

Fungal Endophytes from *Tabernaemontana heyneana* Wall. (Apocynaceae), their Molecular Characterization, L-asparaginase and Antioxidant Activities

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Abstract

Endophytic fungi are an interesting group of microorganisms associated with the healthy tissues of medicinal plants used in folk medicines for health benefits. Fifteen fungal endophytes were isolated from the plant parts of the ethnomedicinal plant of Western Ghats, *Tabernaemontana heyneana* Wall., and evaluated for their L-asparaginase and antioxidative potentials. They were identified by rDNA sequencing of the ITS region and characterized as strains of *Fusarium*, *Colletotrichum*, *Curvularia*, *Nigrospora*, *Plectospharella*, *Neocosmospora*, *Trichoderma* and *Endomelanconiopsis*. *Trichoderma hamatum* and *Fusarium* sp., colonized the bark, leaf and stem tissue fragments. Secondary metabolites from the endophytic strains were extracted in ethyl acetate and evaluated for total phenolic content and antioxidant activity. The total phenolic content of the extracts from all the fungal strains ranged from 17.29±12.92 to 90.41±4.56 mg Gallic acid equivalent/g (mg GAE/g) dry extract and IC₅₀ values in DPPH radical scavenging activity ranged from 51.64±8.91 µg/ml to 764.05±27.67 µg/ml. Eleven strains were positive for L-asparaginase enzyme activity in the range of 1.44 to 3.42 IU/ml. *Fusarium tricinctum* (MK752630) strain potentially produced high enzyme activity of 3.423 IU/ml. The highest total phenolic content (90.14±4.56 mg GAE/g dry extract) and scavenging activity (IC₅₀ value of 51.64±8.91 µg/ml) were detected in *F. tricinctum*.

Keywords: Endophytic fungi; L-asparaginase; *Fusarium tricinctum*; total phenolic content, DPPH radical scavenging, Western Ghats

1. Introduction

Plants are known to harbor endophytic fungi that reside in internal tissues without showing any disease symptoms (Frohlich and Hyde, 1999). The infected host tissues are transiently symptomless, and the microbial colonization can be demonstrated through histological means, by isolation from strongly surface disinfected tissue, or, most recently, through the direct amplification of fungal nuclear DNA from colonized plant tissue (Stone *et al.*, 2000). Fungal endophytes represent an important component of fungal biodiversity and are known to have mutualistic association with host plants (Selim *et al.*, 2012). Endophytes are known to influence population dynamics, plant community diversity and ecosystem functions (Saikkonen *et al.*, 1998). This group of microbes not only synthesizes metabolites to compete first with epiphytes and then with pathogens in order to colonize the host, but also regulates metabolism of the host in a balanced association (Schulz *et al.*, 2002). Once inside host tissue, they assume a quiescent state either for the whole time or for an extended period of time until environmental conditions are favorable for the fungus or the ontogenetic state of the host changes to the advantage of the fungus (Sieber, 2007).

Endophytic fungi are known to have profound effects on plant ecology, fitness, evolution (Brundett, 2006) and play vital roles in host plants being chemical synthesizers

inside plants (Owen and Hundley 2004), producing phytohormones (Schulz and Boyle, 2005), withstanding environmental stress (Malinowski and Belesky, 2000) and providing protection against pathogens (Akello *et al.*, 2007). In recent years, endophytes from medicinal plants have demonstrated the ability to produce bioactive compounds which are potential sources of novel natural products (Nalini *et al.*, 2019).

Asparaginase is widely distributed in nature from bacteria to mammals and plays a central role in the amino acid metabolism and utilization. L-asparaginase (EC 3.5.1.1.) belongs to the group of homologous amidohydrolases family, which catalyses the hydrolysis of the amino acid L-asparagine to L-aspartate and ammonia. L-asparaginase is the first enzyme with anti-leukemic activity studied thoroughly in human beings (Savitri *et al.*, 2003). Asparaginase is used in food industry to prevent the formation of acrylamide when foods are processed at high temperatures (Cachumba *et al.*, 2016).

Naturally occurring phenolic compounds are plant secondary metabolites which are good electron donors since their hydroxyl groups can directly contribute to antioxidant action. Currently, there is a great interest in finding natural antioxidants from endophytes associated with plant species. Plant-endophyte interaction and endophyte-endophyte interaction studies provide understandings of metabolite production by fungi. Endophytes are the repository of natural products

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displaying broad spectrum of biological activities like anticancer, antibacterial, antiviral, anti-diabetic, antioxidant and anti-inflammatory (Rajamanikyam *et al.*, 2017).

