

DIVERSITY OF ARBUSCULAR MYCORRHIZAL FUNGI IN SHOLA  
FORESTS OF KODAIKANAL

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# DIVERSITY OF ARBUSCULAR MYCORRHIZAL FUNGI IN SHOLA FORESTS OF KODAIKANAL

## INTRODUCTION

Mycorrhiza is the mutualistic symbiosis (non-pathogenic association) between certain soil-borne fungi and plant roots (Sieverding, 1991). Mycorrhizal fungi often connect plant root systems over broad areas through their mycelial networks. These fungi frequently comprise the largest portion of soil microbial biomass (Högberg and Högberg, 2002). Arbuscular mycorrhiza (AM) is the most widely distributed association in plants. About 80% of all terrestrial plant species form this type of symbiosis (Smith and Read, 1997). Plant roots have evolved to accommodate, utilize and control mycorrhizal fungi. Both molecular and fossil evidence indicate that the earliest land plants were mycorrhizal (Redecker *et al.*, 2000). Plants could not have colonized land without fungal partners capable of acquiring nutrients from the undeveloped soils that existed during the Silurian and Devonian (Pirozynski and Malloch, 1975). There are plants, however, that have been shown to be mycorrhiza free, such as Proteaceae (Nicholson, 1967; Brundrett *et al.*, 1996), Cruciferae, Zygophyllaceae (Varma, 1998) Dipterocarpaceae, Betulaceae, Myrtaceae and Fagaceae (Nicholson, 1967). The reason why some plants do not form mycorrhizas is not fully known, but it may be related to the presence of fungal toxic compounds in root cortical tissue or in root exudates. It may also be due to interactions between the fungus and the plant at the cell wall and (or) middle lamella level (Tester *et al.*, 1987). High concentrations of salicylic acid have been found to reduce mycorrhization (Medina *et al.*, 2003), meaning that plants with a genetic basis for high salicylic acid content have evolved to be non-mycorrhizal.

Arbuscular mycorrhizal morphology is distinguished into *Arum* – type and *Paris* – type. The *Arum*-type association is characterized by intercellular hyphal growth in the root cortex, with short lateral branches into cortical cells forming arbuscules (Smith and Smith, 1997). Intracellular–hyphal coils frequently having intercalary arbuscules spreading cell to cell in the cortex characterize the *Paris*– type association, has been found to be more frequent in natural ecosystems (Yamato and Iwasaki, 2002; Ahlu *et al.* , 2005; Tsuyuzaki *et al.* , 2005) .

Three main components are involved in AM association: the soil, the fungus and the plant. The fungal component involves the fungal structure within the cell of the root and the

extraradical mycelium in the soil. The last may be quite extensive under some conditions, but does not form any vegetative structures (Smith and Read, 1997). Its primary function is the absorption of resources from the soil. The increased efficiency of mycorrhizal roots versus non-mycorrhizal roots is caused by the active uptake and transport of nutrients by mycorrhizae.

AM fungal diversity is the major factor in the maintenance of plant biodiversity and ecosystem stability and function. Several studies indicate that AM fungi alter plant community structure by affecting the relative abundance of plant species and plant-species diversity (Grimme *et al.*, 1987; Gange *et al.*, 1990; Sanders and Koide 1994). Interplant transport of assimilates from the dominant canopy species via a common mycorrhizal network to subordinate plant species, has been suggested as a mechanism by which AM fungi affect the floristic diversity of plant communities (Grimme *et al.*, 1987). Another mechanism by which AM fungi may affect plant community structure is the differential growth response of plant species to colonization by AM fungi, the so called “mycorrhizal dependence” (Gederman, 1975; Plenchete *et al.*, 1983; Habate and Manjunath, 1991).

The species composition and diversity of AM fungal communities has the potential to determine plant population and plant community structure. The fact that plant species vary in the degree of response to AM fungal species has important implications for growth of individual plant species. In turn, this will affect a plant’s ability to coexist with other plant species in a community (Van der Heijden *et al.*, 1998). On the other hand, established mycorrhizal plants may serve as important sources of inoculum for initially non-mycorrhizal, conspecifics, which may affect regeneration and could contribute to patchy distributions of species within the community (Koide and Dickie, 2002).

