



Bioactivity assessment of endophytic fungi associated with *Centella asiatica* and *Murraya koengii*

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ARTICLE INFO

Article history:

Received on: 16/09/2014

Revised on: 30/09/2014

Accepted on: 19/10/2014

Available online: 30/10/2014

Key words:

Centella asiatica; *Murraya*

koengii; Endophytic fungi;

Ethanol extract;

Antioxidant; Antimicrobial.

ABSTRACT

The present study attempted to evaluate the antimicrobial and antioxidant activity of fungal endophytes associated with *Centella asiatica* and *Murraya koengii*, the ethnomedicinal plants used in traditional practices. Two endophytic fungi *Aspergillus oryzae* CeR1 and *Colletotrichum gloeosporioides* MKL1 were isolated from *C. asiatica* and *M. koengii* respectively which were characterized morphologically and by using rDNA- internal transcribed spacer. The total antioxidant power and free radical scavenging activity of the fungal extracts was estimated using the total phenolic content and 2, 2-diphenyl-1-picrylhydrazyl (DPPH) free radical assay along with the flavonoids and alkaloids content. The DPPH free radical assay showed that the ethanolic extract of endophytic fungi *Aspergillus oryzae* CeR1 had higher radical scavenging activity than *Colletotrichum gloeosporioides* MKL1. Antimicrobial activity of fungal extract tested against three bacterial strain namely, *Staphylococcus aureus* MTCC 96, *Streptococcus pyogenes* MTCC1925, *Enterococcus faecalis* MTCC2729 and one fungal strain *Candida albicans* MTCC183 showed inhibition of growth of test organisms except *Staphylococcus aureus* MTCC96.

1. INTRODUCTION

Endophytic microorganisms are recognized as one of the most chemically promising groups of microorganisms in terms of diversity and pharmaceutical potential. These are microorganisms that grow in the intercellular spaces of higher plants without causing visible damage to their hosts and comprise especially of fungi and bacteria [1]. These microorganisms, in some way contribute to the well being of the plant and being associated with living tissues they are not considered as saprophytes. There are reports indicating the endophytic organisms to be the chemical synthesizers inside the host plants [2]. The biosynthesized chemicals include bioactive compounds used by the host as a defense against pathogens. Some of these bioactive compounds have been proven to be a source for novel drug which are reported to be useful as agro-chemical, antibiotics, immunosuppressant, antimicrobial, anti-parasitic, antioxidant, anticancer agents [3]. The wide spectrum of bioactive secondary metabolites that are characterized from endophytic microbes especially fungi include alkaloids, benzopyranones, flavonoids, phenolic acids, quinines, steroids,

terpenoids, tetralones, xanthenes [4]. The ethno-medicinal plants have been recognized as a repository of endophytes with prospects of providing novel metabolites of pharmaceutical importance [3,5,6]. Many natural products associated with endophytic fungi have been to be potential as antifungal, anti-oxidant, anticancer, anti-inflammatory and antimicrobial agents [7].

Preliminary survey of *Murraya koengii* has demonstrated that it has antioxidant, antidiabetic, antibacterial, larvicidal, hypoglycaemic, antiprotozoal, hypolipidemic, antilipid peroxidative, respiratory protective, antihypertensive, cytotoxic and trypsin inhibitor activity [8]. Likewise chemical constituents reported in *Centella asiatica* include asiaticoside, madecassoside, madecassic acid, asiatic acid, glucose, rhamnose, terpenoids, sitosterol, stigma sterol, fatty oils like glycerides of palmitic acid, stearic acid, linoleic acid, linolenic acid, vitamins like ascorbic acid, and calcium, iron, and phosphate [9, 10]. *C. asiatica* has also been reported to be useful in the treatment of inflammations, diarrhea, asthma, tuberculosis and various skin lesions and ailments like leprosy, lupus, psoriasis and keloid [11, 12].

The exploration of plants for endophytic fungi can be of immense value in screening for potential metabolites as was the case with *Taxus brevifolia* which had fungal endophyte that produced metabolites independently of the tree [13]. The present investigation was designed to characterize the endophytic fungi associated with the two ethnomedicinal plants, *Centella asiatica*

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and *Murraya koengii* with the aim to characterize the associated endophytic fungi and investigate their antimicrobial and antioxidant activity. The hypotheses for the study was to assess if the associated endophytic fungi have some properties as already reported for their host plant and if so, the endophytes can be bioprospected as microbial cell factories for metabolites of pharmaceutical importance.