Tabernaemontana heyneana Wall., is endemic to southern Western Ghats and is listed as lower risk/ near threatened species in the "IUCN Red List of Threatened Species" (World Conservation Monitoring Centre, 1998). *T. heyneana* possess antimicrobial and anthelmintic properties against skin venereal, respiratory and nervous disorders (Duraipandiyar *et al.*, 2006) and get used in the traditional Indian system of medicine as a cure for cancer (Baskar *et al.*, 2012). The medicinal plants inhabiting Western Ghats are major sources of traditional pharmaceuticals and therapeutics of significant potentialities. Therefore, we have investigated the endophytic fungi residing in *T. heyneana*, their molecular identification and evaluation of their L-asparaginase and antioxidative capacities by *in vitro* methods.

2. Materials and methods

2.1. Collection of host plant

Healthy plant samples of *T. heyneana* Wall., were collected in the forests of Kodagu district, in the Talacauvery area of Western Ghats (12°17' to 12°27' N and 75°26' to 75°33' E), Karnataka, India, during the month of June 2014. The habit of plant and collection locality is shown in Figure 1. A total of five leaves, stem and bark samples were collected separately from five plants. Bark samples from the trunk were cut 1.5 m above the ground level with help of sterile machete. The samples were placed in pre-sterilized zip-lock polythene bags, stored at 4°C and transported to the laboratory. Fresh plant materials were used for the isolation of fungal endophytes to reduce the chance of contamination. Thus collected plant materials were subjected to surface sterilization within few hours after sampling.

2.2. Surface sterilization and isolation of fungal endophytes

The leaf, stem and bark samples were rinsed gently in running tap water to remove dust and debris. Before surface sterilization, the cleaned stems were cut into pieces of 5 cm length and bark samples were halved. Samples were first immersed in 70% (v/v) ethanol for one minute, 3.5% (v/v) Sodium hypochlorite for two minutes. The samples were rinsed three times with sterile distilled water and dried on sterile blotters under laminar airflow to ensure complete drying (Schulz *et al.*, 1993). Bits of 1.0X0.1 cm size were excised with the aid of sterilized blade. 200 segments of leaf, stem and bark were evenly placed on water agar (WA) medium (15 g/L). The Petri dishes were sealed using Parafilm™ and incubated at 27±2° C in a light chamber with 12 hours of light followed by 12 hours of dark cycles for 4-6 weeks. The Petri dishes were monitored periodically to check the growth of endophytic fungal colonies from the segments. The hyphal tips which grew out from the segments were transferred separately onto fresh Potato Dextrose Agar (PDA) slants with a sterile fine tipped needle under stereo binocular microscope and incubated at 27±2° C for 10-15 days and pure cultures were maintained at 4° C for further use.

2.3. Identification of endophytic fungi

Morphological identification was done by inoculating the endophytic fungi on PDA plates followed by seven days of incubation and observation of colony and spore morphology. The slides of each fungal endophytes were prepared by tease mount method using lactophenol cotton blue staining [NICE, Kerala, India] and observed under the light microscope (Quasmo, Haryana, India) with 400X magnification. The identification was based on the observation of mycelia, fruiting bodies, conidial characters according to the standard identification manuals (Domsch *et al.*, 1980; Singh *et al.*, 1991; Barnett and Hunter, 1998; Mathur and Kongsdal, 2003).

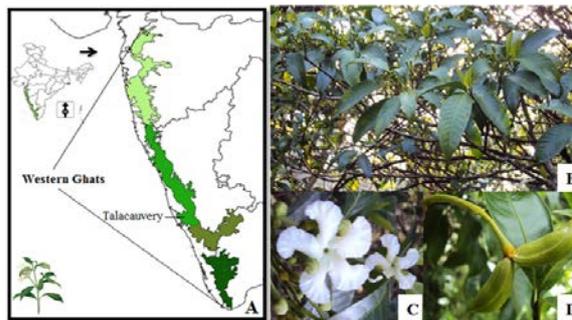


Figure 1. A. Map showing the location of plant sample collection from Talacauvery site of Western Ghats; B. *Tabernaemontana heyneana* Wall. habit; C-Flowers; D- A pair of follicles.

2.4. Molecular characterization of fungal endophytes

Hyphal tips from 15 morphologically different endophytic fungi, each representing individual morphospecies, were inoculated into conical flask containing Potato Dextrose Broth (PDB, Hi Media, Mumbai, India) and incubated at 27±2 °C for 7-10 days. The resulting mycelia were separated and placed in polythene zip lock covers. The mycelia were then freeze dried before being ground in a pestle and mortar containing liquid nitrogen. The powdered mycelia were stored at -20 °C until required. The genomic DNA was extracted using cetyltrimethylammonium bromide (CTAB) method with minor modifications (Ausubel *et al.*, 1994). The DNA concentration was measured using Nanodrop Spectrophotometer (Thermo Fischer 2000c, Bangalore, India) at 260 and 280 nm.

