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## Research Article

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# **Harnessing the biocontrol and phytostimulatory potential of endophytic fungi against damping -off disease in okra**

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## **Abstract**

Okra (*Abelmoschus esculentus* L. Moench) is an economically important vegetable crop, with summer cultivation critical for early market supply. Seedling damping-off, primarily caused by *Rhizoctonia solani*, poses a major constraint, often leading to severe yield losses. This study evaluated the potential of fungal endophytes for managing damping-off under summer cultivation in Himachal Pradesh, India. Field surveys (2023–2024) across Kangra, Mandi, and Hamirpur districts recorded disease incidence of 31.7–84.3%. Among 44 isolates recovered, *R. solani* JPO1 was identified as the predominant pathogen via morpho-molecular characterization. Forty-two fungal endophytes were screened for antagonism and plant growth-promoting (PGP) traits; eight isolates with strong activity were further characterized: *Coniothyrium* sp. E1P, *Xylaria juruensis* E4P, *Aspergillus terreus* E10P, *Epicoccum* sp. E13P, *Aspergillus welwitschiae* E16P, *Fusarium proliferatum* E25P, *F. solani* E26P, and *Epicoccum* sp. JPE2. Glasshouse and field evaluations identified *F. proliferatum* E25P as the most effective isolate, significantly enhancing plant growth and suppressing damping-off. Root colonization, confirmed by plate counts and confocal laser scanning microscopy, demonstrated robust endophytic establishment in treated plants. These

results highlight the potential of fungal endophytes as sustainable, eco-friendly alternatives to chemical fungicides for effective damping-off management in okra.

**Keywords:** Okra; damping-off; endophytic fungi; *Rhizoctonia solani*; root colonization; biocontrol

#### **Statements and Declarations**

**Availability of data and material:** All sequences generated in this study are available in GenBank with accession numbers cited in Table 1. The remaining datasets not included here will be made available from the corresponding author on reasonable request

**Competing interests:** The authors declare that they have no competing interests

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## Introduction

Okra (*Abelmoschus esculentus* L.) is an important vegetable crop cultivated widely in tropical and subtropical regions, particularly across the Indian subcontinent, including Himachal Pradesh (Gaur et al. 2025; Kumar et al. 2025). The crop is known by several local names such as “lady’s finger” in England, “gumbo” in the United States, “bhindi” in India, and “bamia” in Ethiopia (Benchasri 2012; Gemedede et al. 2015). Okra pods are valued for their nutritional richness, containing carbohydrates, proteins, dietary fiber, vitamins, and essential minerals. They also possess bioactive compounds associated with antioxidant and other health-promoting properties (Mishra et al. 2017; Wu et al. 2022; Gaur et al. 2025). Owing to its nutritional value and low caloric content, okra contributes significantly to dietary health and food security (Obomeghei et al. 2022). India is the largest producer of okra in the world, contributing more than 72% of global production, with an annual output of about 6.35 million tonnes cultivated over 0.52 million hectares (FAOSTAT 2020; Karmakar et al. 2025). The crop can be grown during both summer and rainy seasons, although summer cultivation is often preferred because early harvests generally fetch better market prices (Rubatzky and Yamaguchi 1997).

Okra cultivation is affected by several diseases incited by fungi, bacteria, and viruses, including root rot, wilt, leaf spot, anthracnose, powdery mildew, and yellow vein mosaic virus (Patel et al. 2019). Among these, damping-off is particularly damaging during the early stages of crop establishment. The disease affects seeds and young seedlings and is caused by a complex of soil-borne pathogens, including *Pythium* spp., *Phytophthora nicotianae*, *Rhizoctonia solani*, and *Macrophomina phaseolina* (Diwakar et al. 1986; Jiskani et al. 2007; Matny 2013; Jukte et al. 2016). Infection at this stage often results in seed rot, poor germination, and collapse of young seedlings, leading to significant crop loss in the field.

Among the pathogens associated with damping-off, *Rhizoctonia solani* Kühn is considered one of the most aggressive, particularly under warm conditions typical of summer cultivation. The pathogen causes seed rot as well as pre- and post-emergence damping-off, resulting in rapid seedling collapse and reduced plant stand (Abbas et al. 2017; Muimba-Kankolongo 2018; Patel et al. 2019; Nasimi et al. 2024). In later stages of growth, *R. solani* may also cause root rot, further affecting crop productivity (Mimma et al. 2023). Its persistence in soil is aided by the formation of sclerotia, which enable the pathogen to survive in soil and crop residues for extended periods, making disease management difficult (Akber and Fang 2024).

Damping-off in okra is commonly managed through fungicidal seed treatments or soil applications. Although these treatments can suppress the disease, their frequent use has raised concerns about environmental contamination, development of resistant pathogen populations, adverse effects on beneficial soil microorganisms, and possible chemical residues in edible produce (Pandey et al. 2018; Bhattarai et al. 2022). These limitations have encouraged the search for safer and more sustainable approaches to disease management.

Biological control using beneficial microorganisms has gained increasing attention as an alternative to chemical control. Among these, fungal endophytes have attracted particular interest. Endophytic fungi colonize internal plant

tissues without causing visible harm and often form beneficial associations with their host plants (Murthy et al. 2014; Alam et al. 2021; Attia et al. 2022a, b; Abd Alhakim et al. 2022). By occupying internal plant niches, these microorganisms can limit the establishment of invading pathogens through competition for space and nutrients (de Lamo et al. 2018).

Endophytic fungi may also contribute to plant protection by enhancing plant defense responses. They can stimulate defense-related enzymes, promote the accumulation of phenolic compounds, strengthen plant cell walls, and improve the plant's ability to cope with oxidative stress (Llorens et al. 2017; Constantin et al. 2019; Daroodi et al. 2021; El-Sharkawy et al. 2022). In addition, they may inhibit pathogens through mechanisms such as nutrient competition, production of antimicrobial metabolites, induction of systemic resistance, and mycoparasitic interactions involving hyphal coiling, penetration, and secretion of cell wall-degrading enzymes (Waghunde et al. 2017; Sharaf et al. 2022). Many endophytic fungi also produce diverse secondary metabolites that contribute to disease suppression and plant growth promotion (El Enshasy et al. 2019). Their potential for managing plant diseases has been reported in several crop systems (Rashad et al. 2020; Pal et al. 2020; Campos and Jacob 2021; Kushwaha et al. 2024).

Despite these promising attributes, the role of endophytic fungi in managing damping-off in okra has received limited attention, particularly under the agro-climatic conditions of the Himalayan region. Considering the economic importance of okra and the destructive nature of damping-off pathogens such as *R. solani*, identifying effective endophytic fungi that can suppress disease while promoting plant growth is important for sustainable crop production. The present study therefore aimed to isolate and characterize fungal endophytes associated with okra and to evaluate their biocontrol and phytostimulatory potential against the damping-off pathogen of summer-grown okra. The findings are expected to contribute to the development of environmentally sound strategies for managing this disease.

## **Materials and methods**

### **Sampling, isolation and purification of the microbial culture (s)**

Extensive surveys were conducted in three districts of Himachal Pradesh (Kangra, Mandi, and Hamirpur) during the summer season (April–June 2023) to determine the prevalence of damping-off in okra. Surveyed locations ranged in altitude from 350 to 2251 m.a.s.l. Healthy and symptomatic okra roots exhibiting typical damping-off symptoms (water-soaked lesions or brown discoloration at the collar region) were carefully uprooted, placed in sterilized polythene bags, and transported to the laboratory for processing within 24–48 hours. These samples were used to isolate both the causal pathogen(s) and fungal endophytes.

### **Isolation and purification of causative pathogen(s)**

Infected root segments were washed under running water, dried on Whatman No.1 filter paper, and dissected into 0.5–1 cm pieces from the junction of healthy and diseased tissue. Tissue fragments were surface-sterilized with 70% ethanol for 1 min, followed by 1% NaOCl for 1 min, and rinsed four times with sterile distilled water. Excess

moisture was removed on sterilized filter paper, and fragments were placed (four per plate) onto Potato Dextrose Agar (PDA) supplemented with antibiotics to suppress bacterial growth. Plates were incubated at  $25 \pm 2^\circ\text{C}$  in a BOD incubator for 4 days. Emerging mycelia were purified using the hyphal tip method, and axenic cultures were maintained on PDA slants at  $4^\circ\text{C}$  (Ghaffar et al. 2018).

Colonization rate (CR) of pathogens was calculated as per Madhi et al. (2020):

$$\text{Colonization Rate (CR)} = \frac{\text{Total number of segments yielding fungus}}{\text{Total number of segments plated}} \times 100$$

Morpho-cultural identification was performed using a compound microscope (Olympus BX50) and reference descriptions (Waterhouse 1967; Parmeter 1970; Booth 1971; Ho 1981; Nelson et al. 1983; Boerema et al. 2004; Samson et al. 2011).

### **Pathogenicity bioassay**

To confirm pathogenicity, inoculum of *R. solani* was prepared on sand–wheat meal (8:2) medium in 500 ml Erlenmeyer flasks (Sharma 2011). Ten mycelial discs (5 mm) from 3-day-old cultures were inoculated into sterilized medium and incubated at  $25 \pm 2^\circ\text{C}$  for 12 days, with shaking every 2–3 days. The colonized medium was air-dried and stored at  $4^\circ\text{C}$  for subsequent use.

Glasshouse bioassays were conducted using the susceptible okra cultivar ‘Palam Komal’. Sterilized pots (20 cm diameter) were filled with clay:sand:FYM (2:1:1) mixture and inoculated with *R. solani* inoculum (100 g/kg). Surface-sterilized seeds were sown, and disease development was monitored. Pathogen re-isolation from diseased seedlings fulfilled Koch’s postulates. The most aggressive isolate from Palampur was selected for further experiments.