Arbuscular mycorrhizal fungi occur in all kind of landforms including mountains (Shi *et al.*, 2007), plateaus (Pan *et al.*, 1997), hills, plains (Gai *et al.*, 2006), islands (Liu *et al.*, 2001) and basins (Wang *et al.*, 2006). Generally, the AM fungal spores are isolated from field soil and identified based on their morphology and sub cellular characters. More than 200 AM fungal species are described based on spore morphology (Schüßler *et al.*, 2001), but characterization of spore morphology requires considerable experience (Clapp *et al.*, 2001). Spore counts may not reflect the true composition of the AM fungal or plant communities (Turnau *et al.*, 2001).

This lack of reliability arises from the taxon-specific differences existing between sporulation and root colonization rates. The most common and widely distributed, AM fungal genus in the tropics is *Glomus*, followed by *Acaulospora* and *Scutellospora*. Therefore, information regarding the active AM fungi in roots is crucial for any ecological field studies. Understanding AM morphology with in roots at best allows discriminating AM fungi at the family or genus level (Merryweather and Fitter, 1998).

The unique combination of forests and grassland comprise the Shola forest. They are stunted evergreen forest found as patches in grasslands especially in Valleys. The Sholas are dark damp throughout the year, because Shola soil absorbs and retains water like a sponge. However, wide diversity and unique floral distribution, no systematic investigation has been carried out to explore the root fungal associations in plant species of shoal forests. When compare to other ecosystems, shoals are poorly explored for AM fungal distribution. In shoals of Western Ghats region, 29 plant species has been studied for AM fungal association (Bagyalakshmi et al., 2010). The root fungal associations of 107 medicinal and aromatic plant species have been assessed in Western Ghats region (Muthukumar et al., 2006). Six plant species in shoal forest of Velliangiri hills, western Ghats, Southern India, has been examined for AM fungal association and spore numbers (Muthukumar et al., 2018). Mycorrhizal status of sixteen epiphytic and terrestrial ferns has been explored from Kodaikanal Hills of Southern India (Raju et al., 1995). Arbuscular mycorrhizal association of 60 ferns and lycophytes were observed from Palni hills, Western Ghats region southern India (Muthukumar et al., 2014). These studies insist that the importance of mycorrhizal research which deserves much attention is the investigation of more plant species for their mycorrhizal status. In addition, the results of this investigation are primarily used for revegetation programs in shoal forest. The seasonal dynamics of AM fungi is essential to quantify the functioning and ecological significance of AM in natural ecosystems. The increase in the AM fungal spore numbers suggests a period in which the fungi act as a carbon sink (Smith and Read, 2008). Therefore the present investigation was carried out to fulfill the following objectives, (i) To assess the incidence and the types of AM association in shola plant species, (ii) To evaluate the arbuscular mycorrhizal (AM) fungal diversity, (iii) To record the seasonality of AM fungi in shola forest, (iv) To observe if any relationship between plant diversity and AM fungal diversity, (v) To identify the nutrient uptake mechanisms of shola species.

## **Materials and Methods**

### **Study site**

The study site, Kodaikanal (longitude 77° 26' to 77° 33' E and latitude 10° 12' to 10° 15'N) is located within the Eastern offshoot of the Western Ghats and the spur aligned on a east west and north south axis. The shoal forest is occupied by upper elevations of the Palani hills. Sholas are patches of jungles isolated from forests varied with plant species composition and size. The streams running through the shoals and trees showed stunted growth. The research was conducted among shoals with altitudes ranging from 360m – 2550m. The annual rainfall is quite variable in the hills (1300 mm) with temperatures ranging from 13 to 24°C in summer and winter ranged from 7 to 16°C in the summer.