PCR amplification in ITS rDNA region of isolates was conducted using PCR 18 kit (Chromous Biotech Pvt. Ltd. Bangalore, India). The target ITS 1, ITS 2 regions and 5.8s gene were amplified using ITS 1 (5'-TCCGTAGGTGAACCTGCG - 3') and ITS 4 (5'-TCCTCCGCTTATTGATATGC - 3') primers. Amplification reactions were performed in a thermal cycler (Master cycler gradient, Eppendorf, Germany) with the following cycling conditions 94°C for 2 min (Initial denaturation), 35 cycles of 94°C for 1 min (Denaturation), 47°C for 15 sec (Primer annealing), 72°C for 30 sec (Primer extension), followed by 10 min of final extension at 72°C. Following amplification, the PCR products were analyzed by horizontal agarose gel electrophoresis through 1% agarose gel supplemented with ethidium bromide along with 100bp DNA marker. DNA bands on the gel were visualized under UV light trans-illuminator (Geldoc XRT, BioRad USA) and documented.

The PCR products were sent to Chromous Biotech Pvt. Ltd., Bangalore, India for purification and sequencing. Sequencing similarity searches were achieved for the obtained fungal sequences and compared with ITS sequence data from strains available in the database GenBank (National Centre for Biotechnology Information website; <http://www.ncbi.nlm.nih.gov/>) by using the BLAST sequence match routines.

2.5. Data Analysis

The relative colonization frequency (%CF) was calculated as the number of segments colonized by a fungal divide by total number of segments plated x 100.

2.6. L-asparaginase activity of fungal endophytes

Fifteen fungal endophytes isolated from leaf, stem and bark of *T. heyneana* were subjected to rapid screening of primary plate assay method using modified Czapek Dox (MCD) agar medium using phenol red as an indicator (Gulati *et al.*, 1997). The composition of the prepared medium was as follows: glucose- 2.0 g/l, L-asparagine- 10 g/l, potassium dihydrogen phosphate (KH_2PO_4)- 1.52 g/l, potassium chloride (KCl)- 0.52 g/l, magnesium sulphate ($\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$)-0.52 g/l, copper nitrate ($\text{CuNO}_3 \cdot 3\text{H}_2\text{O}$)-0.001 g/l, zinc sulphate ($\text{ZnSO}_4 \cdot 7\text{H}_2\text{O}$)- 0.001 g/l, ferrous sulphate ($\text{FeSO}_4 \cdot 7\text{H}_2\text{O}$)- 0.001 g/l and agar 20g/l. 2.5% (w/v) stock solution of the phenol red dye was prepared and MCD medium was supplemented with 0.009% phenol red dye indicator to check the ability of test fungi that grow in the medium. Final pH of the medium was adjusted to 6.2. The control plates were prepared with KNO_3 -0.001 g/l as the nitrogen source and lacking in phenol red dye indicator. The prepared media was autoclaved and poured into pre-sterilized plates and marked into four quadrants. Mycelial plugs from four different endophytic fungi were inoculated and incubated at $27 \pm 2^\circ\text{C}$ for five days. The colonies showing pink zones were inoculated individually on MCD agar plates to approve the activity of enzyme.

2.7. Quantitative assay for enzyme production

For the production of L-asparaginase under liquid state, the mycelial discs from positive agar plates were inoculated into modified Czapek Dox's broth and incubated at 30°C in orbital shaker (GeNei™, Bangalore) for 5 days at 120 rpm. The crude enzyme sample was extracted by centrifugation and supernatant that contained the enzyme used for further analysis by Nesslerization. The reaction mixture consisting of 0.5 ml of 0.04 M L-asparagine, 0.5 ml of 0.5 M TrisHCl buffer (pH 8.2), 0.5 ml of enzyme was obtained from the culture filtrate and 0.5 ml of distilled water. The samples were incubated for 30 min at 27°C . To stop the enzymatic reaction, 0.5 ml of 1.5 M trichloroacetic acid (TCA) was added. This was followed by pipetting 0.1 ml of the mixture into fresh tubes containing 3.7 ml of distilled water and 0.2 ml of Nessler's reagent and incubated at 27°C for 20 minutes, after which the absorbance of the samples was measured at 450 nm using UV-Visible spectrophotometer (TPL Technology

Pvt. Ltd., Bangalore). Blank was prepared by adding TCA followed by enzyme sample. One international unit (IU) of L-asparaginase was expressed as the amount of enzyme that catalyzes the formation of $1\mu\text{mol}$ of ammonia per minute at 27°C (Imada *et al.*, 1973).