2. MATERIALS AND METHODS

2.1. Sampling

Mature healthy plant parts (leaves, roots, shoots/branches) with no visible symptoms of disease were carefully selected from different regions based on their traditional usage, brought to the laboratory in sterile bags and processed within 24 hr after sampling.

2.2. Surface sterilization and isolation of the endophytic fungi

Isolation of endophytic fungi was done according to the method described by Hallmann *et al.* [14] with minor modifications. The plant samples were rinsed gently in running water to remove adhered dust and debris and were cut into 2 mm segments which were surface sterilized with 70% ethyl alcohol for 1 min. The segments were soaked in 4% sodium hypochlorite solution for 3 min, and then rinsed with 70% ethyl alcohol for 1 min, rinsed with sterile distilled water and blot dried on sterile filter paper followed by drying under laminar airflow chamber. The segments were then inoculated into potato dextrose agar (PDA, Himedia, India) plates and incubated at $25 \pm 2^\circ\text{C}$ for 5-7 days. The growth of endophytic fungi from the plant segments were observed once a day. Hyphal tips growing out of the plated segments were transferred into a fresh PDA plates and incubated at $25 \pm 2^\circ\text{C}$ for 5 days to obtain pure cultures. The pure fungal cultures were preserved on the agar slant at 4°C .

2.3. Morphological features

Microscopically, the endophytic fungal isolates were tentatively identified on the basis of their hyphal features, arrangement of spores and reproductive structures.

2.4. Molecular characterization of endophytic fungi

Genomic DNA of the fungal endophytes was isolated using the HiPurA fungal DNA isolation kit (Himedia). The DNA samples were stored at 4°C for immediate use and stored at -20°C for long-term storage. Fungal rDNA-ITS region was amplified using the fungal domain specific ITS1 and ITS4 [15]. The PCR reaction mixture comprised of 10 μl fungal DNA, 5 μl $10 \times$ PCR buffer, 1 μl of 10 mM dNTP, 0.25 μl Taq polymerase, 2 μl each of the forward and the reverse primers in a total reaction volume of 50 μl . PCR was performed in a Gene Amp 9700 Thermal Cycler (Applied Biosystems, USA) with an initial denaturation step at 94°C for 5 min, followed by 30 cycles of 94°C for 1 min, 52°C for 30 sec, and 72°C for 1 min, with a final extension step of 72°C for 10 min. The amplified ITS was analysed in 1.5% agarose gel

electrophoresis in 1X Tris-acetate-EDTA with a marker ladder of 100-bp and ethidium bromide staining. The amplified ITS products were purified using QIA Quick Gel Extraction Kit (Qiagen, Hilden, Germany) according to manufacturer's instructions. Big dye ready reaction dye deoxy terminator cycle sequencing kit (Applied Biosystems) was employed and sequenced in Applied Biosystems 3700 Genetic Analyzer. The obtained sequences were then analyzed using the BLAST algorithm and closely related phylogenetic sequences obtained from the National Centre of Biological Information (NCBI) database. The phylogenetic tree was constructed using Neighbor Joining method in MEGA 5 [16].

2.5. Preparation of fungal extract

The cultivation of fungus was done on 100 ml Potato dextrose broth by placing agar blocks of actively growing pure culture (3 mm in diameter) in 250 ml Erlenmeyer flask. Each flask was incubated at $25 \pm 2^\circ\text{C}$ for 3 weeks with periodical shaking at 150 rpm. The culture was then filtered through three layers of muslin cloth to remove fungal mycelia. The culture filtrate was then filtered thrice with equal volumes of solvent ethanol. The organic phase was collected and the solvent removed by evaporation under reduced pressure at 45°C using rotary vacuum evaporator. The dry solid residue was re-dissolved in ethanol (5mg/ml) and evaluated for its antimicrobial and antioxidant activities.

2.6. Test microorganisms

Staphylococcus aureus MTCC 96, *Candida albicans* MTCC183, *Streptococcus pyogenes* MTCC1925, and *Enterococcus faecalis* MTCC2729 procured from Institute of Microbial Technology (IMTECH, India) were used as indicator organisms for antimicrobial activity. Test bacteria were subcultured every two weeks on fresh nutrient agar slants and incubated at 37°C , whereas *C. albicans* was subcultured every four week on fresh PDA slant and incubated at 37°C . The cultures were kept at 4°C until further use.

