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Distribution and diversity of foliar endophytic fungi in the mangroves of Andaman Islands, India



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ABSTRACT

Fungal endophytes represent a major component of plant microbiomes. Various aspects of these fungi such as their diversity and technological potential have been studied in detail. However, their distribution and diversity in a mangrove community has not been addressed. In this study, we report the presence of culturable fungal endophytes from 20 obligate mangrove hosts from south Andaman Islands. *Phomopsis/Diaporthe* was isolated from all the mangrove species studied while *Xylaria*, *Colletotrichum* and *Phyllosticta* were recorded from the majority of the mangroves studied. A phylogenetic analysis of representative *Phomopsis/Diaporthe* isolates clearly indicated the broad host range of this genus. Our study also highlighted the fact that leaf endophytes of mangroves are not unique with reference to their species diversity and frequency of occurrence when compared to those of terrestrial plants. These observations suggest that the extraordinary success of some fungal endophytes in colonizing taxonomically disparate hosts could be due to development of traits specific to their ecosystem.

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1. Introduction

Endophytic fungi infect and live within plant tissues without inducing any disease. They are ubiquitous and constitute an integral component of the plant microbiome. Endophyte association increases plant host fitness by enhancing its tolerance to abiotic (Yamaji et al., 2016; Lata et al., 2018) and biotic (Estrada et al., 2015; Suryanarayanan et al., 2018) stressors; these filamentous fungi are thus being investigated for their potential to improve crop fitness (Vega et al., 2008; Raghavendra and Newcombe, 2013). Furthermore, endophytic fungi represent a source of novel bioactive molecules (Suryanarayanan et al., 2009; Kharwar et al., 2011; Kusari et al., 2012) and industrially important enzymes (Thirunavukkarasu et al., 2011, 2015; Suryanarayanan et al., 2012; Sengupta et al., 2017). There are many studies on endophytes residing in the leaves of individual angiosperm plants and a few of them address the status of the foliar endophyte of plant communities (Suryanarayanan et al., 2003, 2011; Arnold and Lutzoni, 2007; Sudhakara Reddy et al., 2016). Here, we report on the distribution

and diversity of culturable foliar endophytes of mangroves of Andaman Islands, India.

Mangroves are plants of the tidal habitats and survive in the ecotone between the terrestrial and marine ecosystems. Mangroves constitute an unique ecosystem which provides crucial ecosystem services including fisheries, shoreline shield, carbon sequestration and bioremediation of wastes (Lee et al., 2014). Furthermore, mangrove forests support a wide biodiversity and constitute the most carbon rich forests of the tropics (Donato et al., 2011). Current satellite data confirm that anthropogenic activity is leading to the loss of mangrove forests globally (Thomas et al., 2017).

While macroscopic basidiomycetes (Gilbert and Sousa, 2002; Maekawa et al., 2003; Sakayaroj et al., 2012; Nogueira-Melo et al., 2014) and marine fungi (Sarma and Hyde, 2001) of mangroves have been studied for their diversity and distribution at the community level, investigations on endophytes of mangroves pertain only to a few mangrove species (Suryanarayanan et al., 1998; Kumaresan and Suryanarayanan, 2001; Costa et al., 2012; de Souza Sebastianes et al., 2013; Li et al., 2016) or to their ability to produce novel bioactive metabolites and extracellular enzymes (Maria et al., 2005; Aly et al., 2010; Debbab et al., 2013). To our knowledge, there are no studies involving simultaneous sampling of many mangrove species to address the diversity and distribution of foliar



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endophytes in them. Furthermore, according to Niranjan and Sarma (2018) there are no studies on the endophytes of Andaman and Nicobar Islands. We chose to investigate this facet of endophytes of Andaman Islands (in the Bay of Bengal, east of India, 11.7401° N, 92.6586° E) since 13% of the total mangrove cover of India is present here (Forest Survey of India, 2013), and the mangroves of this region exhibit the highest density and growth among the mangroves of the country (Dagar et al., 1991).

