNPs+ZnONPs, Luria Bertani broth with zinc oxide nanoparticles and chitosan nanoparticles; $LB + P$, Luria Bertani broth with *Paenibacillus* sp. CCB36; LB+CNPs+P, Luria Bertani broth with chitosan nanoparticles and *Paenibacillus* sp. CCB36; LB+ZnONPs+P, Luria Bertani broth with zinc oxide nanoparticles and *Paenibacillus* sp. CCB36; LB+CNPs+ZnONPs+P, Luria Bertani broth with zinc oxide nanoparticles, chitosan nanoparticles, and *Paenibacillus* sp. CCB36)

Fig. 6 Bioflm formation of *Paenibacillus* sp. CCB36 in the presence of diferent concentrations of supplemented chitosan nanoparticles (P, *Paenibacillus* sp. CCB36; P+1C, *Paenibacillus* sp. CCB36 with 1 mg/mL chitosan nanoparticles; P+2.5C, *Paenibacillus* sp. CCB36 with 2.5 mg/mL chitosan nanoparticles; P+5C, *Paenibacillus* sp. CCB36 with 5 mg/mL chitosan nanoparticles; P+7.5C, *Paenibacillus* sp. CCB36 with 7.5 mg/mL chitosan nanoparticles; P+10C, *Paenibacillus* sp. CCB36sp. CCB36with 10 mg/ mL chitosan nanoparticles)

Fig. 7 Biocontrol activity of *Paenibacillus* sp. CCB36 supplemented with chitosan nanoparticles against *Rhizoctonia solani* infection on potato: **A** distilled water, **B** nutrient broth control, **C** chitosan nanoparticles treated, **D** *Paenibacillus* sp. CCB36 treated, **E** *Paenibacillus* sp. CCB36 cultured with supplemented chitosan nanoparticles treated

Microscopic Analysis of Potato

Microscopic evaluation of antiphytopathogenic activity of CCB36 against *R. solani* on potato was also conducted. Samples from potato pieces treated with distilled water, nutrient broth, and CNPs exhibited the presence of *R. solani* hyphae abundantly near the starch grains of potato. The treatment with suspension of CCB36 slightly reduced the number of *R. solani* hyphae. The samples from potato pieces treated with suspension of CCB36 cultured with CNPs showed minimum hyphae of *R. solani* which confrmed the enhanced biocontrol effect of CCB36 when cultured with CNPs supplementation (Fig. [8\)](#page-11-0).

Discussion

Plants have been documented as meta-organisms, holding a diverse microbiome with mutual symbiotic relationship. Each plant is associated with various species of microbial strains and majority of them perform signifcant ecological functions benefcial to the plant [[3\]](#page-13-10). These groups of organisms encourage the plant growth and also provide resistance to both biotic and abiotic stress conditions $[20]$. The beneficial microbiome of medicinal plants are also known to secrete distinctive bioactive secondary compounds having broad range of applications [[9\]](#page-13-11).

In the study, rhizome of *Curcuma caesia* Roxb. was selected for the isolation of endophytic bacteria as it was traditionally used as a medicine due to rich content of bioactive molecules. Hence, its associated microbiome might also expect to have remarkable application. Among all the endophytic bacteria isolated, CCB36 identifed as *Paenibacillus* sp. exhibited strong antifungal activity against phytopathogens. *Paenibacillus* spp. are often considered antimicrobial factories due to its ability to synthesize biologically active molecules which are inhibitory to phytopathogens. Various strains of *P. polymyxa* have formerly been found to antagonize *Phytophthora* sp., *Rhizoctonia solani*, *Pseudomonas syringae*, and *Xanthomonas campestris*. The chemical basis of the same has also been considered as antimicrobial compounds such as polymyxin, fusaricidin A, gavaserin, and saltavalin. These metabolites can mechanistically cause the cell lysis and leakage of ions, disruption of the structural integrity of membranes, inhibition of mycelial growth, inhibition of spore germination, and protein synthesis [\[37,](#page-14-20) [39\]](#page-14-21).