2.7. Antimicrobial activity by well-diffusion method

The antimicrobial activity of the crude extracts of fungi was carried out on Mueller Hinton agar medium for test bacteria and Potato dextrose agar medium for test fungus. Using the sterile cork borer 7 mm wells were made in which 20 μl of crude ethanol fraction was loaded on inoculum swapped plates. The plates were observed for zone of inhibition after incubation for 24 hr at 37°C for bacteria and at 25°C for fungus.

2.8. Radical scavenging activity using DPPH method

Different concentrations of the extracts (10 μl , 25 μl , 50 μl , 75 μl and 100 μl) were taken in test tubes and the volume adjusted to 100 μl with ethanol. Three ml of 0.1 mM ethanolic solution of DPPH was added to these tubes. The mixture was shaken vigorously, left to stand for 30 min in the dark, and the absorbance was measured at 517 nm. The percentage inhibition of

DPPH radical by the sample extract was calculated using the following relation:

$$\text{DPPH scavenging effect (\%)} = [(A_0 - A_1/A_0) \times 100]$$

Where, A₀ is the absorbance of the control reaction and A₁ is the absorbance in presence of the extract.

2.9. Estimation of Total Phenolic Content (TPC)

The total polyphenolic content in the fungal culture extract was determined calorimetrically using the Folin-Ciocalteu (FC) method as described by Singleton *et al.* [17] with minor modifications. 0.2 ml of test sample was mixed with 0.2 ml of FC reagent and allowed to stand for 10 min to which 0.6 ml of 20% sodium carbonate (Na₂CO₃) was added and mixed thoroughly. The reaction mixture was incubated at 40°C for 30 min and absorbance measured at 765 nm with gallic acid taken as standard.

2.10. Phytochemical analysis

2.10.1. Flavonoid Tests

Presence of flavonoids in ethanolic extracts of endophytic fungi was carried out using the method of Brain and Turner [18] and Trease and Evans [19]. Shinoda's test for Flavonoids: About 5 mg of the fungal extract concentrate was dissolved in ethanol to which 3 mg magnesium powder was added followed by few drops of concentrated hydrochloric acid (HCl). An orange coloration indicated the presence of Flavonoids. Ferric Chloride test for Flavonoids: About 5 mg of the extract concentrate was dissolved in 2 ml of ethanol to which a few drops of 10% ferric chloride solution were added. A green-blue coloration indicated the presence of a phenolic hydroxyl groups.

Sodium Hydroxide (NaOH) test for Flavonoids: The fungal extract was warmed and filtered, and 200 µl of 10% aqueous NaOH was added to the filtrate. This produced a yellow coloration. A change in color from yellow to colorless on addition of dilute HCl was an indication of the presence of flavonoids.

2.10.2. Alkaloid Test

Wagner's test: About 5 mg of the fungal extract was treated with Wagner's Test reagent [1.27 g of Iodine and 2 g of Potassium iodide in 100 ml of distilled water]. A reddish brown color indicated the presence of alkaloid [20].

3. RESULTS

3.1. Isolation and identification of fungi

A total of nine and seven isolates from *Centella asiatica* and *Murraya Koenigii* respectively were isolated from different parts of the plants. Based on microscopic and morphological features, one isolate each from the plants (CeR1 from *C. asiatica* and MKL1 from *M. koengii*) were considered for molecular characterization and phytochemical study. The rDNA-ITS region was amplified, sequenced and submitted to the NCBI GenBank with accession numbers KF358716 and KF358717. BLAST search of ITS gene sequences affiliated the endophytic fungal isolates CeR1 and MKL1 to be closest homolog of *Aspergillus oryzae* and *Colletotrichum gloeosporioides* respectively (Fig.1, 2).

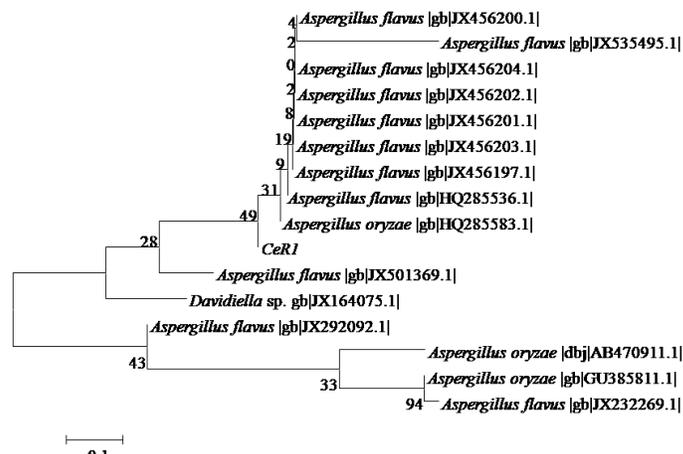


Fig. 1: Evolutionary positions of the endophytic fungal isolate CeR1 with other related fungal species based on internal transcribed spacer sequence similarity.