2. Materials and methods

2.1. Sample collection

We chose South Andaman for our study as it supports 23 of the 25 mangrove species and all the 10 mangrove families distributed in the Andaman and Nicobar Islands (Goutham-Bharathi et al., 2014). Mature and healthy leaves of 20 obligate mangrove species (Goutham-Bharathi et al., 2014) belonging to 10 families were collected from the following seven different locations in South Andaman Island (Fig. 1, Table 1) and sampled for their endophyte presence - Burmanallah, Chidiyatapu, Corbyn's Cove, Manjeri, Shippighat, Shoal Bay and Wright Myo. The leaves were transported to the laboratory in sterile bags and screened for endophyte presence within 48 h of collection, by surface sterilizing and plating them on nutrient agar medium. The study involved a one time sampling between May 2016 and February 2017.

2.2. Surface sterilization

For each mangrove species, a total of 40 leaves were collected from 4 individual plants (10 leaves/individual). From these, 120 tissue segments $(0.5 \text{ cm}^2 \text{ each})$ were cut from the midrib region (including the lamina portion) -one each from the apical, middle and the basal region of the leaf. Of these, 100 leaf segments were surface sterilized and screened. These leaf segments were surface sterilized by immersing them in 70% ethanol for 5 s, followed by treatment with sodium hypochlorite (4% available chlorine) for 90 s and rinsing in sterile distilled water for 10 s (Suryanarayanan et al., 1998). The tissue segments were then plated in Petri dishes (9 cm dia.) containing antibiotic-amended (Chloramphenicol, 150 mg/l) Potato Dextrose Agar (PDA) medium (20 ml). We had previously used this sampling design for studying leaf endophytes of trees of different forest types occurring in the Western Ghats (Suryanarayanan et al., 2002, 2011; Govinda rajulu et al., 2013; Sudhakara Reddy et al., 2016).

2.3. Incubation procedure and isolation of endophytes

The efficacy of the sterilization protocol in removing the surface microbes was confirmed (Schulz et al., 1998) and each Petri dish with ten leaf segments was incubated in a light chamber (12 h light: 12 h dark cycle, 2200 lux of light) at 26 °C for 4 weeks (Suryanarayanan, 1992). The endophytes which grew out of the tissue segments were isolated, cultured on PDA slants and identified using standard manuals (Barnett and Hunter, 1998; Ellis, 1971, 1976; Ellis and Ellis, 1988; Sutton, 1980; Onions et al., 1981). Isolates which failed to sporulate were given codes based on culture characteristics such as growth rate, colony surface texture and hyphal pigmentation (Suryanarayanan et al., 1998) and were assumed to be different taxonomic species (Bills and Polishook, 1994). Since the identification was done primarily based on spore morphology, the anamorph (asexual state) and teleomorph (sexual state) were enumerated separately.



Fig. 1. Map of Andaman Islands showing the places of sample collection.

2.4. Genomic DNA extraction

The genomic DNA was extracted from fresh mycelia collected from 7 d old cultures growing in PDA medium. For this, the phenolchloroform method was used (Sudhakara Reddy et al., 2016). The extracted DNA was resuspended in 50 μ l of sterile distilled water and stored at -80 °C for further studies. For PCR amplification, the samples were thawed, concentration of the genomic DNA was checked in a 1% agarose gel and then used.

2.5. PCR amplification and sequencing of ITS region

A polymerase chain reaction was performed to amplify the ITS region and its flanking sequences. Fungal specific primers ITS4 and ITS5 were used for the reaction (White et al., 1990). The PCR reaction mix consisted of PCR buffer, forward and reverse primers, dNTPs, Taq Polymerase, DMSO, MgCl₂ and fungal DNA and was

 Table 1

 Mangrove species of Andaman Islands, India studied for foliar fungal endophytes.

