ORIGINAL CONTRIBUTION

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Evaluation of bioactivity and phytochemical screening of endophytic fungi isolated from *Ceriops decandra* (Griff.) W. Theob, a mangrove plant in Bangladesh



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Abstract

Background: Due to increasing number of individuals within the world having health issues caused by varied cancers, drug-resistant microorganism, parasitic protozoans, and fungi could be a cause for alarm. Endophyte's are a unique source of medicinal compounds which requires an intensive seek for newer and simpler agents to handle these sickness issues is currently advancing. This study explores the description of endophytic fungi of *Ceriops decandra* (Griff.) W. Theob., a mangrove plant growing in mangrove forest, Sundarban, Bangladesh.

Methods: The endophytic fungi of experimental plants were identified by DNA amplification and sequencing of the internal transcribed spacer (ITS) region Moreover, cytotoxicity, antioxidant and antimicrobial tests were assessed. In addition, the screening of extracts was performed by visual detection using UV light.

Results: Three fungi namely CEDBE-1, CEDLE-6 and CEDLE-10 (internal strain no) were isolated as endophytic fungi and identified as *Fusarium oxysporum*, *Clonostachys* spp. and *Fusarium solani* respectively on the basis of morphological characteristics and DNA sequencing. The isolates were cultured at 28° C±2 for 21 days in Potato Dextrose Agar (PDA) media. The extracts of fungal isolates were examined at 100 μg/disc concentration for the screening of antibacterial activity and exhibited no activity against tested human pathogenic bacteria. The strain CEDBE-1 showed highest antioxidant activity of IC₅₀ value of 31.07 μg/mL comparing with the positive control ascorbic acid (0.74) and BHA (7.79) μg/mL. All isolates exhibited strong cytotoxic activity using brine shrimp lethality bioassay compared with the positive control tamoxifen. Preliminary chemical screening of the fungal extracts by thin layer chromatography technique revealed the presence of various compounds.

Conclusions: The findings indicate that endophytic fungal isolates explored from *C. decandra* has diversified biological activities which may be potential source of leads for new drug development.

Keywords: Ceriops decandra (Griff.) W. Theob, endophytic funqi, molecular identification, phylogenetic analysis

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Introduction

Sundarban is a leading single source of mangrove forest in the world [1]. More than a number of its plant species have massive applications within the convention of folk medicine; numerous components of those plants are applied for the native folks as cure for numerous ailments [2-4]. Yet such colossal potential, few reports are available on these species relating to their biological activities and therefore, the active principles liable to such activities. An endeavor has been created during this analysis to extend the notice for the therapeutic significance as conservation and utilization of those mangrove species as natural affluent sources of novel bioactive agents. As mangroves thrive in enormously traumatic and unfriendly environment of high salinity, high and low tides of water, heat and humidity, sturdy winds and muddy anaerobic soil that the group of plants ought to have extremely developed morphological and physiological variations to such extreme conditions. These exceptional characteristics may result in the synthesis of novel chemical compounds; these chemical compounds recommend protection to those plants against numerous biotic and abiotic stresses [5]. Endophytic microflora resides in the host plants while not inflicting any ailment symptoms. Solely a number of plants are studied for their endophyte variety and their potential to provoke bioactive secondary metabolites [6]. Approximately, all the plants are well- known to harbor for endophytes. The selection of the plant may be the resource of secondary metabolites [7]. Exploration of endophytes and their bioactivities is very important because the endophytic fungi have a very important relationship in some co-existing endophytes and their host plants, which might considerably influence the formation of metabolites in plants, then have an effect on quality and extent of crude medicine derived from healing plants over the long era of evolution [8].

C. decandra is a mangrove plant of tropical Asia belonging to the family Rhizophoraceae. It is grown as a shrub or small tree up to 15 m (50 ft) tall with a trunk diameter of up to 30 cm (12 in) found naturally in India and Bangladesh (including the Sundarbans), Burma, Thailand and Peninsular Malaysia [9]. This plant has strong antioxidant activity.