### **Sampling**

Root and soil samples for each species were collected from five individuals at different stages of growth (vegetative and reproductive). Care was taken during collection that roots of shrubs and tree species could be positively identified. For this reason, samples of herbs were usually made by uprooting the plants. Roots were washed and stained within 24h or preserved in formalin acetic acid-alcohol before staining. Rhizosphere soil from roots and adjacent to plants was collected. Soil samples collected from different individuals of a species were mixed to form a composite sample. These composite soil samples were used for the isolation of AM fungal spores.

### **Preparation of roots and AM assessment**

Fixed roots were washed free of formalin acetic acid alcohol (5:5:90; V/V) (FAA) and examined under a dissection microscope (X 20) for AM fungal spores attached to roots. After examination, the roots were cut into 1-cm fragments, cleared in 2.5% KOH (Koske and Gemma 1989), acidified with 5 N HCl and stained with trypan blue (0.5% in lactoglycerol) overnight. Roots that remained dark after clearing were bleached in alkaline H<sub>2</sub>O<sub>2</sub> prior to the acidification. The stained roots were examined with a compound microscope (X 200–400) for AM fungal structures and the percentage of root length colonization was estimated according to the magnified intersection method (McGonigle et al. 1990). In addition, the number of hyphae, arbuscule and vesicle intersections were noted. It was thus possible to quantify both the root length colonized by AM structures and total root length colonization. Only species in which arbuscules found were considered to have arbuscular mycorrhizae.

The AM-morphology was classified as *Arum*- or *Paris*-type based on whether the fungal hyphae were present mainly as hyphae running through intercellular spaces or within cells as coils respectively following descriptions of Dickson (2004). Since we examined whole and squashed roots, we could not reliably distinguish among the intermediate sub-type morphologies as described and classified by Dickson (2004). However, wherever the parallel running hyphae were seen intracellular, the morphology was designated as the Intermediate-type.

### **Isolation, enumeration and identification of AM fungal spores**

The soil samples from rhizosphere soil were used for enumerating AM fungal spores. One hundred grams of the soil samples were dispersed in 1L water and the suspension was decanted through 710 to 37 $\mu$ m sieves. The residues in the sieves were washed into beakers. The sievates were dispersed in water and filtered through gridded filter papers. Each filter paper was then placed spread on a Petri dish and scanned under a dissection microscope at X40 magnification and all intact spores (non-collapsed spores with cytoplasmic contents and free from parasitic attack) were counted. Sporocarps and spore clusters were considered as one unit. The soils of the pot culture were used for identification of AM fungi. After isolation of the spores as described above, the intact spores were transferred using a wet needle to polyvinyl alcohol-lacto glycerol with or without Melzers reagent on a glass slide for identification. Spores were identified from spore morphology and sub cellular characters and compared to original descriptions (Schenck and Perez 1990). Spore morphology was also compared to the culture database established by INVAM (<http://invam.cag.wvu.edu>).

### **Trap cultures**

Rhizosphere soil samples were mixed thoroughly to form a composite soil sample. Two-liter capacity pots were filled with 1 L of thrice pasteurized (120°C for 60 min) sandy soil followed by 500ml of the mixed composite rhizosphere soil sample. The pots were seeded with *Eleusine coracana* (L.) Gaertn., and the seedlings were thinned to 5 seedlings per pot after germination. A total of 12 pots, were arranged in a randomized block design. At the end of the growth period the soil samples were taken from each pot and AM fungal spores were isolated by a modified wet-sieving and decanting method as detailed above.

### **Identification of AM fungal spores**

Intact and crushed spores in polyvinylalcohol-lactophenol and in Melzer's reagent were examined and identified according to Schenck and Perez (1990). Spore colour was examined under a dissection microscope on fresh specimens immersed in water. Classification, spore wall characters and the spelling of scientific names are as suggested by Morton and Benny (1990), Walker (1983, 1986) and Walker and Trappe (1993).