2.8. Submerged fermentation and extraction of secondary metabolites

The sequential steps involved in fermentation and extraction of secondary metabolites is represented in Figure 2. At first the endophytic fungi were cultivated by inoculating the actively growing mycelia from 7-days-old pure culture into Erlenmeyer flask containing 500 ml of Potato Dextrose Broth. The flasks were then incubated for 21 days at $27 \pm 2^\circ\text{C}$. After incubation, the culture filtrates were filtered through muslin cloth to separate mycelia. To the culture filtrate, equal volume of ethyl acetate was added and mixed well for 10 minutes and kept for 5 minutes till two clear immiscible layers formed. The upper layer of the solvent containing extracted compounds was then separated, evaporated and dried in Rotary flash evaporator (Superfit Model PBU-6D, India). The dry solid residue was dissolved in ethanol and stored as crude extract in glass vials.

2.9. Determination of total phenolic content

The total phenolic content of ethyl acetate extracts of endophytic fungi was estimated using Folin-Ciocalteu (FC) reagent based assay using Gallic acid standard (Liu *et al.*, 2007). One ml of FC reagent and two ml of (20%, w/v) Na_2CO_3 was mixed with methanolic crude extracts. The mixture was incubated at room temperature in the dark for 45 minutes, and the absorbance of the developed color was recorded at 765 nm using UV-vis spectrophotometer (DU 730 "Life sciences", Beckman Coulter). Gallic acid was used as a reference standard for plotting the calibration curve. The TPC of the extracts was expressed as mg of Gallic acid equivalent (GAE)/ g of the dry extract.

2.10. Determination of antioxidant capacity by DPPH (2, 2-diphenyl-1-picrylhydrazyl) assay

The antioxidant activity of endophytic fungal extracts was determined by free radical scavenging activity of DPPH by the method of Pannangpetch *et al.* (2007) with slight modifications. A total of two ml of DPPH solution (DPPH, Sigma-Aldrich, Bangalore, India) (0.001 mM) was added into the endophytic fungal extracts and incubated for 20 minutes at room temperature in the dark. After incubation, the absorbance was measured at 517 nm. Ascorbic acid was used as reference antioxidant compound. The scavenging activity was expressed as inhibitory concentration 50% (IC_{50}) ($\mu\text{g}/\text{ml}$). The scavenging activity was calculated using the following equation:

$$\% \text{ Scavenging} = \frac{\text{Ac} - \text{As}}{\text{Ac}} \times 100$$

Ac= Absorbance of control; As= Absorbance of test sample

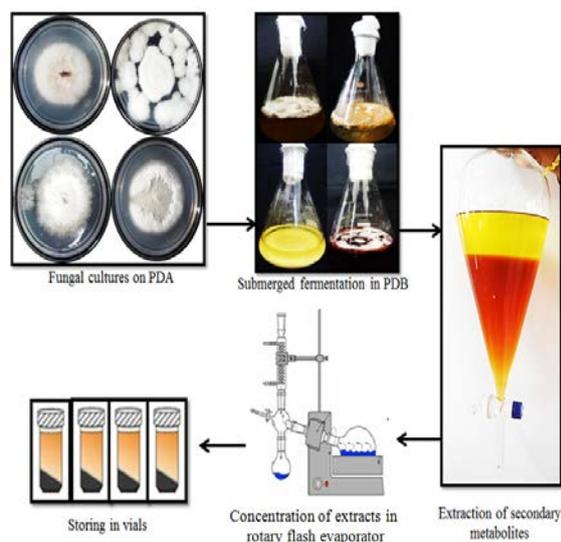


Figure 2. Fermentation and extraction of secondary metabolites from the fungal endophytes of *T. heyneana*

3. Results

3.1. Isolation and the identification of fungal endophytes

A total of 600 tissue segments yielded 477 isolates were obtained, which were distributed in 15 endophytic species. The results revealed that leaf fragments were colonized by more number of fungal endophytes (186) followed by stem (171) and bark segments (120). The analysis of distribution frequencies of endophytes indicated that the fungal communities in the host contained a few frequent genera and some infrequent species.