Furthermore, this plant has reputation for its folk-loric uses in the treatment of gastrointestinal disorders, infection, snakebites, inflammation, and cancer. Various phytoconstituents which include diterpenoids (ceriopsin A-G), triterpenoids (lupeol, α -amyrin, oleanolic acid, ursolic acid), and phenolics (catechin, procyanidins) are present in different parts of the plant. These phytoconstituents and their derivatives could form a new basis for developing new drugs against various diseases [10].

Profuse use of profitable antibiotic and manufactured pesticides for human and crop protection is detrimental to human health, ecology and atmosphere. Attention has additionally been drawn to the antimicrobial properties of plants and their metabolites owing to the growing incidences of drug-resistant pathogens of each clinical and agricultural importance [11]. Thus, this research was carried out to study the bioactivity of endophytic fungi associated with *C. decandra* so that it could be a potential source leading to the development of new drug.

Materials and methods

Isolation of fungal material

The plant samples were accumulated from conservation forest of Sundarban, Bangladesh in February, 2018. The plant material was recognized and authenticated by a taxonomist of Bangladesh National Herbarium (BNH), Dhaka, Bangladesh. A voucher specimen of this compilation was retained at BNH under the accession number DACB – 45,945. Fungal strains were isolated from the fresh plant parts following the procedure described by Khan et al. 2018 [12, 13].

Taxonomical identification of fungal cultures Morphological identification

Slides were prepared from cultures by staining with lactophenol cotton blue examined with a bright-field and phase contrast microscope for the identification of endophytic fungal strains (Sadananda, Govindappa et al., 2014). Likewise, the standard taxonomic keys such as growth pattern, hyphae, the color of the colony and medium, margin character, aerial mycelium, surface, texture, sporulation and production of acervuli, the size and coloration of the conidia were used to identify morphological characteristics [14, 15].

Molecular identification

Fungal strains were identified by DNA amplification and sequencing of the internal transcribed spacer (ITS) region [16]. A section of fungal hyphae (0.5-1.0 cm²) was assembled from the Petridish and lyophilized in an Eppendorf tube (2 mL, Eppendorf, Germany). With the help of glass beads the lyophilized fungal mycelia were pulverized and agitated. In keeping with the manufacturer's protocol, fungal DNA isolation was acquired by using DNeasy Plant Mini Kit (QIAgen, USA). Cell lysis, digestion of RNA by RNase A, removal of precipitates and cell debris, DNA shearing; precipitation and purification were assimilated in the method. The isolated DNA was then amplified by polymerase chain reaction (PCR). The PCR was conceded utilizing HotStarTaq Master Mix Kit (QIAgen, USA). ITS 4 (with base se-TCCGTAGGTGAACCTGCGG) 5(with base sequences TCCTCCGCTTATTGATATGC)

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(Invitrogen, USA), as primers, were mixed with HotStar-Taq Master Mix Kit and DNA template in a total volume of 50 µL. The mixture was then applied to the thermal cycler (BioRad, USA) using the programmed PCR cycle as outlined below: Initial activation step in 95 °C for 15 min to activate HotStarTaq DNA polymerase, cycling steps which were restated 35 times, denaturing: 1 min at 95 °C, annealing: 1 min at 56 °C, extension: 1 min at 72 °C, final extension for 10 min in 72 °C. The PCR product was purified using 2% Agarose-Gel-Electrophoresis at 75 V for 60 min in 1 X TBE buffer. The agarose gel was then stained using 1% ethidium bromide. An approximate size 550 bp stained DNA fragment was then wiped out from the agarose gel. By following manufacturer's protocol, **PCR** purification was carried out using Perfect Prep Gel Cleanup Kit (Eppendorf, USA). The amplified fungal DNA (PCR product) was then submitted for sequencing by a viable service and the base sequence was evaluated with publicly accessible databases such as GenBank with the help of Blast-Algorithms.