Code	Mangrove Plant	Family	Collection Site	Co-ordinates
AE	Acanthus ebracteatus Vahl	Acanthaceae	Shippighat	11° 39' 52.3368" N92° 44' 11.04" E
AI	Acanthus ilicifolius L.	Acanthaceae	Wright myo	11° 47' 19.0896" N92° 43' 34.3884" E
AC	Aegiceras corniculatum (L.) Blanco	Myrsinaceae	Corbyn's Cove	11° 38' 40.7976" N92° 44' 51.3636" E
AM	Avicennia marina (Forssk.) Vierh.	Avicenniaceae	Burmanallah	11° 33' 27.3924" N92° 43' 47.3448" E
AO	Avicennia officinalis L.	Avicenniaceae	Corbyn's Cove	
BC	Bruguiera cylindrica (L.) Blume	Rhizophoraceae	Chidiyatapu	11° 30' 21.9564" N92° 42' 6.0948" E
BG	Bruguiera gymnorhiza (L.) Lam.	Rhizophoraceae	Burmanallah	
BP	Bruguiera parviflora (Roxb.) Wight &Arn. exGriff.	Rhizophoraceae	Shoal Bay	11° 53' 52.53" N92° 46' 32.1204" E
CT	Ceriops tagal (Perr.) C.B. Rob.	Rhizophoraceae	Manjeri	11° 32' 33.36" N92° 39' 8.7588" E
EA	Excoecaria agallocha L.	Euphorbiaceae	Burmanallah	
LL	Lumnitzera littorea (Jack) Voigt	Combretaceae	Shoal Bay	
LR	Lumnitzera racemosa Willd.	Combretaceae	Manjeri	
NF	Nypa fruticans Wurmb	Arecaceae	Shippighat	
PP	Phoenix paludosa Roxb.	Arecaceae	Shippighat	
RA	Rhizophora apiculata Blume	Rhizophoraceae	Burmanallah	
RM	Rhizophora mucronata Lam.	Rhizophoraceae	Burmanallah	
RS	Rhizophora stylosa Griff.	Rhizophoraceae	Chidiyatapu	
SH	Scyphiphora hydrophyllacea C.F. Gaertn.	Rubiaceae	Shoal Bay	
SA	Sonneratia alba Sm.	Lythraceae	Wright myo	
XG	Xylocarpus granatum J. Koenig	Meliaceae	Burmanallah	

carried out in a 25 µl reaction volume. The reaction conditions were: 94 °C for 3 min; 34 cycles consisting of 94 °C for 30 s, 54 °C for 30 s and 72 °C for 60 s; followed by 72 °C for 10 min. The PCR amplicons were run in agarose gel and gel eluted for purification. The sequencing reaction was carried out using one of the primers and sequenced in ABI 3130 Genetic Analyzer. After manual editing, an initial search was carried out using BLAST to identify the closest matches. The sequences obtained in the current study were then aligned with sequences of type materials and sequences showing higher similarity in Blast search using ClustalW. The aligned sequences were then manually realigned again and the phylogenetic tree was constructed using MEGA6 (Tamura et al., 2013). A maximum likelihood analysis was performed to understand the phylogenetic relationship between the isolates. A bootstrap analysis with 1000 replicates was performed and the consensus tree was constructed using MEGA6. The sequences were submitted to GenBank and accession numbers were obtained (MH371241-MH371256).

2.6. Statistical analysis

The colonization frequency (CF%) of each endophyte was calculated by the method of Hata and Futai (1995).

$\label{eq:CF} CF\% = \frac{\text{Number of segments colonized by each endophyte}}{\text{Total number of segments observed}} \\ \times 100$

Fisher's α was used to estimate the species diversity of endophytes. The formula for estimating this value is S = a*ln (1 + n/a), where S is the number of taxa, n is the number of individuals, ln is the natural logarithm and a is the Fisher's alpha. This index was preferred as it has been shown to be less affected by the abundance of common species (Magurran, 2004). Statistical programmes Pro version 2 (The National History Museum and The Scottish Association for Marine Science) and EstimateS software version 9.1.0 (Robert K. Colwell, University of Connecticut) [http://viceroy.eeb. uconn.edu/estimates] were used for calculating the ecological parameters. To avoid the influence of sample sequence on the progress of species accumulation, unique species and singleton curves, the data were randomized 100 times before analysing and plotting the curves (Suryanarayanan et al., 2011).