### **Isolation and quantification of fungal endophytes**

Fungal endophytes were isolated from asymptomatic roots following Sharma and Roy (2015). Roots were washed, cut into 0.5 cm fragments, and surface-sterilized sequentially with 70% ethanol (1 min), 4% NaOCl (5 min), 70% ethanol (30 s), and rinsed five times with sterile distilled water. Fragments were macerated in 10 ml sterile water and serially diluted to  $10^{-5}$ . Aliquots (100  $\mu\text{l}$ ) were plated on PDA ( $10^{-1}$  to  $10^{-5}$ ), in triplicate, and incubated at  $25 \pm 2^\circ\text{C}$  for 2 weeks. Outgrowing hyphal tips were purified, maintained on PDA slants at  $4^\circ\text{C}$ , and sub-cultured every 9–10 weeks. Fungal populations were quantified using the plate count technique (Agarwal and Hasija 1986):

$$\text{Cfu/g} = \frac{\text{No. of colonies} \times \text{Dilution factor}}{\text{Weight of root sample taken (g)}}$$

### **Morpho-molecular characterization**

#### **Morpho-cultural identification**

Pure cultures of pathogens and selected endophytes were assessed on PDA after 7 days for colony color, hyphal septation, spore morphology, pigmentation, and sclerotia formation. Microscopic observations were performed using

slide culture stained with lactophenol cotton blue at 40× magnification (Olympus CX51) and compared with standard keys (Christensen and Raber 1978; Parmeter 1970; Booth 1971; Nelson et al. 1983; Samson et al. 2011; Mathur and Olga 2003; Hussain et al. 2014; Song et al. 2016; Ying et al. 2019; Shehabeldine et al. 2024).

#### **Molecular identification**

Pure cultures of *R. solani* and seven efficient endophytes (excluding JPE2) were sequenced for 18S rRNA (Eurofins Genomics, India). Sequences were analyzed using BLAST (NCBI), and Maximum Likelihood phylogeny was constructed in MEGA v.11 (Tamura et al. 2021). Consensus sequences were submitted to GenBank.

#### ***In vitro* confrontation assay of fungal endophytes**

Dual culture assays were performed to assess antagonism of 42 fungal endophytes against *R. solani* (Coskuntuna and Ozer 2008). Agar plugs (5 mm) of both pathogen and endophyte were placed diametrically opposite on PDA plates (6 cm apart, 1 cm from edge) and incubated at 25 ± 2°C for 7 days. Pathogen radial growth was measured, and inhibition (%) was calculated (Gaigole 2011):

$$\text{Inhibition (\%)} = \frac{R1 - R2}{R1} \times 100$$

Where;

R1 = Radial growth of pathogen in control and R2 = Radial growth of pathogen in the presence of endophyte

#### **Microscopic observation of pathogen-fungal endophyte interactions**

Fungal endophytes exhibiting significant mycelial inhibition of *Rhizoctonia solani* under *in vitro* were examined under compound microscopy to evaluate their interaction with the pathogen (Yurnaliza et al. 2014; Kushwaha et al. 2024). Mycelium from the interaction zone was collected using a sterilized needle and placed onto a clean microscope slide with a drop of distilled water. A coverslip was gently applied, and alterations in hyphal growth at the interaction zone were observed at 40× magnification, comparing treated and control samples.

#### **Characterization of endophytic fungi for plant growth promotion, stress tolerance, and phytochemical profiling**

Endophytic fungi exhibiting ≥40% *in vitro* antifungal activity were selected for further characterization.

#### ***In Vitro* Plant Growth-Promoting Profiling**

**Siderophore Production:** Isolates were precultured on PDA for 7–10 days at 25 ± 2°C. Agar plugs (3 mm) were placed on Chrome Azurol S (CAS) agar and incubated at 25 ± 2°C for 5–6 days. Color changes from blue to yellow, purple, red, or orange indicated positive siderophore production. Zone diameters were measured by subtracting colony diameter from total halo zone (Schwyn and Neilands 1987).

**Phosphate Solubilization:** Fungal stubs were inoculated onto Pikovskaya's agar (Himedia, India) containing  $\text{Ca}_3(\text{PO}_4)_2$  and incubated for 5–6 days at  $25 \pm 2^\circ\text{C}$ . Formation of a clear halo indicated phosphate solubilization; zone diameter calculated as total zone minus colony diameter (Potshangbam et al. 2017).

**Cellulase Activity:** Cultures were inoculated on YEPD medium with 0.5% Na-carboxymethyl cellulose (CMC), incubated at  $25 \pm 2^\circ\text{C}$  for 5–6 days, stained with 0.2% Congo red, and destained with 1 M NaCl for 15 min. Clear zones indicated cellulase production (Toghueo et al. 2017).

**Amylase Activity:** Fungi were grown on Glucose YEPD agar amended with 0.2 g starch, incubated at  $25 \pm 2^\circ\text{C}$ , and flooded with 1% iodine solution. Clear zones indicated starch hydrolysis (Hankin and Anagnostakis 1975).

**Protease Activity:** 3 mm agar plugs from 10-day-old cultures were inoculated onto Casein Starch Agar with 1% skimmed milk, incubated at  $25 \pm 2^\circ\text{C}$  for 5–6 days. Clear halos indicated proteolytic activity (Hankin and Anagnostakis 1975).

**HCN Production:** Isolates were inoculated on PDA slants supplemented with 4.4 g/L glycine. Whatman No.1 filter paper saturated with 2% sodium carbonate in 0.5% picric acid was placed in tubes, sealed, and incubated at  $28^\circ\text{C}$  for 7 days. Yellow discoloration indicated HCN production (Loreck 1948).

### ***Abiotic Stress Tolerance***

**Salt Stress:** Isolates were spot-inoculated on PDA amended with 1–5% NaCl and incubated at  $25 \pm 2^\circ\text{C}$  for 2 weeks; growth was recorded.

**pH Tolerance:** Potato Dextrose Broth adjusted to pH 2, 5, and 8 using glacial acetic acid; cultures incubated for 2 weeks; growth monitored via optical density.

**Temperature Stress:** Isolates incubated at  $4^\circ\text{C}$ ,  $10^\circ\text{C}$ , and  $40^\circ\text{C}$  on PDA; growth observed after 2 weeks.

### **Phytochemical profiling**

Crude fungal extracts were screened for alkaloids, phenols, tannins, triterpenoids, saponins, and flavonoids following standard protocols (Bhardwaj et al. 2015; Thorati and Mishra 2017; Jagadevi Shivaputrapa 2018). Reactions were visualized by colorimetric changes as described in the original protocols.

### ***In vivo* Biocontrol Efficacy**

Eight fungal endophytes with highest *in vitro* antagonistic activity were selected for *in vivo* testing.

### **Preparation of liquid spore suspensions and seed treatment with endophytic fungi**

**Spore Suspension and Seed Treatment:** One-week-old cultures were transferred aseptically to PDB (300 ml in 500 ml flasks) and incubated on a rotary shaker at 180 rpm,  $25 \pm 2^\circ\text{C}$  for 7 days. Spore suspensions were filtered, quantified using a Neubauer's haemocytometer, and standardized to  $1 \times 10^6$  spores/ml. Surface-sterilized okra seeds (*Palam Komal*) were soaked overnight and air-dried for 3 h. Soil in pots was inoculated with *R. solani* mass culture at 100 g/kg.

## 235 Glasshouse Evaluation

236 Treated seeds were sown in 20 cm diameter pots containing sterilized clay:sand:FYM (2:1:1). Soil sterilization  
237 involved 5% formaldehyde treatment under polythene cover for 10–15 days, followed by aeration for 1 week. Six  
238 seeds were sown per pot; experiment arranged in CRD with five replicates. Glasshouse conditions: 25°C, 14/10 h  
239 light/dark, 85% RH. Regular irrigation was provided. Data recorded at 10 days' post-inoculation The following  
240 parameters were recorded:

## 241 Disease Incidence

$$242 \quad \text{Disease incidence (\%)} = \frac{\text{Number of diseased plants}}{\text{Total number of plants observed}} \times 100$$

## 243 Disease Control

$$244 \quad \text{Disease control (\%)} = \frac{[\text{Incidence in control} - \text{Incidence in treatment}]}{\text{Incidence in control}} \times 100$$

## 245 Plant growth parameters

246 **Germination (%):** Germination was recorded 10 days after sowing and per cent was estimated as the number of  
247 seeds sown and the number of seeds germinated expressed in percentage.

248 **Vigour index (cm):** Vigour index (%) = (Mean root length + Mean shoot length) x Germination (%)

249 **Shoot length (cm):** Shoot length was recorded in centimeters from the ground level to the apical bud stem.

250 **Root length (cm):** The length of tap root was recorded in centimeters using measuring scale by placing it  
251 horizontally on the ground.

## 252 Field Evaluation

253 Field trials were conducted at CSKHPKV Research Farm (2024) using RBD with three replications, plot size 2.25  
254 m<sup>2</sup> (4 rows × 20 plants). Seeds were soaked in endophytic spore suspensions (1 × 10<sup>6</sup> spores/ml). Natural epiphytic  
255 conditions were utilized; PDB-treated seeds served as control. Disease and growth parameters were recorded as  
256 described above.

## 257 Root Colonization Assay

258 Colonization of roots by representative endophytes was assessed through re-isolation at 7 and 14 days after  
259 inoculation (DAI). The fungal population was enumerated at 10<sup>-2</sup> dilution (cfu/g) on PDA following the methods of  
260 Agarwal and Hasija (1986), Barrett et al. (2008), and Miles et al. (2012). Further, for confirmation of endophytic  
261 colonization using laser scanning confocal microscopy (LSCM), the fungal treated roots were rinsed 3–4 times with  
262 autoclaved distilled water and then treated with 10% KOH solution overnight at 4°C, followed by washing with 1×  
263 PBS (2–3 times), the roots were then kept at RT for 10 minutes, followed by vacuum infiltration for 30 minutes  
264 after being incubated in 1mL of staining solution containing aniline blue, alexafluor 488-WGA, propidium iodide &

Tween 20 in PBS buffer and images were captured using STELLARIS 5, Leica Microsystems, Germany and represented following the protocol of Bhardwaj et al. (2025).