Bioactivity screenin

Brine shrimp lethality bioassay

According to Meyer et al. Brine shrimp lethality bioassay was implemented to inspect the cytotoxicity of the extract [17]. Serial dilution was then carried out in order to obtain the concentration of 1.25 µg/mL to 320 µg/ mL. 5 mL of artificial sea water was added into all the test tubes. Artificial seawater (prepared by using sea salt 38 g/L and adjusted to pH 8.5 using 1 N NaOH) was used to hatch the eggs of the brine shrimps (Artemia salina) under constant aeration for 24 h under the light. To get shrimp larvae, the hatched shrimps were permitted to grow by 48 h. Suspension containing 10 nauplii was added into each test tube and was incubated at room temperature under the light. The tubes were then examined after 24 h and the number of surviving larvae in each test tube was counted with the aid of a magnifying glass. The percentage of mortality was plotted against the logarithm of concentration. The concentration that would kill 50 % of the nauplii (LC₅₀) was determined from probit analysis as well as linear regression equation using the software "Microsoft Excel-2007". Tamoxifen was used as standard in this bioassay [18].

Antimicrobial screening

The antimicrobial and antifungal activities were investigated by the procedure explained by Bauer et al. [19]. The test microorganisms used in the antimicrobial study incorporated four pathogenic bacterial strains *Staphylococcus aureus* (ATCC 25,923), *Escherichia coli* (ATCC 28,739), *Bacillus megaterium* (ATCC 259), *Salmonella typhi* (ATCC 28) and one fungal strains *Aspergillus*

niger. After 18–24 h of incubation at 37 °C for bacteria and 48–96 h of incubation at 28 °C for fungi, the zones of growth inhibition around the discs were measured. By determining the diameter of inhibitory zones in millimeter compared to kanamycin (30 $\mu g/disc$) as a positive control and ketoconazole (30 $\mu g/disc$) as standard antibiotics for antibacterial and antifungal screening, correspondingly the sensitivities of the microorganism species to the fungal extract were calculated.

Initial screening by Thin Layer Chromatography (TLC)

This screening of the extract was performed by visual detection, UV light (in short and long wavelength) and vanillin- H_2SO_4 spray reagent.

Chemical screening

For identification of chemical constituents of endophytic fungal crude extracts the extracts were subjected to thin layer chromatographic technique. Commercially available precoated silica gel (Kiesele gel 60 PF 254) plates were used with solvent system of toluene: 10 % ethyl acetate for the initial chemical screening of the crude extracts.

Statistical analysis

 IC_{50} values between groups were compared using independent student's t-tests. For the analysis of the cytotoxicity results for each extract, continuous variables between groups were compared with one-way analysis of variance (ANOVA) with post hoc Tukey's test. Mean values between groups were compared using independent student t-test for equality of variances. Statistical significance was accepted when P < 0.001.

Results

Isolation and morphological grouping of endophytic fungi from C. decandra

In this investigation, a total 30 segments (14 leaves and 16 barks) of *C. decandra* were inoculated in the plate and a total 25 isolates were purified on the basis of morphological characteristics using standard protocol [15]. These fungal isolates were successfully identified up to genus level as two species Fusarium and one species Clonostachys. They exhibited characteristic colony and microscopic morphology that could be used to differentiate them. The colonies of Fusarium appeared yellowish or dull white in color (Fig. 1A & B). In microscopy, Fusarium spores exhibited 1-celled, cylindrical, 6-15 × 3-4 μm in size, conidia was elliptical to cylindrical, with 0 to 2 septa curved (Fig. 1C). Acervuli was in groups, sometimes appeared like circular bodies emerging from yellowish white spots. Conidial mass was yellow in the middle and the rest portion was white. Mycelium was also yellowish white (Fig. 2). Strain no CEDLE-6 was Munshi et al. Clinical Phytoscience (2021) 7:81 Page 4 of 10