The isolates were identified using morphological and microscopic characteristics with the support of molecular analysis. 15 endophytic species belonged to Sordariomycetes (87%) and Dothideomycetes (13%) were obtained. Molecular characterization of isolated endophytic fungi with GenBank accession numbers, the closest match of ITS sequence in the NCBI database and their sequence similarity is depicted in Table 1. Among the isolated endophytic fungi, seven different species of *Fusarium* and two species of *Colletotrichum* were recovered. Five species of *Fusarium* viz., *Fusarium begoniae*, *Fusarium commune*, *Fusarium culmorum*, *Fusarium tricinctum*, *Fusarium babinda* were associated with the stem segments. In addition, *Nigrospora sphaerica* and *Curvularia coicis* were also recovered from the stem segments.

Neocosmospora haematococca, *Plectosphaerella cucumerina*, *Colletotrichum fruticola* exhibited tissue specificity by colonizing the leaf segments, while *Fusarium petroliphilum*, *Endomelanconiopsis endophytica* were isolated from bark segments. *Colletotrichum gloeosporioides* colonized both leaf and the stem fragments, whereas *Fusarium* sp. and

Trichoderma hamatum showed colonization in all plant segments. The percent colonization frequency of endophytic fungi in bark, leaf and stem parts along with their isolation code are represented in Table 2. The highest percent colonization was exhibited by *Fusarium* sp. (15.3) followed by *T. hamatum* (14.7) whereas *F. commune* showed the least percent colonization of 2.0. All endophytes from *T. heyneana* were assembled into eight different genera and assigned to Nectriaceae, Glomerallaceae, Plectosphaerellaceae, Hypocreaceae, Pleosporaceae, Botryosphaeriaceae and Trichosphaeriaceae.

3.2. Qualitative analysis of endophytic fungi for L-asparaginase activity

Of the 15 fungal endophytes screened for enzyme activity by preliminary plate assay method, 11 could grow on MCD agar producing pink zone with phenol red, a dye indicator that changes from yellow to pink (acidic to alkaline condition). The formation of pink zone around each fungal colony specified variations in pH due to the accumulation of ammonia in the medium.

3.3. Quantitative analysis by Nesslerization

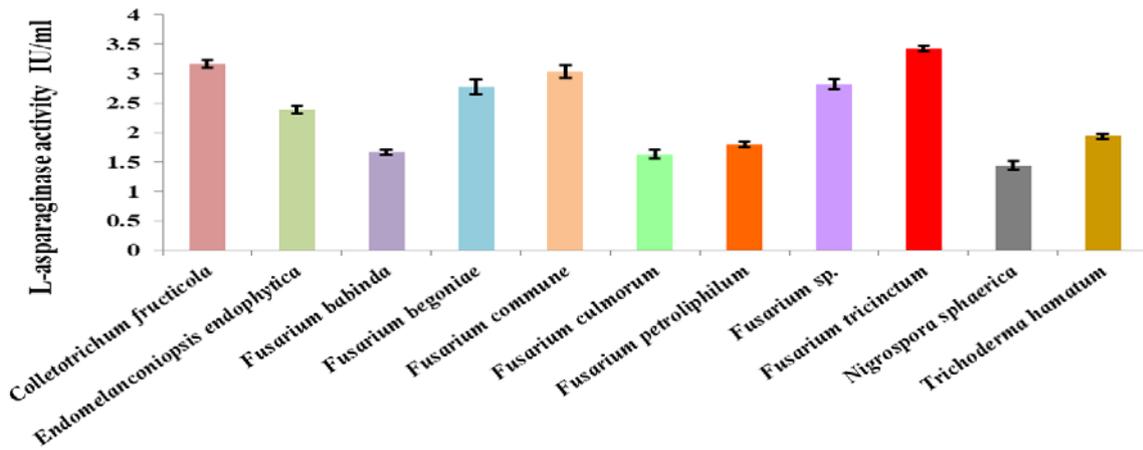
Positive isolates from preliminary screening were further evaluated by Nesslerization using spectrophotometric methods. Eleven positive isolates exhibited asparaginase activity in the range of 1.448 ± 0.07 to 3.423 ± 0.05 IU/ml (Figure 3). *F. tricinctum* from stem showed highest asparaginase activity among all the tested endophytic fungi with 3.423 ± 0.05 IU/ml of enzyme. *C. fruticola* from stem and *F. commune* from leaf also demonstrated high asparaginase activity of 3.166 ± 0.07 and 3.036 ± 0.11 IU/ml respectively. *Fusarium* sp., *F. begoniae* and *E. endophytica* showed moderate L-asparaginase activity in the range of 2.827 ± 0.09 – 2.388 ± 0.06 IU/ml of enzyme.