## Statistical Analysis

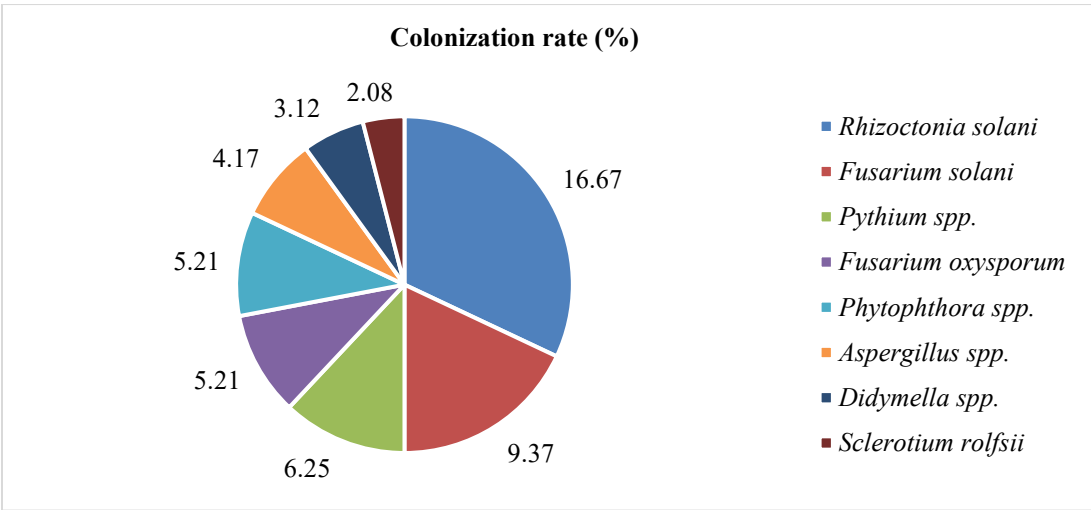
Laboratory and field data were analyzed using ANOVA (OPSTAT software; Gomez and Gomez, 1984). CRD and RBD designs with a minimum of three replicates were applied. Treatment means were compared using LSD at 5% significance level ( $p < 0.05$ ).

## Results

### Isolation and Identification of Pathogens Associated with Damping-Off of Okra

Surveys conducted during April–June 2023 across Kangra, Mandi, and Hamirpur districts of Himachal Pradesh revealed widespread damping-off in all okra-growing localities, predominantly at post-emergence stages. The highest disease incidence was observed in Tal, Hamirpur (84.33%), followed by Bhota, Hamirpur (81.67%) and Kanaid, Mandi (81.00%), whereas the lowest incidence was recorded in Naun, Mandi (31.67%). District-wise, Hamirpur showed the highest mean incidence (65.46%), followed by Mandi (59.04%), with Kangra having the lowest (53.33%). Overall, the incidence ranged from 31.67% to 84.33%, and no surveyed fields were disease-free, indicating a critical constraint on okra cultivation in Himachal Pradesh.

A total of 44 pathogenic isolates were recovered. *Rhizoctonia solani* exhibited the highest colonization frequency (16.67%), followed by *Fusarium solani* (9.37%). In total, eight fungal genera were associated with okra damping-off (Fig. 1), including *R. solani* (16 isolates), *F. solani* (8), *Pythium* spp. (6), *F. oxysporum* (4), *Phytophthora* sp. (4), *Aspergillus* sp. (3), *Didymella* sp. (2), and *Sclerotium rolfsii* (1). These data indicate *R. solani* as the predominant causal agent of damping-off in the region.



**Fig. 1** Frequency percentage of pathogenic fungal isolates associated with okra damping-off disease in different districts of Himachal Pradesh

## Pathogenicity Bioassay

Among the 44 isolates, *R. solani* was selected for further experiments. Under controlled pot conditions using the susceptible cultivar ‘*Palam Komal*’, soil inoculated with *R. solani* produced typical damping-off symptoms within seven days’ post-inoculation, primarily at the 2–3 leaf stage. Infected seedlings showed irregular, water-soaked, brown lesions at the basal stem, leading to wilting, leaf drooping, and seedling collapse. Koch’s postulates were fulfilled through re-isolation, and morphological comparison confirmed the pathogenicity of *R. solani*.

## Isolation and Quantification of Root Endophytic Fungi

Healthy okra roots harbored diverse indigenous fungal populations. Maximum endophyte densities were recorded in Palampur and Sunehar (Kangra) and Naun (Mandi) at  $4.33 \times 10^2$  cfu/g root, followed by Tikkar, Mandi ( $4 \times 10^2$  cfu/g root). The lowest populations were recorded in Tal, Nalti, and Bhota (Hamirpur;  $1 \times 10^2$  cfu/g root). Overall, Kangra contributed the highest number of isolates (18), followed by Mandi (13) and Hamirpur (10), yielding 41 endophytic isolates. These findings suggest that host genotype and geographical location influence root-associated microbial communities.

## Identification of Predominant Pathogen and Selected Fungal Endophytes

### Morpho-Molecular Identification of *Rhizoctonia solani*

The predominant isolate of *R. solani* formed cream to light brown colonies with floccose aerial hyphae on PDA after 5–6 days at  $25 \pm 1^\circ\text{C}$ . Within 20 days, dark brown sclerotia (0.2–1.0 mm) developed. Microscopically, hyphae exhibited right-angled branching with constriction and septation (width 6.7–7.9  $\mu\text{m}$ ; mean 7.3  $\mu\text{m}$ ). ITS sequence analysis confirmed 99.36% identity with *R. solani* (isolate JPO1; GenBank PQ605661). Phylogenetic analysis (Maximum Likelihood, MEGA 11) revealed close relation to *R. solani* isolate GPB (MK621284.1) (Fig. 2).

### Morpho-Cultural and Molecular Identification of Selected Endophytes

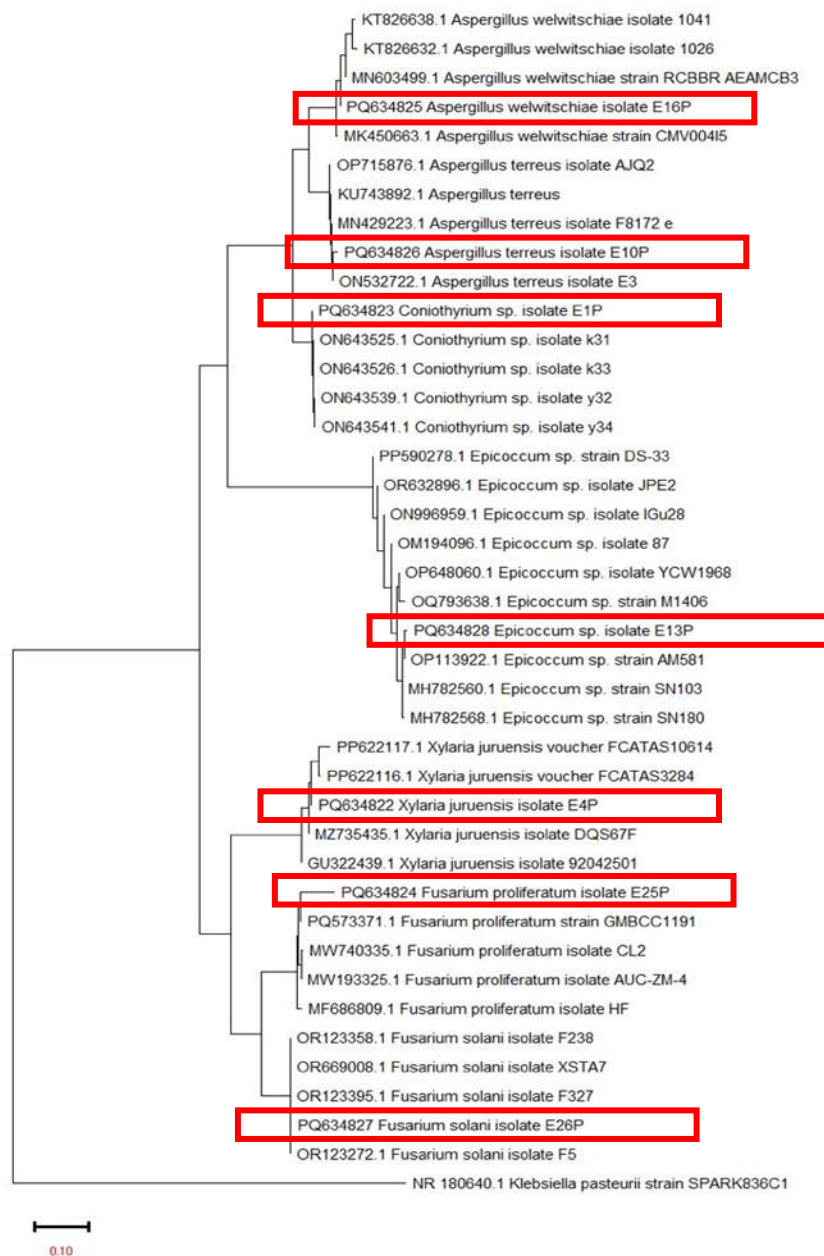
Eight endophytic fungi exhibiting  $\geq 40\%$  pathogen inhibition were characterized. Colony morphologies ranged from creamish white to dark brown or dull green, with septate hyphae and varying sporulation. Growth rates ranged from 6 to 24 days, fastest in E16P (6 days), slowest in E4P (24 days). The isolates E10P and E16P were marked as highly sporulating, E13P, E25P, E26P and JPE2, were recorded as less sporulating. However, E1P and E4P were categorized in to very less sporulating group of endophytic fungi. The sequence data from these fungal isolates were deposited in GenBank (the accession numbers are provided in Table 1). Phylogenetic analysis confirmed evolutionary relationships with top BLAST hits (Fig. 2).

**Table 1** Identification of endophytic fungi recovered from different districts of Himachal Pradesh

| Isolate | Identified fungi                    | Closest Neighbour                              | GenBank accession | Match identity (%) |
|---------|-------------------------------------|------------------------------------------------|-------------------|--------------------|
| E1P     | <i>Coniothyrium</i> sp. isolate E1P | <i>Coniothyrium</i> sp. isolate k31 ON643525.1 | PQ634823          | 98.28              |

|             |                                              |                                                                 |          |       |
|-------------|----------------------------------------------|-----------------------------------------------------------------|----------|-------|
| <b>E4P</b>  | <i>Xylaria juruensis</i><br>isolate E4P      | <i>Xylaria juruensis</i> isolate DQS67F MZ735435.1              | PQ634822 | 99.12 |
| <b>E10P</b> | <i>Aspergillus terreus</i><br>isolate E10P   | <i>Aspergillus terreus</i> isolate E3 ON532722.1                | PQ634826 | 98.03 |
| <b>E13P</b> | <i>Epicoccum</i> sp.<br>isolate E13P         | <i>Epicoccum</i> sp. strain AM581 OP113922.1                    | PQ634828 | 98.31 |
| <b>E16P</b> | <i>Aspergillus welwitschiae</i> isolate E16P | <i>Aspergillus welwitschiae</i> strain RCBBR_AEAMCB3 MN603499.1 | PQ634825 | 99.42 |
| <b>E25P</b> | <i>Fusarium proliferatum</i> isolate E25P    | <i>Fusarium proliferatum</i> strain GMBCC1191 PQ573371.1        | PQ634824 | 92.13 |
| <b>E26P</b> | <i>Fusarium solani</i><br>isolate E26P       | <i>Fusarium solani</i> isolate F5 OR123272.1                    | PQ634827 | 98.60 |
| <b>JPE2</b> | <i>Epicoccum</i> sp.<br>isolate JPE2         | <i>Epicoccum sorghinum</i><br>OR397965.1                        | OR632996 | 100   |