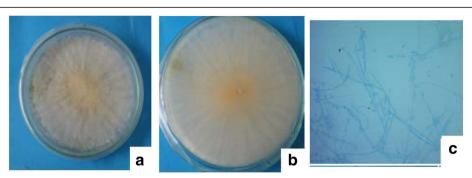


Fig. 1 Macroscopic front (A) macroscopic (B) and microscopic (C) view of Fusarium sp. The pictures (A, B) represent macroscopic view of Fusarium sp. from leaf of C. decandra after 12 days of cultivation on PDA media and also represent microscopic view (40X) (C)

presumed the new strain of Clonostachys sp. as the BLAST results showed 91.58 % similarity between Clonostachys sp. and Clonostachys rosea. Some of the differences were observed between Clonostachys sp. and C. rosea. Colonies of Clonostachys sp. were irregularly cottony and usually gray to black where the dark brown mature conidia eventually formed, whereas the colonies of C. rosea was grayish green to olivaceous green and the conidial mass was also grayish to olivaceous green (Fig. 3). Moreover, Conidia were $3-5 \times 5-10 \mu m$ in size, rough; fairly thin walled, oblong and usually two cells in studied Clonostachys sp. in as much as conidiophores of C. rosea were $25-200 \times 3.5-5.5 \mu m$ in size. Furthermore, the conidiophores of C. rosea were verticillium-like, while the conidia of Clonostachys sp. were spindle-like [18]. In addition, hyphae were septate, phialides were erect, unbranched, tapering and form directly on the fine, narrow hyphae in case of Clonostachys sp.

Molecular identification and phylogenetic analysis of isolates

Molecular characterization of fungal endophytes provides evidence about the separation of morphologically identical fungi which may vary enough genetically to be considered separate species, and in some cases to be placed into different clades. Based on BLAST analysis, the sequence likeness considered 95–100%. Three

species were identified as representing two genera which is shown in Table 1.

The generic name of fungal strain CEDBE-1 was analyzed on the basis of 5.8 S-rRNA-ITS regions and examined the phylogenetic relationship with the published data. As seen in Fig. 4 fungal isolates CEDBE-1exhibited 100% similarity to other fungal isolates *F. oxysporum* (accession no. MK828120.1) that itself was identified as *F. oxysporum*, similarly to other species (100%, accession numbers LT841236.1, LT841222.1 etc.) deposited in the U.S. National Center for Biotechnology Information (NCBI). Finally, CEDBE-1 was identified as *F. oxysporum* by comparing with the morphological key features.

Next fungal strain CEDLE-6 was presumed to be *Chlonostachys* sp. based on the phylogenetic relationship for 5.8SRNA-ITS regions as shown in Fig. 5 and comparison of the 5.8SRNA-ITS regions with available sequences in the database, namely the base sequences which showed 98 % similarities with *Hypomyces corticolia* (accession no MH860037.1) and 91.58 % similarities with *C. rosea* (accession no MH862790.1). On the basis of morphological characteristics finally it was concluded that strain CEDLE-6 *Chlonostachys* sp. registered as KY320599.1.

Based on the analysis of 5.8 S-rRNA-ITS regions of CEDLE-10 this fungal strain showed 100% similarity with the genus *F. solani* (accession no. KC577186.1)

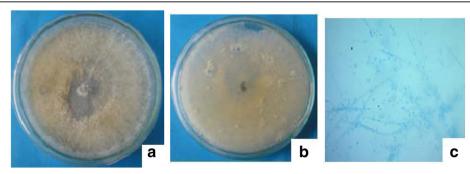


Fig. 2 Macroscopic front (**A**) macroscopic (**B**) and microscopic (**C**) view of *Fusarium* sp. The pictures (**A, B**) represent macroscopic view of *Fusarium* sp. from bark of *C. decandra* after 12 days of cultivation on PDA media and also represent microscopic view (40X) (**C**)