### **Life-history attributes and plant nomenclature**

Each plant species recorded during the survey was categorized for life-form and life-cycle attributes as determined from the literature (Parin 1981a,b; Toby and Hodd 1982; Nair and Henry 1983; Henry et al. 1987,1989) or field observations. Nomenclature and authorities are as used by Nair and Henry (1983) and Henry et al. (1987, 1989).

## **RESULTS**

### **Occurrence of AM association**

Of the 71 plant species (in 35 families) examined, all the families were colonized by AM fungi except two species in a genus *Psychotria*(Table 1). AM association was observed in members of supposedly non-mycorrhizal families Commelinaceae, Cleomaceae and Convolvulaceae. Only those species in which arbuscules or arbusculate coils were found were considered to have AM association. The fungal entry into roots was characterized by the presence of appressorium (Plate 1.). Further invasion of the roots varied depending upon the AM types.

### **AM morphology**

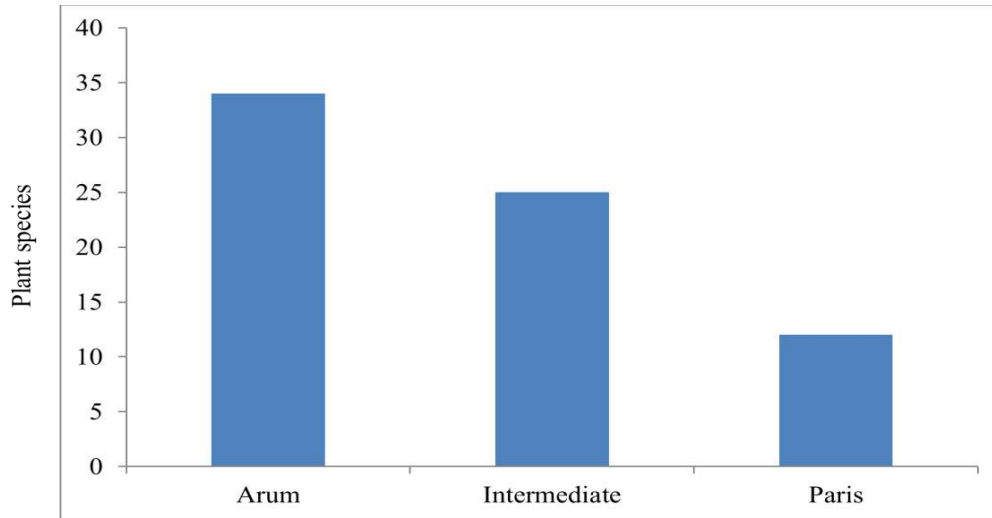
Thirty four of the plant species had *Arum*-type morphology, 25 had Intermediate- type and 12 had typical *Paris*-type morphology. The *Arum*-type was characterized by the presence of intercellular hyphae, vesicles and intracellular arbuscules. Intracellular hyphal coils, arbusculate coils and intracellular vesicles characterized the *Paris*-type morphology. The Intermediate-type had intracellular hyphal coils, as well as intercellular hyphae, arbuscules / arbusculate coils and inter / intracellular vesicles (Figure 1).