319



**Fig. 2** Phylogenetic tree drawn using the Maximum Likelihood method showing relationship between selected endophytic fungi and similar other sequences deposited in the NCBI Genbank. (Evolutionary analysis were conducted in MEGA vs. 11)

### ***In Vitro* Antagonism of Fungal Endophytes**

Dual culture assays of 42 endophytes against *R. solani* showed varying degrees of inhibition (Fig. 3, Table 2). Maximum inhibition was observed with E16P (79.63%), followed by E25P (63.70%). The lowest inhibition was E36P (15.93%). Furthermore, the representative fungal antagonists (showing  $\geq 40\%$  inhibition) were subjected to microscopic examination of the interaction zone formed during the *in vitro* confrontation assay. Various

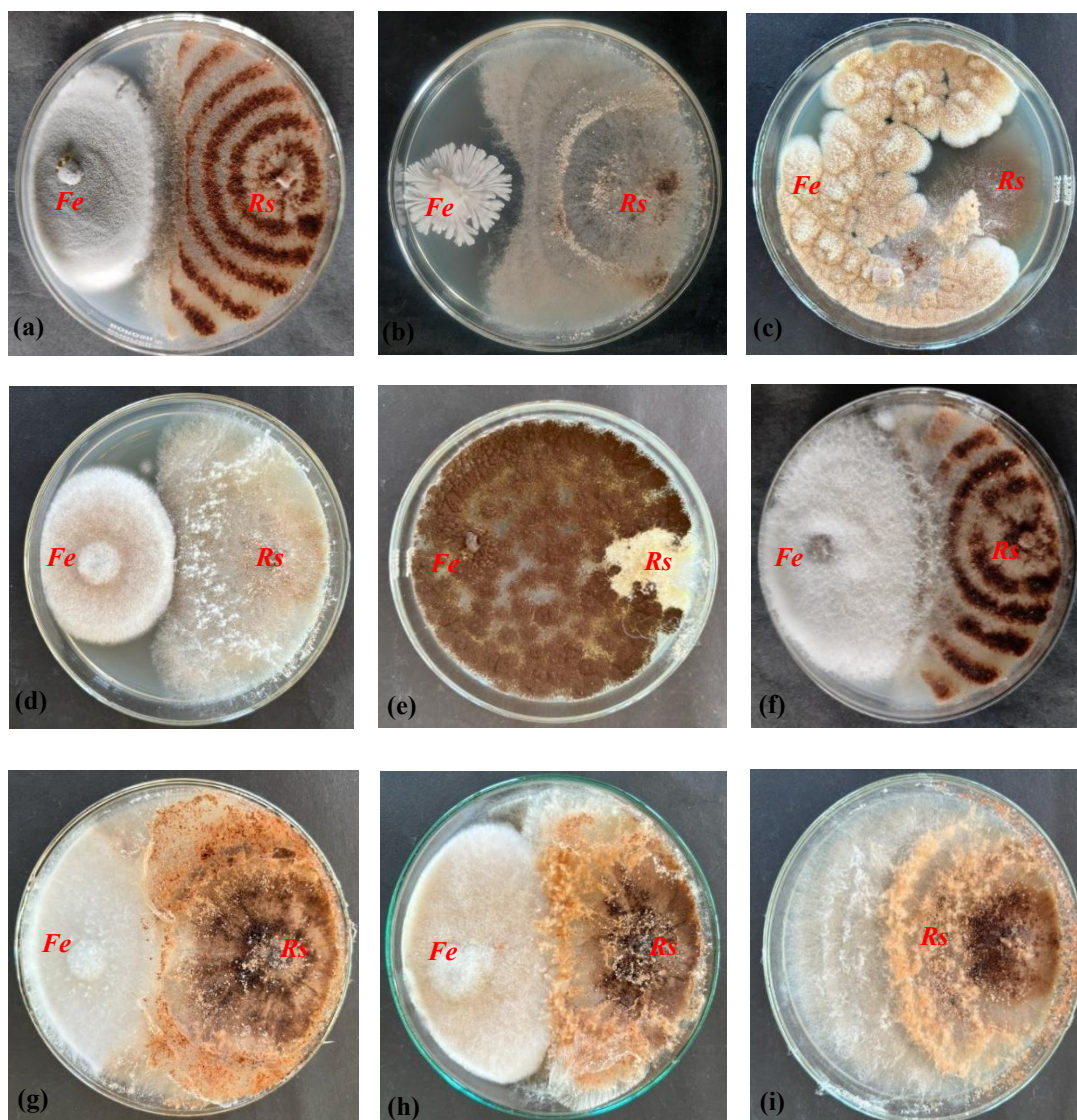
morphological abnormalities and alterations were observed in the mycelium of test pathogen due to its interaction with the fungal endophytes (Supplementary file 1).

**Table 2** *In vitro* antagonism of efficient fungal root endophytes against *Rhizoctonia solani*, causing damping-off of okra

| Isolate                                      | Mycelial growth (mm) | Mycelial growth inhibition (%) * | Changes in mycelial growth at interaction zone                   |
|----------------------------------------------|----------------------|----------------------------------|------------------------------------------------------------------|
| <i>Coniothyrium</i> sp. isolate E1P          | 44.00 <sup>d</sup>   | 51.11 <sup>de</sup><br>(45.62)   | Swelling and deformation of the pathogen mycelium                |
| <i>Xylaria juruensis</i> isolate E4P         | 47.33 <sup>e</sup>   | 47.41 <sup>e</sup><br>(43.50)    | Abnormalities and penetration in the pathogen hyphae             |
| <i>Aspergillus terreus</i> isolate E10P      | 40.33 <sup>c</sup>   | 55.19 <sup>cd</sup><br>(47.96)   | Penetration of the endophytic conidia inside the pathogen hyphae |
| <i>Epicoccum</i> sp. isolate E13P            | 42.67 <sup>cd</sup>  | 52.59 <sup>d</sup><br>(46.47)    | Swelling and deformity of the pathogen mycelium                  |
| <i>Aspergillus welwitschiae</i> isolate E16P | 18.33 <sup>a</sup>   | 79.63 <sup>a</sup><br>(63.18)    | Breakage and encircling over the pathogen hyphae                 |
| <i>Fusarium proliferatum</i> isolate E25P    | 32.67 <sup>b</sup>   | 63.70 <sup>b</sup><br>(52.94)    | Distortion and penetration in the pathogen mycelium              |
| <i>Fusarium solani</i> isolate E26P          | 51.33 <sup>f</sup>   | 42.96 <sup>f</sup><br>(40.93)    | Penetration in the pathogen mycelium                             |
| <i>Epicoccum</i> sp. isolate JPE2            | 37.67 <sup>bc</sup>  | 58.15 <sup>c</sup><br>(49.68)    | Bending and twisting of the pathogen mycelium                    |
| Control                                      | 90.00 <sup>g</sup>   | 0.00 <sup>g</sup>                |                                                                  |
| CD (p=0.05)                                  | 4.54                 | 3.00                             |                                                                  |
| SE(m)                                        | 0.913                | 1.688                            |                                                                  |
| SE(d)                                        | 1.291                | 2.387                            |                                                                  |
| C.V.                                         | 3.740                | 5.837                            |                                                                  |

\*Figures in parentheses are arc sine transformed value

Figures denoted by same letter do not differ significantly

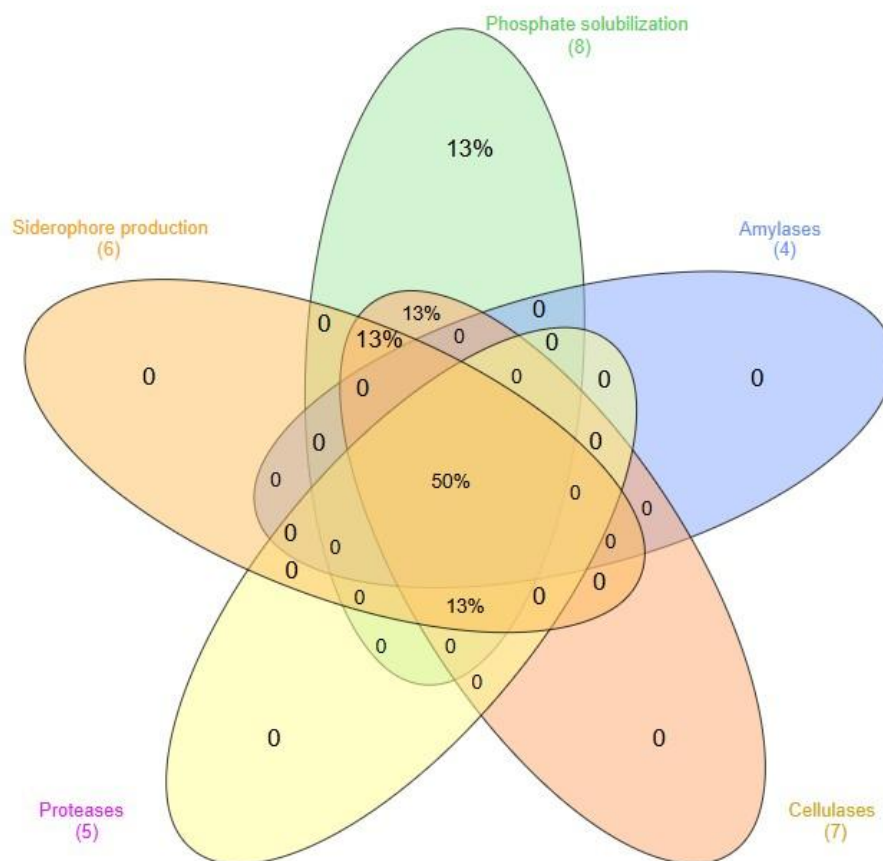


**Fig. 3** *In vitro* confrontation activity of endophytic fungi [a. *Coniothyrium* sp. isolate E1P, b. *Xylaria juruensis* isolate E4P, c. *Aspergillus terreus* isolate E10P, d. *Epicoccum* sp. isolate E13P, e. *Aspergillus welwitschiae* isolate E16P, f. *Fusarium proliferatum* isolate E25P, g. *Fusarium solani* isolate E26P, h. *Epicoccum* sp. isolate JPE2, i. Control] against *Rhizoctonia solani* isolate JPO1, inciting okra damping off (Left side-Fungal endophyte (Fe); Right side-*Rhizoctonia solani* (Rs))

### Plant Growth-Promoting Traits, Stress Tolerance, and Phytochemical Profiling

Selected endophytes exhibited variable PGP traits: all isolates solubilized phosphate, with *F. solani* E26P showing maximum activity (29.67 mm). Siderophore production ranged 7.67–18.33 mm; HCN production was observed in 62.5% of isolates. Lytic enzyme activities among various isolates varied: cellulase (87.5%), protease (62.5%), amylase (50%). Notably, 50% of the isolates exhibited both PGP traits and lytic enzyme activities (Figs. 3–4). Stress tolerance assays indicated growth at pH 2–8, NaCl 1–5% (maximum at 1%), and temperature up to 40°C for most

isolates. *E. jruensis* E4P and *F. solani* E26P showed reduced tolerance at extreme pH/salt. From the results obtained on stress tolerance assay (Table 3), it is inferred that all representative fungal antagonists were able to withstand temperature and salinity stress which could enhance the plant growth under drastic climatic conditions. Phytochemical screening of crude extracts revealed alkaloids, phenols, tannins, triterpenoids, saponins, and flavonoids. *A. terreus* E10P and *A. welwitschiae* E16P contained all tested metabolites; other isolates varied in compound profiles (Fig. 5).