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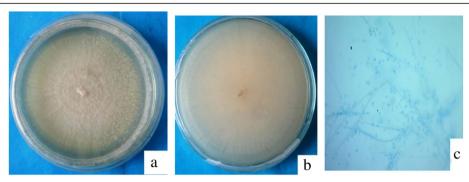


Fig. 3 Macroscopic front (A) macroscopic (B) and microscopic (C) view of Clonostachys sp. The pictures (A, B) represent macroscopic view of Clonostachys sp. from leaf of C. decandra after 12days of cultivation on PDA media and also represent microscopic view (40X) (C)

where other texa *F.solani* (accession no. AB190389.1) showed very close phylogenetic relationship with the *F. solani* which is shown in Fig. 6. By comparison with the available sequences in the database and morphological keynotes it has been clarified CEDLE-10 as *F. solani*.

Bio-screening of isolated fungal isolates Cytotoxicity test

Extracts of the plant *C. decandra* and its associated endophytic fungal strains CEDBE-1, CEDLE-6 and CEDLE-10 were screened for probable cytotoxic activities using brine shrimp lethality bioassay and the result is represented as a bar graph in Fig. 7. The LC₅₀ value of crude methanol extract of the plant was found to be 8.67 μ g/mL. On the other hand, extracts of the fungal strains CEDBE-1, CEDLE-6 and CEDLE-10 were found to be 0.87 μ g/mL, 1.03 μ g/mL and 2.68 μ g/mL respectively. It is noticed that all the test samples were lethal to brine shrimp nauplii. By contrasting these values with tamoxifen as positive control, it was noticed that crude fungal extracts had remarkable cytotoxicity to brine shrimp.

Antioxidant test

The free radical scavenging activity (Fig. 8) of different extracts from C. decandra and its associated endophytic fungal strains were assayed by using DPPH scavenging method (Dame et al., 2016). In this assay, fungal strain CEDBE-1 showed highest antioxidant activity with IC₅₀ value of 31.07 μ g/mL compared with IC₅₀ values of

positive control 0.74, 7.79 for ascorbic acid and BHA respectively whereas CEDLE-6 and CEDLE-10 exhibited moderate antioxidant activity with IC_{50} values of 88.22 and 141.86 µg/mL, respectively. These results indicate that CEDBE-1 has immense antioxidant capacity which may have potentiality to release antioxidant metabolites.

Antimicrobial test

All isolates were screened for antimicrobial activity adjacent to a wide variety of human pathogenic bacteria and fungi. The results of the antimicrobial tests of ethyl acetate extracts of laboratory culture of three endophytic fungi are listed in Table 2. Isolate CEDLE-10 exhibited potential antimicrobial activity against all tested human pathogenic bacteria except *S. aureus*. Other isolates CEDBE-1 and CEDLE-6 showed moderate activity against only *S. typhi*.

Chemical screening

All the extracts qualitatively tested for presence of chemical constituents. The screening of extracts was performed by visual detection, UV light (both in short at 254nm and in long at 365nm) and all of these results are presented in Table 3. Thin layer chromatographic technique was used in solvent system (Toluene: Ethyl acetate = 90 %: 10 %) to identify the presence of different secondary metabolites. This result indicates the presence of a good number of compounds like flavonoids, terpenoids, carotenoids, anthocyanins etc.