Fig. 8 Microscopic images showing biocontrol efect of *Paenibacillus* sp. CCB36 against *Rhizoctonia solani* infection on potato: **a** *Rhizoctonia solani*, **b** potato treated with distilled water, **c** potato treated with nutrient broth, **d** potato treated with chitosan nanoparticles, **e** potato treated with *Paenibacillus* sp. CCB36, and **f** potato treated with *Paenibacillus* sp. CCB36 cultures with supplemented chitosan nanoparticles (red arrows indicate the presence of *Rhizoctonia solani* hyphae)

The endophytic CCB36 with antiphytopathogenic property was used as the candidate organism in the study to demonstrate its colonization through bioflm and enhanced antagonism with the supplementation of CNPs and ZnONPs. Even though colonization on plant tissue is the primary requirement for the functional establishment of endophytes, most of them cannot colonize and establish bioflm formation on non-host plant tissues. In the screening for bioflm formation, the CCB36 was found to form increased bioflm in the presence of chitosan nanoparticles when compared to the control. However, ZnONPs suppressed the formation of bioflm. The optimum bioflm was formed in the presence of 2.5 mg/mL CNPs and then declined when concentration was increased. Previous study has already reported the coupling of chitosan with plant growth-promoting rhizobacteria to have enhancement efect on its plant benefcial traits [[1\]](#page-13-12). But the augmented bioflm forming property of plant benefcial microorganism due to coupling with chitosan nanoparticles has not yet been studied in detail. This indicates the novelty of the study and can have signifcant applications in delivery and feld performance of plant benefcial microorganisms.

Atomic force microscopy (AFM) was used to visualize the bioflm forming property of CCB36 on a non-living surface. AFM has been used in the study for the direct nanoscale observation on the surface morphology of CCB36 supplemented with nanosized particles and bacterial bioflms formed. More aggregated bacterial cells were observed in the AFM topography of CCB36 treated with chitosan nanoparticles. The

average roughness value of surface of the same treatment was higher than other treatments and control. In the presence of chitosan nanoparticles, the bacteria exhibited intense adhesion and aggregation. This suggests that it may be utilized to boost the colonization of selected bacteria on plants.

The biocontrol property of CCB36 was also evaluated in the presence of chitosan nanoparticles on potato against *R. solani*. Potato pieces treated with the cell-free supernatant of CNP-treated CCB36 were found to have enhanced protection from the infection caused by *R. solani* when compared with CCB36 alone treatment. Many previous studies have already demonstrated the antimicrobial applications of chitosan in agriculture, food, and pharmaceutical industries [\[22,](#page-14-22) [30](#page-14-23), [31,](#page-14-24) [35](#page-14-25), [40](#page-14-26)]. As the CNP-treated tubers were not protected from the infection by *R. solani*, the chitosan nanoparticles alone used in the present study could not provide any direct antagonism to *R. solani*. But, these CNPs functioned efectively to boost up the biocontrol property of CCB36. Here, the used CNPs might have induced the CCB36 as a pathogen, because chitin is a major component of cell wall of most of the fungal phytopathogens.

The fnding was further substantiated by a microscopic analysis of samples collected from potato pieces treated with CCB36 supernatant where CCB36 was cultured in the presence of CNPs. This demonstrated the suppression of mycelia growth on potato samples. This might be due to the activity of induced metabolite production by CCB36 in response to the supplemented chitosan nanoparticles. *Paenibacillus* spp. have already been reported to offer protection against diseases affecting a wide variety of crops [[29](#page-14-5)]. Hence, the results obtained in the current study can have application to improve the functioning of other *Paenibacillus* spp. also. The isolate CCB36 with CNPs supplementation can have signifcant applications in the agricultural feld.

Conclusions

Endophytic bacteria ubiquitously colonize the internal tissues of plants and provide resistance from drought, heavy metals, salt, and phytopathogens and also promote the host growth through various mechanisms*.* In the present study, *Paenibacillus* sp. CCB36 with potential antifungal activity was isolated and found to have an increased ability to form bioflm in the presence of chitosan nanoparticles. Bioflm formation of plant benefcial bacteria is a crucial primary phase for the colonization on plant tissues. In addition to the bioflm formation, CNPs enhanced the biocontrol property of CCB36 which provided better protection to the potato pieces against *Rhizoctonia solani*. Microscopic analysis confrmed the presence of abundant fungal mycelia in potato pieces treated with control and enhanced suppression of pathogens on potato treated with CCB36 cultivated with supplemented CNPs. These results indicate the modulating efect of chitosan nanoparticles on plant benefcial traits of CCB36 such as bioflm formation and antagonism against *R. solani*. Hence, the coupling of CCB36 with chitosan nanoparticles will be an effective ecofriendly means to control crop plant disease in the agricultural feld.

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Author Contribution Radhakrishnan Edayileveetil Krishnankutty contributed to the study conception and design. Experiments and analysis were performed by Jishma Panichikkal, Sreejith Sreekumaran, Ashitha Jose, Anju Kanjirakandi Ashokan, and Cimmiya Susan Baby. The frst draft of the manuscript was written by Jishma Panichikkal. The manuscript was revised, and corrections were included by Radhakrishnan Edayileveetil Krishnankutty, Jishma Panichikkal, and Ashitha Jose. All authors commented on previous versions of the manuscript. All authors read and approved the fnal manuscript.

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Declarations

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