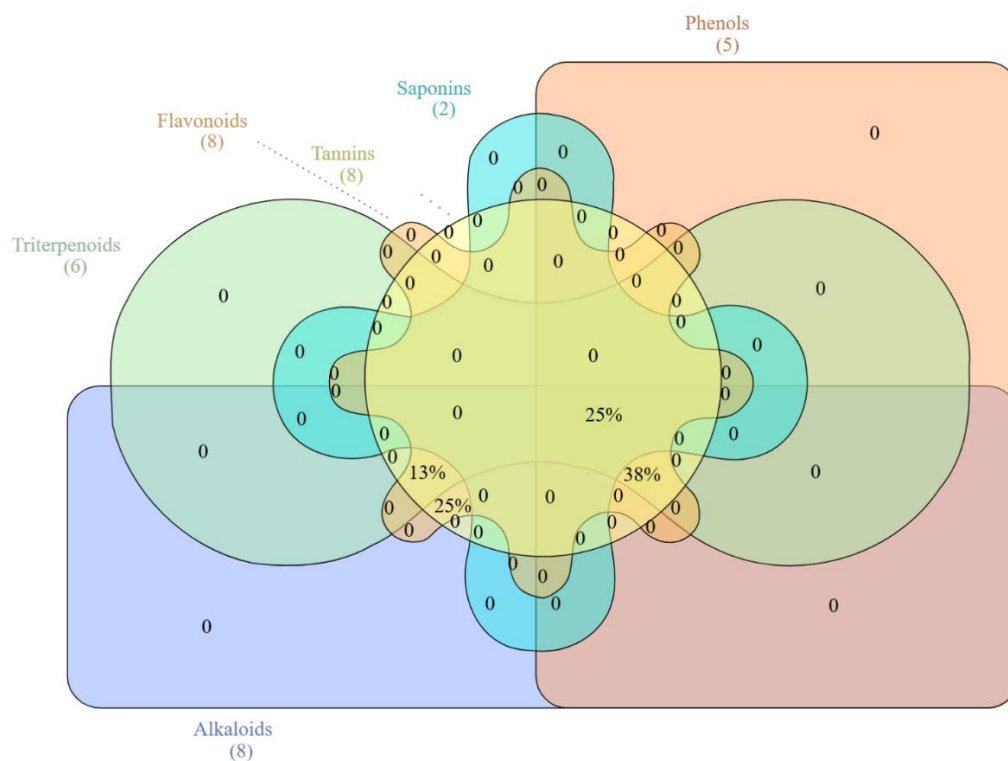
**Fig. 4** Venn diagram representing various plant growth promoting (PGP) traits and lytic enzymes activity of selected endophytic fungi of okra

**Table 3** *In vitro* screening of selected fungal root endophytes for stress tolerance traits

| Isolate                                  | HCN<br>production | Stress Tolerances |   |   |          |   |   |                  |    |    |
|------------------------------------------|-------------------|-------------------|---|---|----------|---|---|------------------|----|----|
|                                          |                   | pH                |   |   | NaCl (%) |   |   | Temperature (°C) |    |    |
|                                          |                   | 2                 | 5 | 8 | 1        | 3 | 5 | 4                | 10 | 40 |
| <i>Coniothyrium</i> sp. -<br>isolate E1P | -                 | +                 | + | + | ++       | + | + | +                | +  | -  |
| <i>Xylaria jruensis</i> -<br>isolate E4P | -                 | -                 | + | - | +        | + | - | +                | +  | -  |

|                                              |   |  |    |     |    |     |    |   |  |   |    |   |
|----------------------------------------------|---|--|----|-----|----|-----|----|---|--|---|----|---|
| <i>Aspergillus terreus</i> isolate E10P      | + |  | ++ | +++ | ++ | +++ | ++ | + |  | + | +  | + |
| <i>Epicoccum</i> sp. isolate E13P            | + |  | +  | ++  | +  | ++  | +  | + |  | - | +  | + |
| <i>Aspergillus welwitschiae</i> isolate E16P | + |  | ++ | +++ | ++ | +++ | ++ | + |  | + | ++ | + |
| <i>Fusarium proliferatum</i> isolate E25P    | + |  | +  | ++  | ++ | ++  | +  | + |  | + | +  | + |
| <i>Fusarium solani</i> isolate E26P          | - |  | +  | ++  | ++ | ++  | +  | - |  | - | +  | - |
| <i>Epicoccum</i> sp. isolate JPE2            | + |  | +  | ++  | +  | ++  | +  | + |  | - | +  | + |
| <b>CD</b>                                    |   |  |    |     |    |     |    |   |  |   |    |   |
| <b>(p=0.05)</b>                              |   |  |    |     |    |     |    |   |  |   |    |   |
| SE(m)                                        |   |  |    |     |    |     |    |   |  |   |    |   |
| SE(d)                                        |   |  |    |     |    |     |    |   |  |   |    |   |
| C.V.                                         |   |  |    |     |    |     |    |   |  |   |    |   |

\*-=No activity; +=Low activity; +=Moderate activity; +++=High activity



**Fig. 5** Venn diagram illustration of distribution of various phytochemicals/secondary metabolite present in crude extracts of selected endophytic fungi of okra

### Biocontrol Potential of Fungal Endophytes Against Damping-Off Under Glasshouse Conditions

The eight most promising fungal endophytes identified *in vitro* were evaluated for their effectiveness against *Rhizoctonia solani* under glasshouse conditions (Supplementary file 2). All isolates significantly reduced disease incidence compared to the untreated control.

**Pre- and post-emergence damping-off:** *Fusarium proliferatum* E25P exhibited the lowest disease incidence (16.66% and 11.66%, respectively) at 10 days' post-inoculation. For pre-emergence damping-off, E25P was statistically similar to *Coniothyrium* sp. E1P (20.00%) and *Epicoccum* sp. E13P (21.66%). For post-emergence damping-off, E13P (16.66%) was comparable with E1P (18.33%).

**Disease control:** Maximum disease suppression was observed with *Fusarium proliferatum* E25P (76.2% pre-emergence, 82.06% post-emergence), followed by *Coniothyrium* sp. E1P (71.43% pre-emergence) and *Epicoccum* sp. E13P (74.37% post-emergence). *Fusarium solani* E26P was the least effective (35.72% and 41.03%, respectively) (Table 4).

**Table 4** Efficacy of fungal endophytic formulations against okra damping-off caused by *Rhizoctonia solani* under glass house conditions

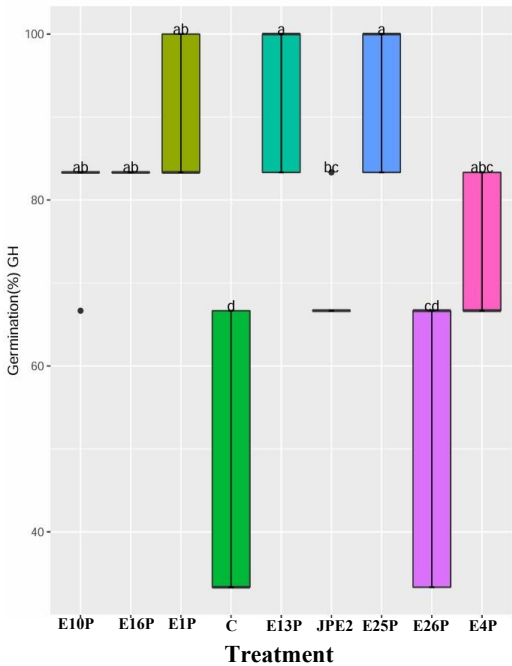
| Treatment                                    | Per cent incidence of damping -off disease |                     |                               |                     |
|----------------------------------------------|--------------------------------------------|---------------------|-------------------------------|---------------------|
|                                              | Pre emergence (%)*                         | Disease control (%) | Post emergence (%)*           | Disease control (%) |
| <i>Coniothyrium</i> sp. isolate E1P          | 20.00 <sup>a</sup><br>(26.44)              | 71.43               | 18.33 <sup>b</sup><br>(25.26) | 71.80               |
| <i>Xylaria juruensis</i> isolate E4P         | 25.00 <sup>ab</sup><br>(29.73)             | 64.29               | 20.00 <sup>b</sup><br>(26.44) | 69.23               |
| <i>Aspergillus terreus</i> isolate E10P      | 33.33 <sup>c</sup><br>(35.25)              | 52.38               | 28.33 <sup>c</sup><br>(32.09) | 56.41               |
| <i>Epicoccum</i> sp. isolate E13P            | 21.66 <sup>a</sup><br>(27.63)              | 69.05               | 16.66 <sup>b</sup><br>(24.08) | 74.37               |
| <i>Aspergillus welwitschiae</i> isolate E16P | 26.67 <sup>b</sup><br>(31.04)              | 61.90               | 25.00 <sup>c</sup><br>(29.99) | 61.53               |
| <i>Fusarium proliferatum</i> isolate E25P    | 16.66 <sup>a</sup><br>(24.08)              | 76.20               | 11.66 <sup>a</sup><br>(19.69) | 82.06               |
| <i>Fusarium solani</i> isolate E26P          | 45.00 <sup>d</sup><br>(42.10)              | 35.72               | 38.33 <sup>d</sup><br>(38.21) | 41.03               |
| <i>Epicoccum</i> sp. isolate JPE2            | 36.66 <sup>c</sup><br>(37.22)              | 47.62               | 26.67 <sup>c</sup><br>(31.04) | 58.97               |
| Control                                      | 70.00 <sup>c</sup><br>(56.82)              |                     | 64.99 <sup>c</sup><br>(53.72) |                     |
| CD (p=0.05)                                  | 3.85                                       |                     | 3.31                          |                     |
| SE(m)                                        | 1.33                                       |                     | 1.15                          |                     |
| SE(d)                                        | 1.89                                       |                     | 1.63                          |                     |
| C.V.                                         | 8.66                                       |                     | 8.25                          |                     |