Table 1 Molecular identification using ITS regions of endophytic fungi isolated from Ceriops decandra plant

Fungal Species	Max	Total	Query	E-Value	%	Accession No.
	·	Score	Cover		Similarity	
Fusarium oxysporum	1055	1055	100 %	0	100.00 %	MK828120.1
				0.0		
Clonostachys sp.	760	760	98 %	0.0	91.58 %	KY320599.1
Fusarium solani	1020	1020	99 %	0.0	100.00 %	KC577186.1

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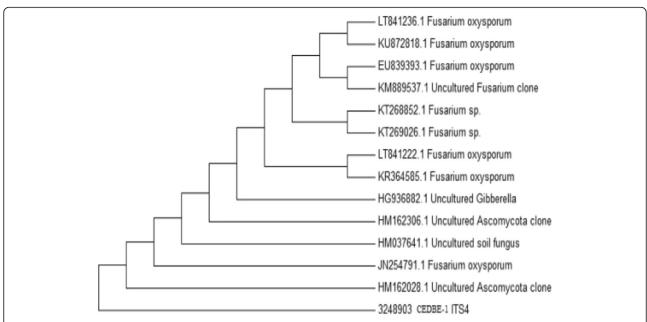


Fig. 4 Phylogenetic relationship for 5.8S-rRNA-ITS regions between the endophytic fungus CEDBE-1 and published data. Phylogenetic tree constructed using Neighbor-Joining (NJ) method with bootstrap support based on 1,000 replicates, and evolutionary distances computed using the Maximum Composite Likelihood method based on 5.8S-rRNA-ITS sequences

Discussion

Plants have been known as latent sources for novel drug compounds and many plant extracts have been used as substitute of medical treatments since the late 1990s Mangroves are pervasive in tropical and subtropical regions, especially in Asia, and they are unique for their saline environment which promises innovation of biologically active compounds, for instance antiviral, antibacterial and antifungal. *C. decandra*, a mangrove plant species, has been known to be a source of many bioactive compounds which could be found within the bark and leaves of the plants. The present study shows that the bioactive

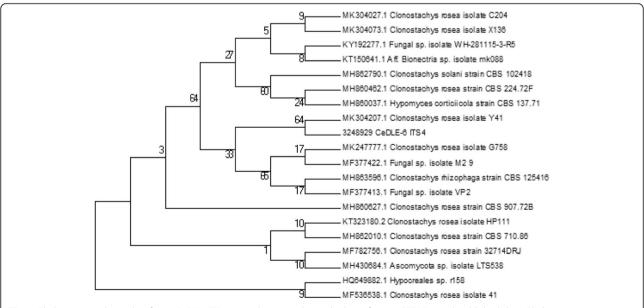


Fig. 5 Phylogenetic relationship for 5.8S-rRNA-ITS regions between the endophytic fungus CEDLE-6 and published data. Phylogenetic tree constructed using Neighbor-Joining (NJ) method with bootstrap support based on 1,000 replicates, and evolutionary distances computed using the Maximum Composite Likelihood method based on 5.8S-rRNA-ITS sequences

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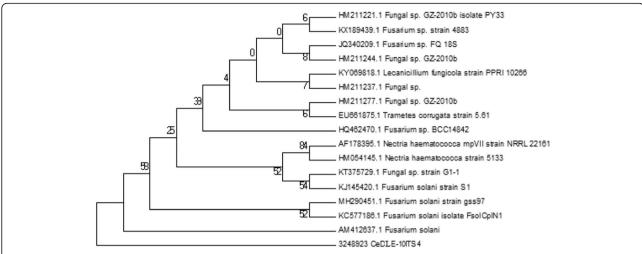
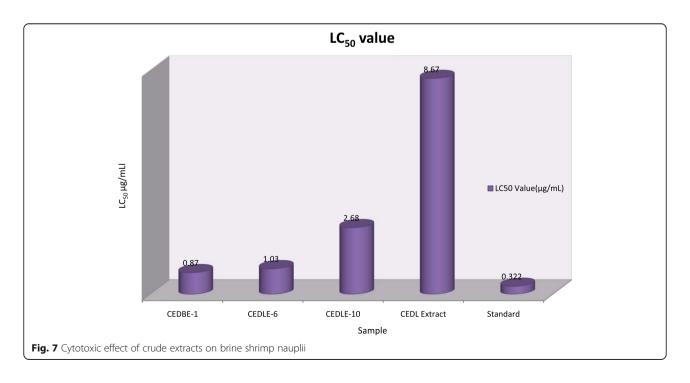