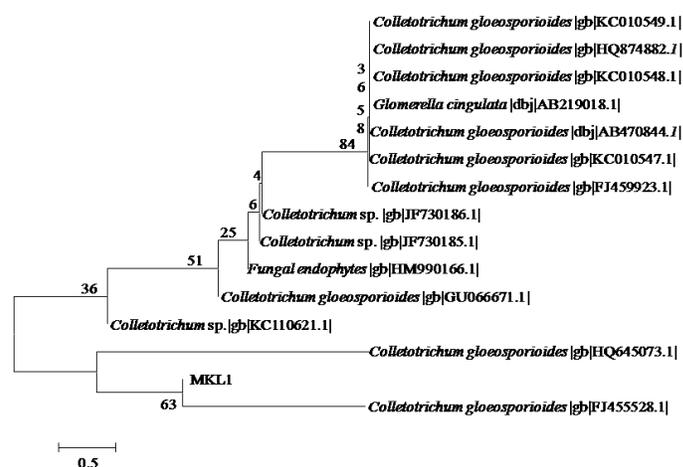


Fig. 2: Evolutionary positions of the endophytic fungal isolate MKL1 with other related fungal species based on internal transcribed spacer sequence similarity.

3.2. Antimicrobial assay

The crude extracts of the fungi displayed considerable antimicrobial activity against some pathogens (Table 1). None of the fungal extracts exhibited inhibition of *Staphylococcus aureus* MTCC 96. The crude extract of *Aspergillus oryzae* CeR1 inhibited the growth of *Candida albicans* MTCC183 but had no effect on *Streptococcus pyogenes* MTCC1925 and *Enterococcus faecalis* MTCC2729. In contrast, the crude extract of *Colletotrichum gloeosporioides* MKL1 had no effect on *Candida albicans* MTCC183 and *Staphylococcus aureus* MTCC96.

3.3. DPPH radical scavenging activity

The comparative analysis of DPPH assay between *Aspergillus oryzae* CeR1 and *Colletotrichum gloeosporioides* MKL1 revealed the latter to have higher radical scavenging activity than the former (Table 2). The fungal extracts showed better activity than ascorbic acid which was used as the reference standard.

Table. 1: Antimicrobial activity as zone of inhibition (in mm) of endophytic extracts by well diffusion method.

Plant parts used for isolation	Characterized isolates	<i>Staphylococcus aureus</i>	<i>Candida albicans</i>	<i>Streptococcus pyogenes</i>	<i>Enterococcus faecalis</i>
Roots	<i>Aspergillus oryzae</i> (CeR1)	-	+	-	-
Leaf	<i>Colletotrichum gloeosporioides</i> (MKL1)	-	-	++	+

-No zone of inhibition; + inhibition zone between 10~14 mm; ++ inhibition zone between 15~20 mm.

Table. 2: Scavenging activity (%) on DPPH radicals of ethanolic extracts of endophytic fungi at different concentration ($\mu\text{g/mL}$).

Fungal Isolates	Concentrations ($\mu\text{g/ml}$)				
	10	25	50	75	100
<i>Aspergillus oryzae</i> (CeR1)	4.66 \pm 0.05	26.16 \pm 0.13	15.29 \pm 0.08	104.88 \pm 0.10	72.88 \pm 0.18
<i>Colletotrichum gloeosporioides</i> (MKL1)	0.67 \pm 0.05	47.89 \pm 0.06	73.84 \pm 0.08	60.09 \pm 0.08	52.77 \pm 0.06
Ascorbic acid	0.15 \pm 0.21	16.05 \pm 0.3	17.75 \pm 0.25	19.56 \pm 0.19	21.36 \pm 0.43

Table. 3: Total phenolic content (TPC) in the ethanolic extracts of endophytic fungi.

Fungal isolates	TPC (mg gallic acid equivalent / g dry wt.)
<i>Aspergillus oryzae</i> (CeR1)	1.4 \pm 0.01
<i>Colletotrichum gloeosporioides</i> (MKL1)	0.72 \pm 0.03

Table. 4: Test results for flavonoids and alkaloids from the endophytic fungi.