3. Results

The number of endophyte isolates that were recovered varied from 39 in Avicennia marina to 236 in Excoecaria agallocha; the number of endophyte species isolated ranged from 8 in Bruguiera parviflora to 23 in Rhizophora apiculata (Supplementary Table 1). The species diversity of the endophytes was lowest for *B. parviflora* (2.1) and highest for *R. apiculata* (11.4) (Supplementary Table 1). Phomopsis was the most common endophyte genus recorded and was isolated from all the 20 mangrove species screened; it accounted for 377 isolates of the total 2180 isolates obtained from all the mangroves and hence isolates belonging to this genus were selected for further molecular characterisation (Fig. 2, Table 2). Species of Xylaria and Colletotrichum were present in the leaves of 19 mangroves. Phyllosticta capitalensis was the densest endophyte as 545 isolates of it were isolated from 18 mangrove species. Colletotrichum gloeosporioides, Phyllosticta capitalensis, Phomopsis spp. (along with the teleomorph Diaporthe spp.), or Xylaria spp. (along with the anamorph Nodulisporium spp.) dominated or occurred as co-dominant species in the endophyte assemblage of different mangrove species studied (Table 3). C. gloeosporioides dominated the endophyte assemblages of AE and XG (abbreviations are expanded in Table 1), P. capitalensis was dominant in AI, BG, EA, LL, LR and SH and co-dominat in AE and SA, Phomopsis spp. was dominant in AC, AM, AO and RM and co-dominant in AI, BG, LR, PP, RA and RS, Xylaria spp. was dominant in BC, BP, CT, NF, PP and RS and co-dominant in AM, AO, BC, BP, CT, LL, RM, SH and XG (Table 3). An ITS sequence analysis of the 16 Phomopsis/Diaporthe isolates showed that these belonged to 10 different species (Table 2). Diaporthe pseudomangiferae was endophytic in the leaves of AC, AM, AO and CT. D. discoidispora was present in BP, LL, and RM while D. hongkongensis was isolated from NF and PP. D. longicolla, D. kyushuensis, Phomopsis heveicola, D. eucalyptorum, D. liquidambaris, D. perseae and D. tectonae were isolated from the leaves of AI, BC, BG, LR, SH, SA, and XG respectively. A bootstrap consensus tree constructed based on maximum likelihood approach showed that the Phomopsis/Diaporthe isolates could be grouped into several distinct clades (Fig. 2). Many species of Phomopsis/Diaporthe were isolated from different mangroves indicating very little host specificity among the endophytic isolates. We also could not discern any clusters based on the location from which these isolates were obtained. Statistical analyses revealed that while the number of endophyte isolates increased with increasing sample size, the





Fig. 2. Bootstrap consensus tree based on Maximum Likelihood method for *Phomopsis/ Diaporthe* isolates obtained in the study. Black squares indicate sequences obtained in the present study, open squares represent type sequences while open triangles represent other sequences.

Phomopsis/Diaporthe species identified based on ITS-sequences and their GenBan
accession number (Refer Table 1 for host code).

Host Code	Isolate	GenBank Accession No.
AI	Diaporthe longicolla	MH371243
AC	Diaporthe pseudomangiferae	MH371242
AM	Diaporthe pseudomangiferae	MH371241
AO	Diaporthe pseudomangiferae	MH371244
BC	Diaporthe kyushuensis	MH371245
BG	Phomopsis heveicola	MH371246
BP	Diaporthe discoidispora	MH371247
СТ	Diaporthe pseudomangiferae	MH371248
LL	Diaporthe discoidispora	MH371249
LR	Diaporthe eucalyptorum	MH371250
NF	Diaporthe hongkongensis	MH371251
PP	Diaporthe hongkongensis	MH371252
RM	Diaporthe discoidispora	MH371253
SH	Diaporthe liquidambaris	MH371255
SA	Diaporthe perseae	MH371254
XG	Diaporthe tectonae	MH371256

numbers of endophyte species (Fig. 3), unique species and singletons (Fig. 4) decreased with increasing sample size.