3.4. Determination of total phenolic content

Total phenolic content of all the fungal extracts ranged from 18.14 ± 8.62 to 90.41 ± 4.56 mg GAE/g dry extract (Table 3). The extracts of *F. tricinctum* displayed highest TPC (90.41 ± 4.56 mg GAE/g dry extract) followed by *C. coicis* (88.99 ± 38.75 mg GAE/g dry extract) and *N. sphaerica* (87.52 ± 34.44 mg GAE/g dry extract).

3.5. Antioxidant activity

The antioxidant activity of the endophytic extracts was determined following DPPH radical scavenging activity. The free radical scavenging activity is represented as 50% scavenging activity (IC_{50}) and is depicted in Table 3. The fungal extracts indicated IC_{50} values ranging from 51.64 ± 8.91 μ g/ml to 764.05 ± 27.67 μ g/ml. *F. tricinctum* extract depicted high scavenging activity with 51.64 ± 8.91 μ g/ml, followed by *F. commune* (124.16 ± 9.12) and *N. haematococca* (129.20 ± 15.12 μ g/ml).



Endophytic fungal isolates

Figure 3. L-asparaginase producers of endophytic fungal species isolated from *T. heyneana*.

Data are reported as mean±SD of three independent analyses (n=3).

Table 1. Molecular characterization of fungal endophytes isolated from *T. heyneana* with GenBank accession numbers, ITS sequence match in the NCBI database and their percent sequence similarity

Fungal endophytes	Isolate code	GenBank accession number	Closest match ITS	Sequence length (bp)	Sequencing similarity
<i>Colletotrichum gloeosporioides</i>	TH-WG-03	MK767027	KU682216	441	441/441 (100%)
<i>Colletotrichum fructicola</i>	TH-WG-11	MK976025	MK041495	516	507/517 (98%)
<i>Curvularia coicis</i>	TH-WG-12	MK976026	MG589634	555	532/553 (96%)
<i>Endomelanconiopsis endophytica</i>	TH-WG-13	MK991793	MK075024	527	527/527 (100%)
<i>Fusarium tricinctum</i>	TH-WG-04	MK752630	MK102656	504	495/504 (98%)
<i>Fusarium sp.</i>	TH-WG-06	MK776870	MH935958	293	290/290 (100%)
<i>Neocosmospora haematococca</i>	TH-WG-07	MK911734	KX099641	512	459/464 (99%)
<i>Nigrospora sphaerica</i>	TH-WG-15	MN066344	MF380852	493	484/500 (97%)
<i>Plectosphaerella cucumerina</i>	TH-WG-08	MK940863	MH063586	499	485/487 (99%)
<i>Trichoderma hamatum</i>	TH-WG-09	MK940897	MK304047	551	553/553 (100%)
<i>Fusarium babinda</i>	TH-WG-10	MK940898	MH862578	404	381/406 (94%)
<i>Fusarium begoniae</i>	TH-WG-01	MK720625	KM577645	483	475/483 (98%)
<i>Fusarium commune</i>	TH-WG-02	MK723994	KX878889	473	473/473 (100%)
<i>Fusarium culmorum</i>	TH-WG-05	MK767026	KU375665	468	468/468 (100%)
<i>Fusarium petroliphilum</i>	TH-WG-14	MN066343	LS999414	506	501/509 (98%)

Table 2. Colonization frequency of endophytic fungi associated with *T. heyneana*

.Endophytic strains	Isolate code	Plant parts						Total % CF
		Bark*		Leaf*		Stem*		
		I	% CF	I	% CF	I	% CF	
<i>Colletotrichum gloeosporioides</i>	TH-WG-03	-	-	29	14.5	3	1.5	5.3
<i>Colletotrichum fructicola</i>	TH-WG-11	-	-	17	8.5	-	-	2.9
<i>Curvularia coicis</i>	TH-WG-12	-	-	-	-	17	8.5	2.9
<i>Endomelanconiopsis endophytica</i>	TH-WG-13	35	17.5	-	-	-	-	5.9
<i>Fusarium tricinctum</i>	TH-WG-04	-	-	-	-	36	18	6.0
<i>Fusarium sp.</i>	TH-WG-06	8	4.0	75	37.5	9	4.5	15.3
<i>Neocosmospora haematococca</i>	TH-WG-07	-	-	18	9.0	-	-	3.0
<i>Nigrospora sphaerica</i>	TH-WG-15	-	-	-	-	19	9.5	3.1
<i>Plectosphaerella cucumerina</i>	TH-WG-08	-	-	15	7.5	-	-	2.5
<i>Trichoderma hamatum</i>	TH-WG-09	49	24.5	32	16	7	3.5	14.7
<i>Fusarium babinda</i>	TH-WG-10	-	-	-	-	27	13.5	4.5
<i>Fusarium begoniae</i>	TH-WG-01	-	-	-	-	13	6.5	2.1
<i>Fusarium commune</i>	TH-WG-02	-	-	-	-	12	6.0	2.0
<i>Fusarium culmorum</i>	TH-WG-05	-	-	-	-	24	12	4.0
<i>Fusarium petroliphilum</i>	TH-WG-14	28	14.0	-	-	-	-	4.7