\*Figures in parentheses are arc sine transformed value

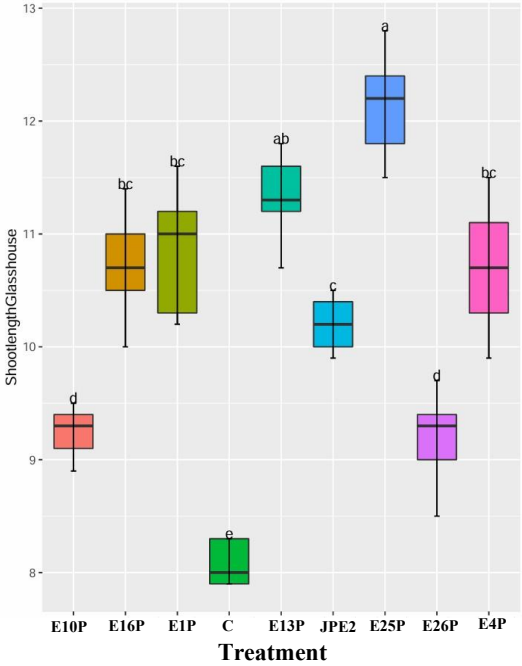
Figures denoted by same letter do not differ significantly

# Evaluation of Endophytic Fungi on Plant Growth Parameters Under Glasshouse Conditions

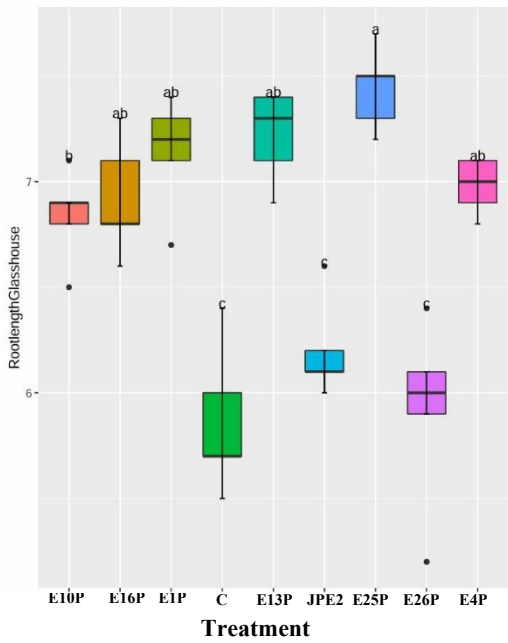
Eight promising fungal endophytes were evaluated under glasshouse conditions for their plant growth–promoting potential in okra. Endophyte-treated seeds showed significant improvements in germination, shoot length, root length, and vigour index compared to the control (Fig. 5).



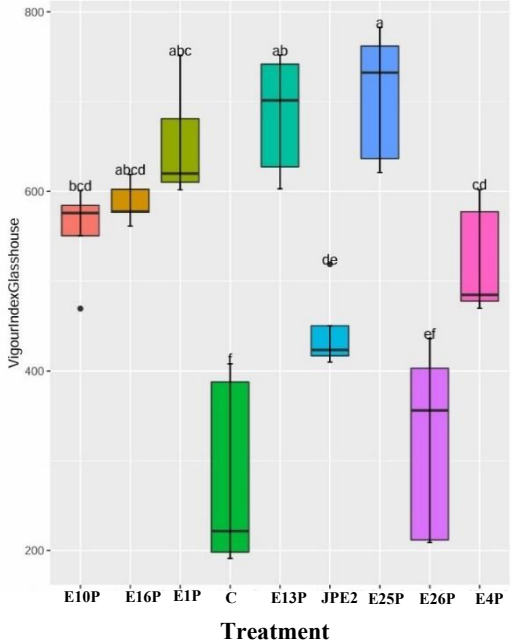
Germination (%)



Shoot length (cm)



Root length (cm)



Vigour Index

**Fig. 5** Effect of fungal endophytic formulations on plant growth parameters of okra under glasshouse conditions [E10P: *Aspergillus terreus* isolate, E16P: *Aspergillus welwitschiae* isolate, E1P: *Coniothyrium* sp. isolate, C: Control, E13P: *Epicoccum* sp. isolate, JPE2: *Epicoccum* sp. Isolate, E25P: *Fusarium proliferatum* isolate, E26P: *Fusarium solani* isolate, E4P: *Xylaria juruensis* isolate]

E25P recorded the highest germination (83.33%), shoot length (12.14 cm), root length (7.44 cm), and vigour index (706.80). E1P, E13P, and E10P were statistically similar for germination and early growth promotion. E26P exhibited the lowest growth performance (germination 66.66%, shoot 9.18 cm, root 5.92 cm, vigour index 323.15), statistically comparable to E4P, JPE2, and E16P. Tukey's post hoc test ( $P \leq 0.05$ ) indicated E25P and E13P consistently formed the superior group across parameters, with moderate responses for E1P, E10P, and E16P, while E26P and control ranked lowest. These results highlight the selective and parameter-specific nature of plant–endophyte interactions, identifying E25P and E13P as the most reliable isolates for enhancing early seedling growth under controlled conditions.

#### Biocontrol and Growth Promotion Under Field Conditions

The most efficient glasshouse isolates were evaluated under field conditions at the CSKHPKV Experimental Research Farm.

**Damping-off incidence:** E25P minimized pre- and post-emergence damping-off (23.33% and 16.66%), followed by E1P (28.33% pre-emergence) and E13P (30.00% pre-emergence; 18.33% post-emergence). E26P was least effective (30.44% and 39.04%, respectively).

**Disease control:** E25P showed the highest suppression (69.57% pre-emergence, 75.62% post-emergence), followed by E1P (63.04%) and E13P (73.18%) (Table 5).

**Table 5** Efficacy of fungal endophytic formulations against damping-off of okra under field conditions

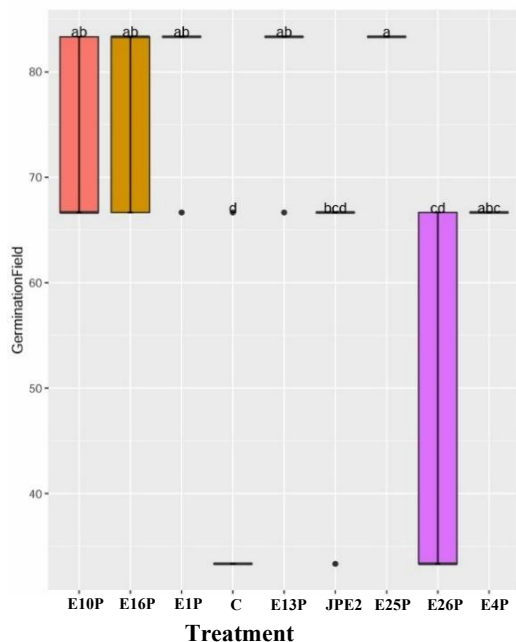
| Treatment                                    | Per cent incidence of damping-off disease |                     |                                 |                     |
|----------------------------------------------|-------------------------------------------|---------------------|---------------------------------|---------------------|
|                                              | Pre emergence (%) <sup>*</sup>            | Disease control (%) | Post emergence (%) <sup>*</sup> | Disease control (%) |
| <i>Coniothyrium</i> sp. isolate E1P          | 28.33 <sup>b</sup><br>(32.09)             | 63.04               | 21.66 <sup>b</sup><br>(27.63)   | 68.29               |
| <i>Xylaria juruensis</i> isolate E4P         | 31.66 <sup>cd</sup><br>(34.20)            | 58.70               | 31.66 <sup>d</sup><br>(34.20)   | 53.66               |
| <i>Aspergillus terreus</i> isolate E10P      | 41.53 <sup>c</sup><br>(40.08)             | 45.83               | 35.00 <sup>c</sup><br>(36.24)   | 48.78               |
| <i>Epicoccum</i> sp. isolate E13P            | 30.00 <sup>bc</sup><br>(33.14)            | 60.87               | 18.33 <sup>ab</sup><br>(25.26)  | 73.18               |
| <i>Aspergillus welwitschiae</i> isolate E16P | 35.00 <sup>d</sup><br>(36.24)             | 54.35               | 26.67 <sup>c</sup><br>(31.04)   | 60.97               |
| <i>Fusarium proliferatum</i> isolate E25P    | 23.33 <sup>a</sup>                        | 69.57               | 16.66 <sup>a</sup>              | 75.62               |

|                                     |                    |       |                     |       |
|-------------------------------------|--------------------|-------|---------------------|-------|
|                                     | (28.81)            |       | (24.08)             |       |
| <i>Fusarium solani</i> isolate E26P | 53.33 <sup>e</sup> | 30.44 | 41.66 <sup>f</sup>  | 39.03 |
|                                     | (46.90)            |       | (40.16)             |       |
| <i>Epicoccum</i> sp. isolate JPE2   | 43.33 <sup>e</sup> | 43.48 | 28.33 <sup>cd</sup> | 58.54 |
|                                     | (41.14)            |       | (32.09)             |       |
|                                     | 76.67 <sup>g</sup> | 0.00  | 68.33 <sup>g</sup>  | 0.00  |
| Control                             | (61.16)            |       | (55.76)             |       |
| CD (p=0.05)                         | 3.47               |       | 3.33                |       |
| SE(m)                               | 1.198              |       | 1.152               |       |
| SE(d)                               | 1.694              |       | 1.629               |       |
| C.V.                                | 6.814              |       | 7.566               |       |