4. Discussion

The leaves of all the mangrove species studied harboured culturable fungal endophytes. In community level investigations, the sample size should be large enough to inform the diversity of the organisms studied. The sample size in the present study was rigorous enough to represent the diversity of foliar endophytes of the mangrove community existing at the time of sampling. This is borne out by the observation that the accumulation of endophyte species, as well as the unique and singleton species among them, although increasing initially, started to decrease with increasing sample size (Longino, 2000; Henderson, 2003; Suryanarayanan et al., 2011). The total CF% of the endophytes (which is also equal to the total number of isolates since 100 tissue segments were screened for each mangrove species) was higher than 100% in nine mangrove species due to the growth of more than one endophyte species from a tissue segment. This indicated that the density of colonization of the leaves by endophytes is high.

Leaf endophytes enhance the tolerance of terrestrial host plant to pathogen (Arnold et al., 2003) and herbivores (Estrada et al., 2015). Mangrove endophytes have not been investigated for such a role in biotic stress tolerance of mangroves. Furthermore, as speculated by Suryanarayanan (2013), dense colonization of leaves by endophytes could affect photosynthesis negatively by draining photosynthates or positively by creating localized zones of low photorespiration in the leaf due to their respiration. This aspect gains importance as the air in mangrove canopy is low in CO₂ and its microclimate is different from that of the terrestrial plants due to tidal inundation (Al-Saidi et al., 2009).

In the present study, species of *Colletotrichum, Phomopsis, Phyllosticta* and *Xylaria* were more commonly isolated as endophytes and were present in different mangrove species. These fungi have a wide host range as foliar endophytes and have been reported from mangroves of Hong Kong (Pang et al., 2008), Thailand (Chaeprasert et al., 2010), Brazil (Wanderley et al., 2012; de Souza Sebastianes et al., 2013) and China (Li et al., 2016). These fungi as endophytes are known to have loose plant host affiliation and occur even in taxonomically unrelated terrestrial plants distributed throughout the world (Pandey et al., 2003; Murali et al., 2006; Govinda rajulu et al., 2013; Unterseher et al., 2016). Although, low host specificity is a rule among several guilds of tropical fungi including the endophytes (Suryanarayanan, 2011), of particular

Dominant and Co-dominant endophyte species present in the leaves of mangrove plants of Andaman Islands, India (Refer Table 1 for host code).

Host Code	Dominant endophyte		Co-dominant endophyte	Co-dominant endophyte	
	Name	Order	Name	Order	
AE	Colletotrichum gloeosporioides	Glomerellales	Phyllosticta capitalensis	Botryosphaeriales	
AI	Phyllosticta capitalensis	Botryosphaeriales	Diaporthe longicolla	Diaporthales	
AC	Diaporthe pseudomangiferae	Diaporthales	Sterile form 4	-	
AM	Diaporthe pseudomangiferae	Diaporthales	Nodulisporium sp. 1	Xylariales	
AO	Diaporthe pseudomangiferae	Diaporthales	Xylaria sp. 2	Xylariales	
BC	Xylaria sp. 1	Xylariales	Nodulisporium sp. 1	Xylariales	
BG	Phyllosticta capitalensis	Botryosphaeriales	Phomopsis heveicola	Diaporthales	
BP	Xylaria sp. 1	Xylariales	Nodulisporium sp. 2	Xylariales	
CT	Xylaria sp. 1	Xylariales	Nodulisporium sp. 1	Xylariales	
EA	Phyllosticta capitalensis	Botryosphaeriales	Graphium sp.	Microascales	
LL	Phyllosticta capitalensis	Botryosphaeriales	Xylaria sp. 1	Xylariales	
LR	Phyllosticta capitalensis	Botryosphaeriales	Diaporthe eucalyptorum	Diaporthales	
NF	Xylaria sp. 1	Xylariales	Penicillium sp. 1	Eurotiales	
PP	Nodulisporium sp. 1	Xylariales	Diaporthe hongkongensis	Diaporthales	
RA	Aspergillus fumigatus	Eurotiales	Phomposis sp. 2	Diaporthales	
RM	Diaporthe discoidispora	Diaporthales	Xylaria sp. 1	Xylariales	
RS	Xylaria sp. 1	Xylariales	Phomposis sp. 1	Diaporthales	
SH	Phyllosticta capitalensis	Botryosphaeriales	Xylaria sp. 1	Xylariales	
SA	Pestalotiopsis sp.	Amphisphaeriales	Phyllosticta capitalensis	Botryosphaeriales	
XG	Colletotrichum gloeosporioides	Glomerellales	Xylaria sp. 1	Xylariales	