* 200 segments were plated from bark, leaf and stem respectively for frequency analysis. I: Number of isolates; CF-Colonization frequency; '-': Not detected

Table 3. Estimation of total phenolic content and radical scavenging potentials of *T. heyneana* endophytes

Fungal strains	Total phenolic content (mg GAE/ g dry extract)	DPPH radical scavenging capacity (IC ₅₀ µg/ml)
<i>C. gloeosporioides</i>	37.82±25.82	145.48±14.73
<i>C. fruticola</i>	33.51±16.02	413.36±26.40
<i>C. coicis</i>	88.99±38.75	374.15±30.50
<i>E. endophytica</i>	64.77±11.39	736.04±31.65
<i>F. babinda</i>	25.30±3.01	175.25±21.35
<i>F. begoniae</i>	37.87±3.44	138.68±5.71
<i>F. commune</i>	76.84±16.20	124.16±9.12
<i>F. culmorum</i>	35.96±21.52	378.04±17.48
<i>F. petroliphilum</i>	78.35±3.80	764.05±27.67
<i>F. tricinctum</i>	90.41±4.56	51.64±8.91
<i>Fusarium</i> sp.	18.14±8.62	336.03±22.71
<i>N. haematococca</i>	17.29±12.92	129.20±15.12
<i>N. sphaerica</i>	87.52±34.44	155.26±33.71
<i>P. cucumerina</i>	31.25±3.88	670.37±26.51
<i>T. hamatum</i>	68.65±12.62	481.24±17.99

Abbreviations: DPPH – 1,1-diphenyl-2-picrylhydrazyl; IC₅₀ – Inhibitory concentration 50%. Data are reported as mean ± SD of three independent analyses.

4. Discussion

The Western Ghats of India has an exceptionally high level of biological diversity and is recognized as one of the world's 'hot spots' of biological diversity. *T. heyneana* is one of the medicinally important species of apocynaceae known to possess active pharmacological principles. This study has aimed to isolate endophytic fungi from *T. heyneana* and evaluate their L-asparaginase activity, total phenolic content and antioxidant activity.

Fifteen endophytic fungal strains such as species of *Fusarium*, *Colletotrichum*, *Neocosmospora*, *Plectosphaerella*, *Trichoderma*, *Curvularia*, *Endomelanconiopsis* and *Nigrospora* were identified using ITS sequencing method. The occurrence of seven different *Fusarium* species is remarkable as these have adapted to a range of geographical sites, climatic conditions, ecological habitats and host plants. Researchers suggest that there are similar positive interactions of endophytic *Fusarium* species with plants (Blok and Bollen, 1995). *Fusarium* sp. and *Colletotrichum* sp. are also reported as endophytes from earlier studies in *T. heyneana* (Manasa and Nalini, 2014). In our quest to identify fungal endophytes from medicinal species of *Rauvolfia*, both *Fusarium* and *Colletotrichum* spp. were documented as endophytes (Bhavana *et al.*, 2019). Preceding findings also reveal the evaluation of *T. divaricata* for the isolation of endophytes like *Colletotrichum*, *Gliocladium*, *Mycelia sterilia*, *Phoma*, *Phomopsis*, *Xylaria* sp. (Huang *et al.*, 2008) as well as for arbuscular mycorrhizal and dark septate endophyte fungal associations (Debnath *et al.*, 2015).

It was interesting to document that some of the fungal genera such as *Plectosphaerella*, *Trichoderma*, *Curvularia*, *Endomelanconiopsis* and *Nigrospora* were not previously reported from *T. heyneana*. The variation in the colonization potential of the endophytic fungi shows that, each endophyte exhibited different degrees of affinity

towards different tissues of the plant. *Fusarium* sp., and *Trichoderma* showed colonization in leaf, stem and bark tissues while the remaining endophytes were specific to tissue types.