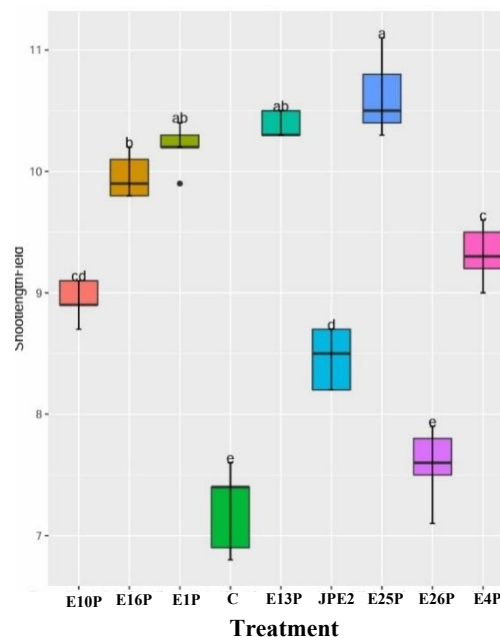
\*Figures in parentheses are arc sine transformed value

Figures denoted by same letter do not differ significantly

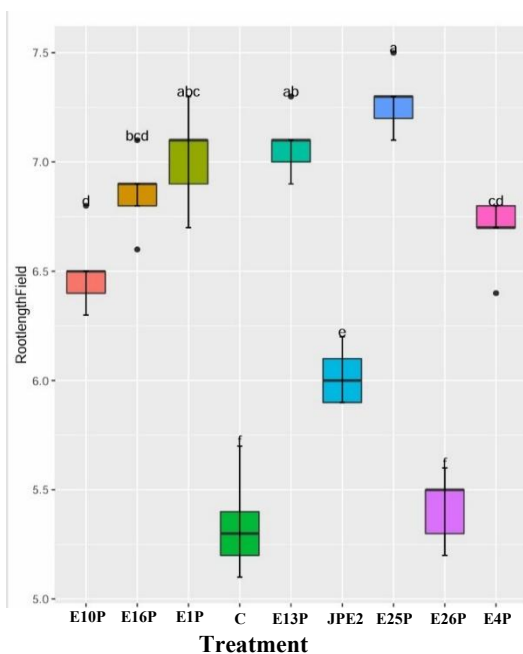
**Plant growth promotion:** Field evaluation demonstrated significant improvements in germination, shoot length, root length, and vigour index (Fig. 6). E25P showed the highest performance (germination 83.33%, shoot 10.62 cm, root 7.28 cm, vigour index 617.26), followed by E13P (80%, 10.38 cm, 7.08 cm, 576.69) and E1P (statistically comparable). Isolates E10P and E16P showed moderate improvements, while E4P, JPE2, E26P, and control exhibited the lowest values. Comparative analysis indicated E25P and E13P maintained superior and stable performance in both glasshouse and field trials.



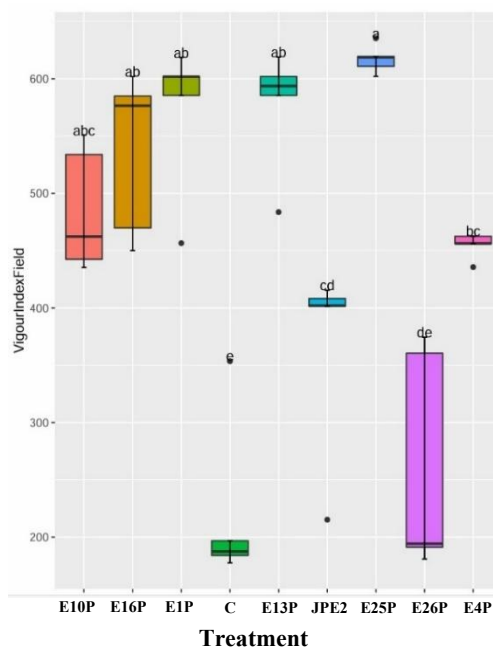
**Germination (%)**



**Shoot length (cm)**



**Root length (cm)**

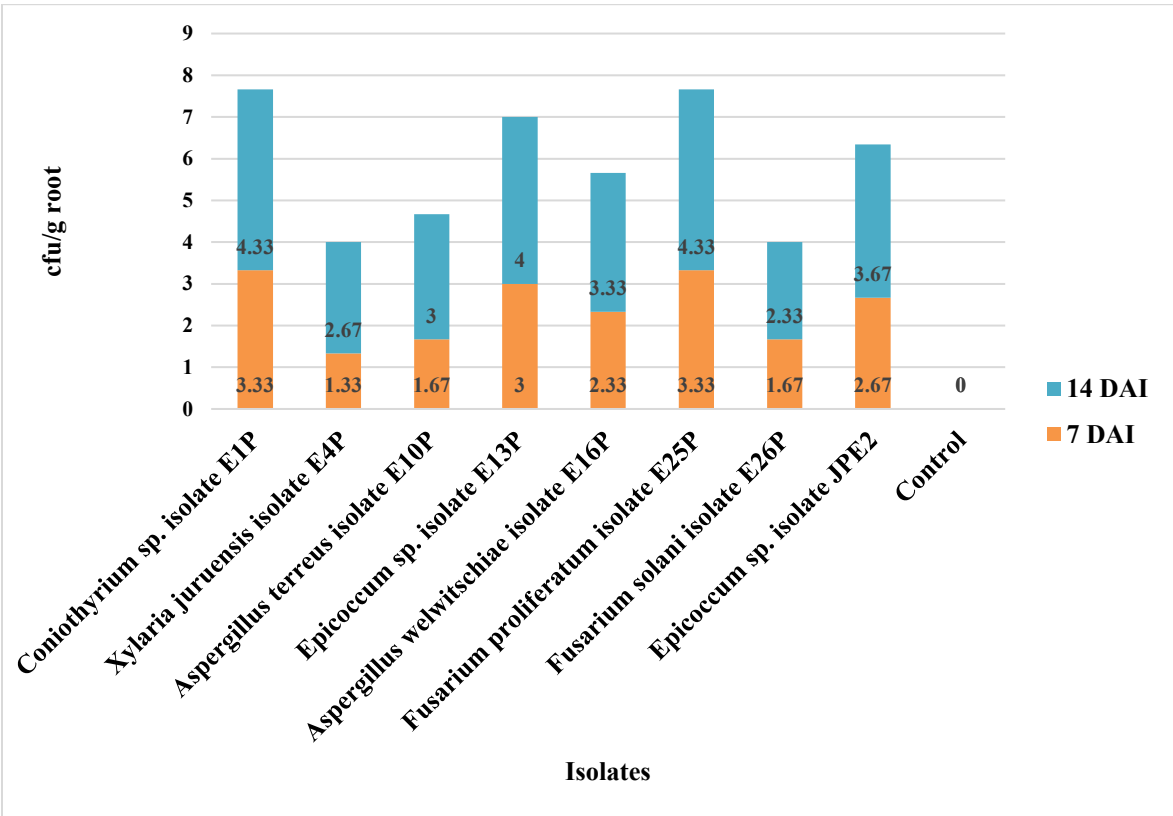


**Vigour Index**

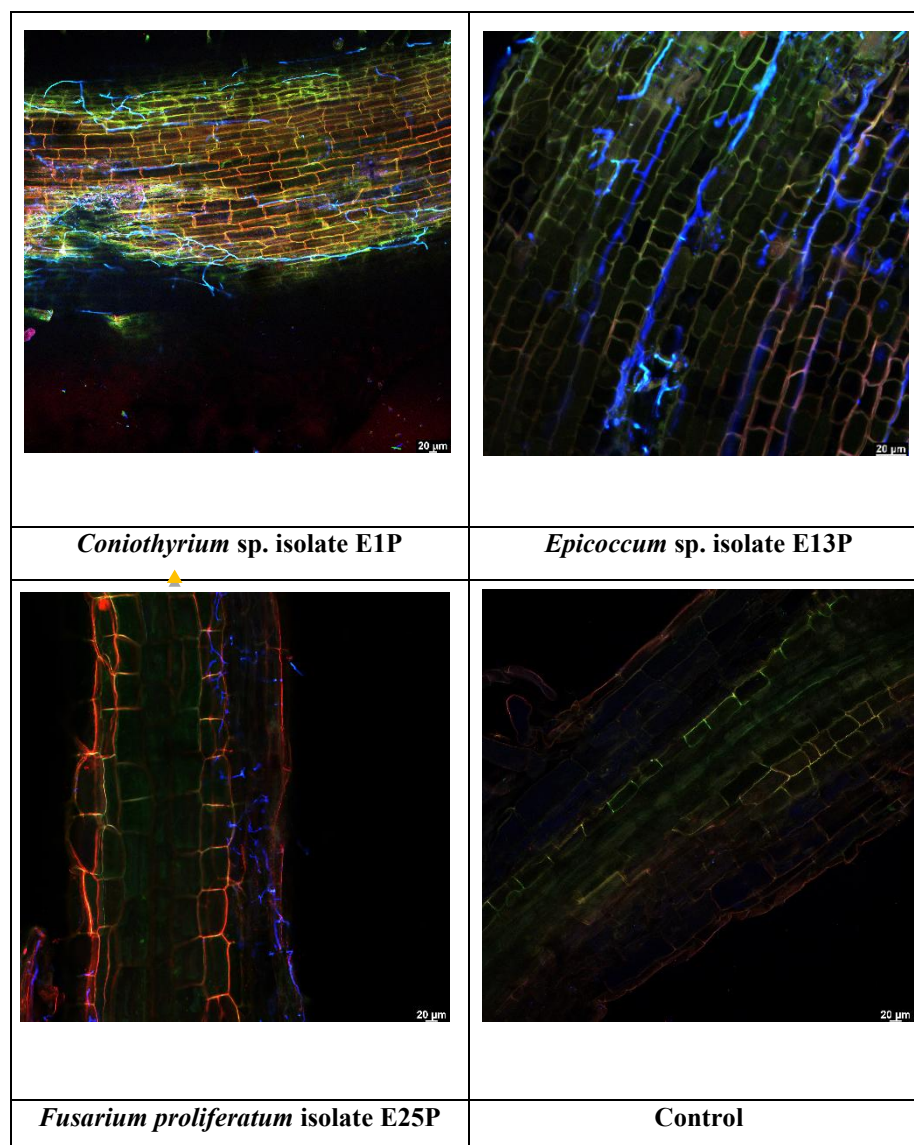
**Fig. 6** Effect of fungal endophytic formulations on plant growth parameters of okra under field conditions [E10P: *Aspergillus terreus* isolate, E16P: *Aspergillus welwitschiae* isolate, E1P: *Coniothyrium* sp. isolate, C: Control, E13P: *Epicoccum* sp. isolate, JPE2: *Epicoccum* sp. Isolate, E25P: *Fusarium proliferatum* isolate, E26P: *Fusarium solani* isolate, E4P: *Xylaria juruensis* isolate]