Fig. 3. Species accumulation curve for foliar endophytes isolated from twenty mangrove species. Dotted lines represent 95% confidence interval limits (upper and lower bounds) for species observed. Data were randomized 100 times for plotting the graph.

interest is the wide host range of *Diaporthe* (*Phomopsis*) spp. observed in the present study which is indicative of their ecological success. For instance, in our study, *Phomopsis* (*Diaporthe*) species were isolated from all the mangrove species. Of the 20 isolates from the 20 mangrove species, 4 (from AE, EA, RA and RS) failed to grow upon further sub-culturing and the other 16 were taken up for identification at the species level using ITS sequencing since morphological characters of this genus are unreliable (Gomes et al., 2013; Dissanayake et al., 2017). Interestingly we did not observe any host specificity among the *Phomopsis* isolates since the same species were isolated from different mangroves. This further attests to the loose host affiliation seen among major fungal endophytes, especially in tropical climates, and might be a major contributing factor in their extraordinary success as an endophyte in wide and disparate hosts.

The present survey involving more mangrove species than earlier studies confirms that leaf endophytes of mangroves are not unique with reference to their species diversity and frequency of occurrence when compared to those of terrestrial plants. Mangrove leaves accumulate salt as they mature (Cram et al., 2002) to levels which are 3–12 times higher when compared to terrestrial plants (Dissanayake and Amarasena, 2009). They are also rich in tannins (Maie et al., 2008) which are inhibitory to fungal growth (Harrison, 1971; Dix, 1979). The common foliar endophytes of mangroves are tolerant of high concentrations of both salt and tannins (Kumaresan et al., 2002). Selection of fungi with traits to survive in the salt and tannin-rich leaves of mangroves could be the reason for the wider ecological amplitude of certain endophyte genera in the mangrove forest.

Fungi in general, along with bacteria and oomycetes, are known



Fig. 4. Number of singleton and unique endophyte species isolated from twenty mangrove species. Data were randomized 100 times for plotting the graph. The curve represents a polynomial trendline.

to decompose mangrove litter (Kristensen et al., 2008). It is now established that foliar endophytic fungi remain in fallen leaves, switch to a saprotrophic mode (Unterseher et al., 2013; Guerreiro et al., 2017) and, owing to their ability to produce different extracellular enzymes, aid in litter decomposition (Suryanarayanan et al., 2012). Kumaresan and Suryanarayanan (2002) reported that the density of colonization of endophytic Glomerella sp. and Pestalotiopsis sp. in the leaves of the mangrove R. apiculata increases in fallen leaves with time and that these fungi produce biomass destructuring enzymes indicating their role in mangrove litter decomposition. Talaromyces stipitatus, a root endophyte in the mangrove Avicennia marina, elaborates salt-tolerant chitin modifying enzymes (Paranetharan et al., 2018). The specific role of mangrove endophytes in litter decomposition has to be addressed to understand their contribution to carbon dynamics in mangrove ecosystem. Since endophyte association confers salt tolerance in some plants, (Rodriguez et al., 2008; Khan et al., 2015), it could be worthwhile to investigate the contribution of endophytes to the fitness of mangroves.

To conclude, the mangrove ecosystem supports diverse microorganisms which perform various functions. The diversity and roles of archaea (Bhattacharyya et al., 2015) and bacteria (Basak et al., 2016; Chen et al., 2016) in this ecosystem are well established. With reference to fungi, only the manglicolous (Sahoo and Dhal, 2009) and mycorrhizal fungi (D'Souza, 2016) have been studied in detail. Considering the importance of endophytes in enhancing host plant's fitness and recycling nutrients, this cryptic ecological group of fungi of the mangroves deserves more attention to get a holistic view of the mangrove ecosystem.

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Supplementary data

Supplementary data to this article can be found online at https://doi.org/10.1016/j.funeco.2018.09.007.

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