Fungal isolates	Flavonoids			Alkaloids
	Shinoda's Test	Ferric chloride Test	Sodium hydroxide Test	
<i>Aspergillus oryzae</i> (CeR1)	-	-	-	-
<i>Colletotrichum gloeosporioides</i> (MKL1)	+	+	-	+

+ indicates present; - indicates absent.

3.4. Total phenolic content (TPC)

Total phenolic content was found to be higher in the culture extract of *Aspergillus oryzae* CeR1 as compared to *Colletotrichum gloeosporioides* MKL1 (Table 3) when expressed as gallic acid equivalent (GAE) i.e. mg gallic acid/g dry wt. The levels of phenolic contents in the endophytic fungal extracts were significantly different from each other.

3.5. Phytochemical analysis

Presence of flavonoid and alkaloid was noted in the extracts of *Colletotrichum gloeosporioides* MKL1 whereas both were absent in the crude fungal extract of *Aspergillus oryzae* CeR1 (Table 4).

4. DISCUSSION

There has been a growing interest in finding the alternative to reduction of the problem of drug resistance among the pathogens and antimicrobial metabolites from fungal endophytes has been explored as one of the alternate approaches [5]. The fungal isolates obtained from both the medicinal plants in the present study i.e. *Aspergillus oryzae* CeR1 and *Colletotrichum gloeosporioides* MKL1 revealed activities and phytochemical contents which can be explored for generation of bioactive metabolites. The endophytic fungus, *Colletotrichum gloeosporioides* MKL1 showed better inhibition of growth of the pathogenic microorganisms when compared to *Aspergillus oryzae* CeR1. This could be due to their higher efficiency in metabolite production indicating that these fungi have differing activity potential. There is a vast diversity of microbes which still remains untapped for evaluation of metabolites production that may possess valuable bioactivities including antioxidant activity [21]. There is a strong relationship between total phenol content,

flavonoid, alkaloid and antioxidant activity, as phenols possess strong scavenging ability for free radicals due to their hydroxyl groups [22]. Therefore, the phenolic content of may directly contribute to their antioxidant action as has been reported for plants [23, 24, 25]. Phenolic compounds are powerful chain breaking antioxidants and are reported to be associated with antioxidant activity and play a crucial role in stabilizing lipid peroxidation [26]. The observation in the study revealed that crude ethanolic extract of *Aspergillus oryzae* CeR1 does not show the presence of both alkaloid and flavonoid, however, it showed highest phenolic content which makes it a potential antioxidant. The DPPH assay has been largely used as a quick, reliable and reproducible parameter to search the *in vitro* general antioxidant activity of pure compounds [27].

The effect of antioxidants on DPPH is thought to be due to their hydrogen donating ability [28, 29]. A freshly prepared DPPH solution exhibits a deep purple color which disappears when an antioxidant is present in the medium. An antioxidant molecules can quench DPPH free radicals and convert them to a colorless product, resulting the decrease in absorbance. The reducing capacity of compounds thus, serve as indicator of potential antioxidant property [30]. Considering the total phenolics, reducing power and the DPPH radical scavenging activity as indices of antioxidant activity of the extract, the present findings reveal the potential of the extract as a source for natural antioxidants [31]. It indicates that the metabolites of endophytic fungi isolated from *Aspergillus oryzae* CeR1 and *Colletotrichum gloeosporioides* MKL1 could be potential agents in scavenging free radicals and treating diseases related to free radical reactions.

A moderate correlation exists between the amount of total phenolics and the biological activity, when a comparison of both antioxidant and antimicrobial activities is made in relation to the amount of total phenolics of different natural extracts [32, 33,

34, 35, 36] In this context, the crude extracts of the endophytes, *Aspergillus oryzae* CeR1 and *Colletotrichum gloeosporioides* MKL1 presented a moderate antioxidant activity and a better greater antimicrobial activity opening scope for their exploration as a natural antioxidant in food industries and other pharmaceutical preparations [37, 38]. This work is the first report on fungal endophytes of *Centella asiatica* and *Murraya koenigii* which provides an insight into understanding some basis of therapeutic properties traditional medicine. However, further studies on the characterization of the metabolites from the culture extracts of endophytic fungi obtained in the study as well as their *in vivo* assays are necessary for their pharmaceutical bioprospection.

5. ACKNOWLEDGEMENT

The support received as research grant from the Department of Biotechnology, Govt. of India is thankfully acknowledged.

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How to cite this article:

Archana Nath, Jyoti Pathak and SR Joshi. Bioactivity assessment of endophytic fungi associated with *Centella asiatica* and *Murraya koengii*. J App Biol Biotech. 2014; 2 (05): 006-011.