Fig. 6 Phylogenetic relationship for 5.8S-rRNA-ITS regions between the endophytic fungus CEDLE-10 and published data. Phylogenetic tree constructed using Neighbor-Joining (NJ) method with bootstrap support based on 1,000 replicates, and evolutionary distances computed using the Maximum Composite Likelihood method based on 5.8S-rRNA-ITS sequences

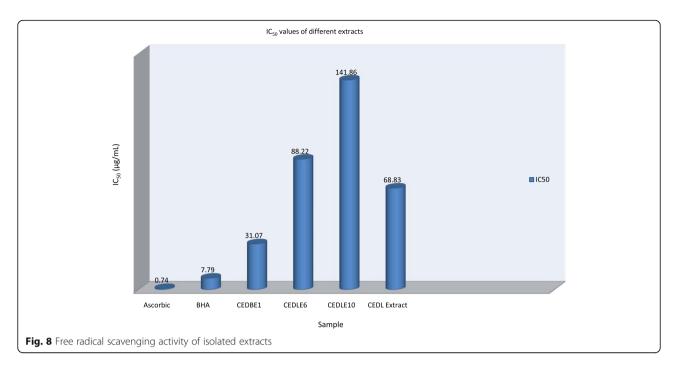
properties of *C. decandra* collected from the mangroves in Sundarban, might be due to the activity of fungal endophytes found within the leaves and barks of the mangrove plant.

This is the first report on the isolation of endophytic fungi isolated from *C. decandra* (mangrove plant). In October 2010, endophytic bacteria from leaves of mangrove and salt-marsh plant species were isolated by Gayathri, Saravanan et al. in India and they proved that the mangroves were the sources of endophytic bacteria with bioprospecting potential. [20].

Extracts of the plant *C. decandra* and its associated endophytic fungal strains CEDBE-1, CEDLE-6 and CEDLE-10 were screened for probable cytotoxic activities using brine shrimp lethality bioassay. The LC50 value of crude methanol extract of the plant was found to be 8.67 μ g/mL. On the other hand, extracts of the fungal strains CEDBE-1, CEDLE-6 and CEDLE-10 were found to be 0.87 μ g/mL, 1.03 μ g/mL and 2.68 μ g/mL respectively. It is apparent that all the test samples were lethal to brine shrimp nauplii. By evaluating these values with tamoxifen as positive control, it was found that



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crude fungal extracts illustrated effective cytotoxicity to brine shrimp. Bhimba et al. (2011) characterized the cytotoxic compound from mangrove-derived fungi Irpex hydnoides VB4. The fungal extract (tetradecan) isolated from the endophytic fungi showed very potent cytotoxic effect, indicating its possible potential for development as an anticancer drug, and warrants further scientific investigation [21].

The fungal strain CEDBE-1 showed highest antioxidant activity with IC $_{50}$ value of 31.07 µg/mL compared with IC $_{50}$ values of positive control 0.74, 7.79 for ascorbic acid and BHA respectively. CEDLE-6 and CEDLE-10 exhibited moderate antioxidant activity with IC $_{50}$ values of 88.22 and 141.86 µg/mL, respectively. These results designate that CEDBE-1 has elevated antioxidant capacity which may have potentiality to be a treatment for averting certain diseases, being either the cause or the consequence of reactive oxygen species. Huang and

coworkers (2007) suggested that the phenolic contents were the major antioxidant constituents of the endophytes [22]. Moreover, Bharathidasan et al. found *Phomopsis amygdali*, an endophytic fungus isolated from the mangrove plant in Karankadu, India, showed potent antioxidant activity against both ABTS and DPPH radicals [23].