## Root Colonization by Fungal Endophytes

The colonization potential of selected fungal endophytes in okra roots was assessed using both plate count techniques and confocal laser scanning microscopy (CLSM). Plate counts revealed that all inoculated endophytes successfully established within the root tissues, with significantly higher populations compared to uninoculated controls. Colonization increased over time, being higher at 14 days after inoculation (DAI) than at 7 DAI (Fig.7). Among the tested isolates, *Epicoccum* sp. E13P and *Fusarium proliferatum* E25P exhibited the highest population densities ( $7.67 \times 10^2$  and  $7.33 \times 10^2$  cfu/g, respectively), while *Fusarium solani* E26P showed the lowest colonization ( $2.33 \times 10^2$  cfu/g). Re-isolation confirmed the persistence of these endophytes, with E25P, E1P (*Coniothyrium* sp.), and E13P showing the strongest establishment within root tissues. CLSM analysis further validated these results, revealing that the endophytes predominantly colonized the intercellular spaces of the epidermis and the outer cortex of okra roots. The colonization pattern varied among isolates, with treated roots showing dense and uniform colonization, whereas control roots displayed no fungal presence (Fig. 8). These findings collectively demonstrate that the selected fungal endophytes not only establish effectively in okra roots but also persist over time, supporting their role as potential biocontrol and plant growth-promoting agents.



**Fig. 7** Re-isolation frequency and colonization confirmation of inoculated endophytic fungi by plate count technique



**Fig. 8** Confocal microscopic analysis of colonized and established fungal endophytes. Blue colored fungal hyphae (yellow arrows) showing colonization in the roots of okra (a, b & c). No fungal colonization observed in control images (d). Images were captured at 10X & 20X objective. (Scale bar- 20 μm)

## Discussion

Okra (*Abelmoschus esculentus* L.) is a nutritionally rich and economically important vegetable cultivated widely across tropical and subtropical regions of the world (De Rosa et al. 2010; Doymaz 2005; Gemede 2015; Wu et al. 2022). Its remarkable nutritional profile and adaptability have contributed to its high market demand, particularly during the summer season in regions such as Himachal Pradesh (Tikoo et al. 2025). However, okra productivity is constrained by multiple soil-borne pathogens, among which damping-off disease is a major bottleneck, particularly affecting early crop establishment. The opportunistic fungal pathogen *Rhizoctonia solani* has emerged as a principal causative agent responsible for considerable losses in okra-growing regions of Himachal Pradesh (Sharma 2011; Patel et al. 2019). Damping-off can lead to yield reductions of 10–30%, and under severe conditions may result in

complete crop failure, depending on varietal susceptibility and environmental factors (Patil et al. 2015; Mohankumar et al. 2016; Wu et al. 2022; Mimma et al. 2023). Traditional strategies, including crop rotation, seed treatments, soil fumigation, and chemical fungicide applications, have shown limited success due to the emergence of resistant pathogen strains, environmental contamination, and negative impacts on non-target microflora (Lamichhane et al. 2017; Pandey et al. 2018; Bhattarai et al. 2022; Akber and Fang 2024). Consequently, there is a pressing need for sustainable, eco-friendly strategies for disease management, especially in high-demand summer crops (Sharma et al. 2015; Abdelaziz et al. 2022).

In this context, fungal endophytes represent a promising avenue for integrated disease management. These microorganisms colonize plant tissues internally, establishing mutualistic associations that confer multiple benefits, including disease suppression, growth promotion, and enhanced resilience against abiotic stresses (Attia et al. 2022a, b; Hashem et al. 2023; Pandian et al. 2024; Zotchev 2024). Unlike rhizospheric microbes, endophytes inhabit the same ecological niche as vascular pathogens, experiencing less antagonism and greater stability under biotic and abiotic stresses (Dutta et al. 2014; de Lamo and Takken 2020; Jiajia et al. 2023). Additionally, their ability to produce diverse bioactive secondary metabolites—such as alkaloids, steroids, terpenoids, flavonoids, quinones, and phenols—enhances their biocontrol and plant growth-promoting potential (El Enshasy et al. 2020; Hazra et al. 2024). To the best of our knowledge, the present study provides the first comprehensive report on endophytic fungi-mediated management of okra damping-off, offering a novel, sustainable approach for mitigating crop losses while improving plant growth and resilience.

Field surveys conducted across Kangra, Mandi, and Hamirpur districts revealed widespread incidence of damping-off, ranging from 31.67% to 84.33%, with early-sown okra (April–May) exhibiting greater susceptibility compared to late-sown crops (May–June). These findings underscore the urgency of effective management strategies and are consistent with earlier reports of widespread damping-off of okra in Himachal Pradesh (Jandaik et al. 2015; Patel et al. 2019). A total of 44 fungal isolates were recovered from symptomatic samples, with *R. solani* emerging as the predominant pathogen (16 isolates). Detailed morpho-molecular characterization confirmed the causal agent as *R. solani* isolate JPO1 (GenBank accession MK621284.1), corroborating previous studies from India and Iraq where *R. solani* was identified as the primary pathogen responsible for okra damping-off (Sharma 2011; Jandaik et al. 2015; Patel et al. 2019; Madhi et al. 2020).

Pathogenicity assays conducted on the susceptible okra cultivar ‘*Palam Komal*’ confirmed the virulence of *R. solani*, with characteristic water-soaked sunken lesions appearing seven days’ post-inoculation, followed by wilting and seedling collapse. These observations align with previous reports of damping-off symptoms caused by *R. solani*, *Fusarium spp.*, and *Macrophomina phaseolina* (El-Mohamedy 2004; Sharma 2011; Gondal et al. 2019; Kushwaha et al. 2024).

Quantitative assessment of root endophytic fungi revealed significant spatial variation across surveyed districts, with 41 isolates recovered. *Fusarium* and *Aspergillus spp.* were the most frequent endophytes, consistent with reports indicating their widespread ecological distribution and dominance across diverse host plants (Maciá-Vicente et al.

2008; Fontana et al. 2021; Singh et al. 2021; Gupta et al. 2023; Zhang et al. 2024; Kushwaha et al. 2024). Diversity was influenced by host genotype, age, soil properties, altitude, and geographic location (Arnold 2007; Katoch et al. 2014; Adnan et al. 2018; Jin et al. 2021).

Eight isolates exhibiting  $\geq 40\%$  in vitro antagonism were selected for further characterization. ITS-based molecular analysis validated morpho-cultural identifications: E1P as *Coniothyrium* sp., E4P as *Xylaria juruensis*, E10P as *Aspergillus terreus*, E13P as *Epicoccum* sp., E16P as *Aspergillus welwitschiae*, E25P as *Fusarium proliferatum*, E26P as *Fusarium solani*, and JPE2 as *Epicoccum* sp. Phylogenetic analysis confirmed these identities.

Dual-culture assays demonstrated pronounced antagonistic activity, with microscopic examination revealing marked deformation of pathogen hyphae, likely mediated by endophyte-produced secondary metabolites. This supports the multifunctional biocontrol potential of fungal endophytes through mechanisms including nutrient competition, antibiosis, and mycoparasitism, consistent with earlier reports (Lahlali and Hijri 2010; Shentu et al. 2014; Hadimani and Naik 2018; Pal et al. 2020; Singh et al. 2021; Alves et al. 2021; Kushwaha et al. 2024; Zhang et al. 2024).

PGP assays confirmed the ability of endophytes to solubilize phosphate, produce siderophores and HCN, and express lytic enzymes. Stress tolerance studies revealed resilience to a range of pH, salinity, and temperature conditions, while phytochemical profiling identified alkaloids, phenols, tannins, triterpenoids, saponins, and flavonoids. These properties are associated with enhanced plant growth and pathogen suppression, consistent with prior findings on multifunctional endophytes (Waqas et al. 2015; Ripa et al. 2019; Husna et al. 2021; Gopane et al. 2024; Narayanan et al. 2024).

Glasshouse and field trials confirmed *Fusarium proliferatum* E25P and *Epicoccum* sp. E13P as the most effective isolates in reducing both pre- and post-emergence damping-off, enhancing germination, shoot/root length, and vigour index. *Coniothyrium* sp. E1P demonstrated moderate but consistent performance. Root colonization, assessed through plate counts and confocal microscopy, revealed extensive establishment within epidermal and cortical tissues, supporting both biocontrol and growth promotion functions. These results are in line with studies demonstrating successful colonization and functional impact of endophytes in crops like sorghum, tomato, pea, and legumes (Favaro et al. 2012; Rajini et al. 2020; Ebrahimi et al. 2023; Kushwaha et al. 2024).

Collectively, this study demonstrates that selected endophytic fungi can simultaneously suppress *R. solani*-induced damping-off and enhance early growth and vigour of okra under glasshouse and field conditions. The isolate-specific efficacy and stable performance of E25P and E13P highlight their potential as bioformulations for sustainable okra cultivation. This research not only expands the knowledge base on endophyte-mediated disease management but also provides a practical, environmentally sustainable strategy to mitigate a major constraint in summer okra production.

This study demonstrates the high potential of endophytic fungi as sustainable biocontrol agents against *Rhizoctonia solani*-induced damping-off in okra. Among the 42 isolates screened, *Fusarium proliferatum* E25P and *Epicoccum* sp. E13P consistently exhibited superior antagonistic activity, enhanced seed germination, and improved early

seedling growth under both glasshouse and field conditions. The isolates successfully colonized okra root tissues, as confirmed by plate counts and confocal microscopy, establishing stable and persistent endophytic populations.

In addition to disease suppression, selected endophytes displayed multiple plant growth-promoting traits, stress tolerance, and production of bioactive secondary metabolites, highlighting their multifunctional role in enhancing crop health and resilience. These findings underscore the feasibility of developing fungal endophytic formulations as eco-friendly, effective, and practical alternatives to chemical fungicides for damping-off management in okra, particularly under summer cultivation.

Overall, this research provides novel insights into endophyte-mediated disease control and growth promotion, offering a promising strategy for sustainable okra production while contributing to the broader understanding of plant–endophyte interactions in agroecosystems.

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