Of the 15 fungal endophytes, the highest percent colonization was recorded in *Fusarium* sp. (15.3) followed by *T. hamatum* (14.7) whereas *F. commune* showed low percent colonization (2.0). *Fusarium* spp. as endophytes are documented from *T. heyneana* (Manasa and Nalini, 2014), *Catharanthus roseus* (Ayob and Simarani, 2016), Himalayan Yew plants (Garyali *et al.*, 2013) and from Malaysian anticancer plants (Chow and Ting, 2015). Similarly, *Trichoderma* spp. as endophytes have been recorded from three medicinal plants ie., *Solanum surattense* (Ikram *et al.*, 2019), *Salvia miltiorrhiza* (Zhou *et al.*, 2019) and *Elettaria* (Munir *et al.*, 2019).

The fungal strains were investigated for L-asparaginase production, total phenolic content and antioxidative potentials. *T. heyneana* revealed the presence of diverse endophytic fungal species with L-asparaginase activity. Of the 15 endophytic strains screened, 11 were positive for L-asparaginase. *F. tricinctum* isolated from stem fragments showed high asparaginase activity among all the tested fungi (3.423±0.05 IU of enzyme).

Fusarium endophytes differ in their ability to produce L-asparaginase. *F. verticilloides* from *T. heyneana*, showed considerably less enzyme activity of 1.136 IU/ml (Manasa and Nalini (2014). *Fusarium* spp. are known producers of L-asparaginase. *F. oxysporum* with 27.10 units/gram dry substrate (Pallem, 2019), *F. equiseti* with 8.51 IU (Hosamani and Kaliwal, 2011), *F. culmorum* with 7.21 units/gram dry substrate (Meghavam and Janakiraman, 2017), *Fusarium* sp. with 11.4 U/min/ml (Gonçalves *et al.*, 2016), *F. solani* and *F. oxysporum* with 0.04 IU/ml and 0.08 IU/ml respectively were also reported for their asparaginase activity (Gulati *et al.*, 1997). Our results are on par with the original work of Theantana *et al.* (2009), wherein quantitative activity for 53 positive/82 strains of endophytes ranged from 0.014 to 1.5 Units/ml. It is interesting to note that *Trichosporon asahii* IBBLA1 isolated from soil and moss samples from Schirmacher hills, Antarctica regions exhibited significant enzyme activity of 20.57 IU/ml (Ashok *et al.*, 2019).

In the present investigation, high TPC of 90.41± 4.56 mg GAE/g dry extract, was detected in *F. tricinctum*. *T. heyneana* is a well-known Indian medicinal plant with various phytochemicals. The total phenolic contents of fresh leaf extract of *T. heyneana* contained 11.4± 0.17 mg GAE/g dry extract (Sathishkumar and Baskar, 2012), whereas in the methanol extract, the TPC was 14.0± 0.45 mg GAE/g dry extract (Manasa and Chandrashekar, 2015). On the basis of these observations, it can be opined that the phenolic content of endophytic fungal extracts are high in comparison with the host plants.

Because of their capability to secrete bioactive metabolites, endophytic fungi can also reduce the oxidative stress (Khan *et al.*, 2017). The anti-oxidative function of endophytic fungi is due to the secretion of phenolic compounds. Of the 15 endophytic fungal strains screened for the antioxidant activity, *F. tricinctum* was found have high antioxidative potentials with IC₅₀ value of 51.64 ± 8.91 µg/ml., whereas the DPPH radical scavenging activity of *T. heyneana* carried out by Baskar *et al.* (2012), Sathishkumar and Baskar (2012) showed IC₅₀ values of

340.17 ±26.04 µg/ml and 507 µg/ml respectively, lesser in comparison with the current study.

Endophytic *F. tricinctum* with unique bioactivity from several medicinal species are reported viz., *Taxus baccata* (Vasundhara *et al.*, 2016) *Salicornia bigelovii* (Zhang *et al.*, 2015), *Aristolochia paucinervis* (Wätjen *et al.*, 2009) *Fritillaria unibracteata* var. *wabuensis* (Pan *et al.*, 2017a), *F. cirrhosa* (Pan *et al.*, 2017b), grasses (Przemieniecki *et al.*, 2019) and *Paris polyphylla* var. *yunnanensis* (Zhang *et al.*, 2011).

5. Conclusion

This study explores the endophytic fungi from *T. heyneana* and their L-asparaginase, total phenolic content and antioxidative potentials. The increasing importance of L-asparaginase in recent years due to its anti-carcinogenic applications encouraged us to screen the extracts of endophytes for L-asparaginase activity. *F. tricinctum* displayed significant L-asparaginase activity with remarkable total phenolic content and antioxidative capacities. Our study has shown that fungal endophytes from *T. heyneana* have potential bioactivities. The extracts are being evaluated for their anti-proliferative potentials in *in vitro* cancer cell lines.

Conflict of interest

The authors declare that there is no conflict of interest

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