Isolate CEDLE-10 exhibited potential antimicrobial activity against all tested human pathogenic bacteria except *S. aureus*. Other isolates CEDBE-1 and CEDLE-6 showed moderate activity against only *S. typhi*. None of the endophytes exhibited any activity next to the tested pathogenic fungi. Some other researchers' also isolated 71 endophytic fungi with antimicrobial potentiality from mangrove plants towards elected bacteria (*Bacillus subtilis, Pseudomonas aeruginosa, E. coli* and *S. aureus*) was hardened using ethyl acetate extracts of fungi cultured under stagnant circumstances. All test bacteria were

Table 2 Antimicrobial Screening of Isolated Compounds

Sample (100 μg/disc)	Diameter of Zone of Inhibition (mm)							
	Bacterial Str	Fungal Strain						
	Gram positiv	ve		Gram negative				
	B. megateriu	ım	S. aureus	S. typhi	E. coli	A. niger		
CEDBE-1				7				
CEDLE-6				8				
CEDLE-10	9			7	7			
Positive control	KM	40	36	47	48			
(30 µg/disc)	KC							

KM – Kanamycin; KC – Ketoconazole; "-" indicates no sensitivity; nd: indicates not done

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Table 3 Chemical screening of crude extracts of C. decandra and its associated fungi

Strain no	Visual	254 nm	365 nm	Remarks on probable compounds
CED	Yellow, Brown, Yellow	No color	Red, Orange,	Flavonoids, Terpenoids, Carotenoid
CEDBE-1 (Organic Part)	Purple, Pink	Dark quenching spot, Light purple, Light purple	Spot purple, Dark quenching purple.	Anthocyanin
CEDBE-1 (Aqueous part)	Purple	No color	Spot purple, Dark quenching purple.	Anthocyanin
CEDLE-6 (Organic part)	Yellow	Dark quenching purple, Light purple	Spot purple, Dark quenching spot, Purple	Anthocyanin
CEDLE-10 (Organic part)	Purple, Pink	Dark quenching purple, Orange, Light purple.	Dark quenching spot	Anthocyanin
CEDLE-10 (Aqueous part)	Pink	Purple, Light brown, Brown	Purple	Anthocyanin

repressed by a *Cladosporium* sp. isolated from the leaves of *T. populneoides* and an endophytic *Xylaria* sp. 1 isolated from *A. ilicifolius* leaves caused significant inhibition to Gram-positive and Gram-negative bacteria [24].

Conclusions

This study supports the growing indication that bioactive substances are generated by fungal endophytes *F. oxysporum*, *Clonostachys* sp. and *F. solani*. In a continuation of seeking bioactive secondary metabolites from natural sources at our laboratory, chemical exploration of ethyl acetate extract of *Clonostachys* sp. and *F. solani* strain isolated from *C. decandra* led to the isolation of effective secondary metabolites. Results of this investigation showed that CEDBE-1, CEDLE-6, CEDLE-10 and crude extract have potentiality as bioactive compounds against cytotoxicity, anti-microbial and antioxidant test. Thus, evidences indicated that *F. oxysporum*, *Clonostachys* sp. and *F. solani* strain are good sources of bioactive compounds.

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Authors' contributions

MM has conducted all the experiments with the assistance of HS, NB and SRR. MM wrote the manuscript. FA and NH have supervised the entire project, experiments and manuscript preparation. AK, FA and NH revised the manuscript and worked to improve the quality of the final manuscript. All authors read and approved the final manuscript.

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Availability of data and materials

Not applicable.

Declarations

Ethics approval and consent to participate

This study was approved by the Biological Research Division, Bangladesh Council of Scientific and Industrial Research (BCSIR), Dhaka, 1205, Bangladesh.

Consent for publication

Not applicable.

Competing interests

The authors declare that there are no conflicts of interest

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