### **AM morphology in life forms**

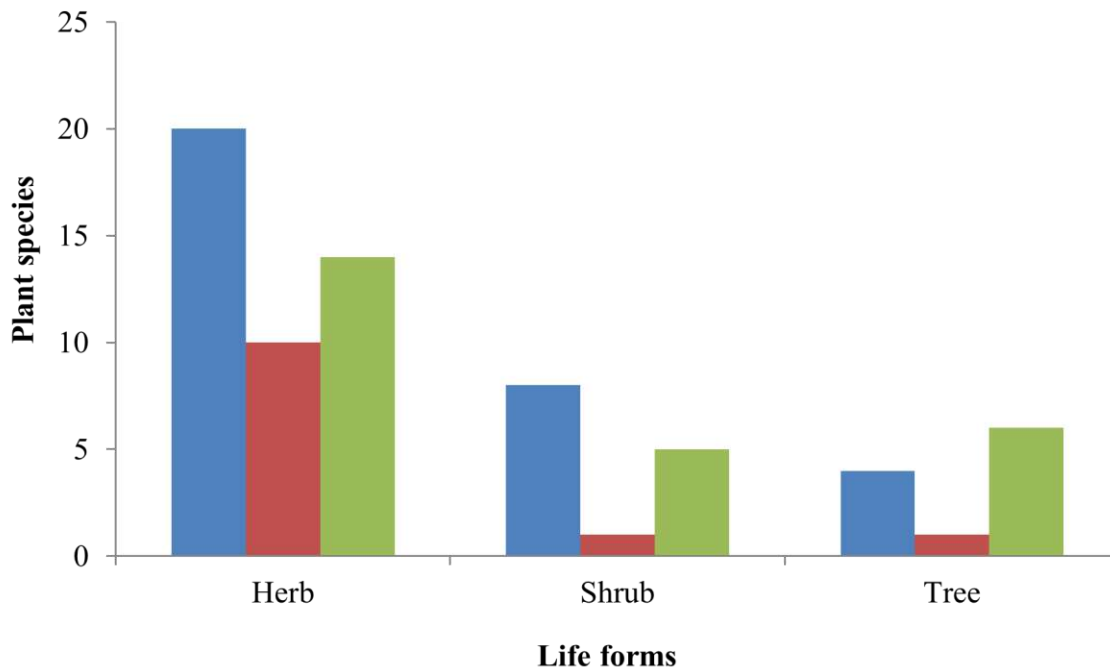
In herbs 20 species had Arum type morphology, 10 had Paris type morphology and 14 had Intermediate type morphology. In Shrubs, 8 species had Arum, one species had Paris and 5

species had Intermediate type morphology. In Tree species 4, 1 and 6 plant species had Arum, Paris and Intermediate type morphology respectively (Table 1; Figure 2).

**Figure 1. Arbuscular mycorrhizal fungal morphology in shoal plant species of Kodaikanal**



**Figure 2. Arbuscular mycorrhizal fungal morphology in various life forms of shoal species in Kodaikanal**





**Table 1.** Arbuscular mycorrhizal (AM) fungal morphology and colonization in shola plant species of Kodaikanal

<b>Family/Plant species</b>	<b>Habit</b>	<b>Fungal association</b>	<b>AM type</b>
<b>Acanthaceae</b>			
<i>Justicia adhatoda</i> F. Muell.	Shrub	AM	<i>Arum</i>
<i>Justicia txanquebariensis</i> Roxb.	Herb	AM	<i>Arum</i>
<i>Rungia repens</i> Nees.	Herb	AM	<i>Arum</i>
<i>Thunbergia fragrans</i> C. Presl,	Herb	AM	<i>Paris</i>
<i>Acorus calamus</i> L.	Herb	AM	<i>Paris</i>
<b>Amaranthaceae</b>			
<i>Achyranthus aspera</i>	Herb	AM	<i>Intermediate</i>
<i>Aerva lanata</i>	Herb	AM	<i>Arum</i>
<b>Hypoxidaceae</b>			
<i>Curculigo orchioides</i> Gaertn.,	Herb	AM	<i>Arum</i>
<b>Annonaceae</b>			
<i>Annona squamosa</i> L.,	Herb	AM	<i>Arum</i>
<b>Apiaceae</b>			
<i>Centella asiatica</i> (L.) Urb.	Herb	AM	<i>Paris</i>
<b>Apocynaceae</b>			
<i>Cascable thevetia</i>	Tree	AM	<i>Intermediate</i>
<i>Holarrhena antidysenterica</i>	Tree	AM	<i>Intermediate</i>
<b>Aristolochiaceae</b>			
<i>Aristolochia bracteolata</i>	Herb	AM	<i>Arum</i>
<i>Aristolochia indica</i>	Tree	AM	<i>Intermediate</i>
<b>Asclepiadaceae</b>			
<i>Gymnema sylvestre</i>	Herb	AM	<i>Intermediate</i>
<b>Asteraceae</b>			
<i>Anaphalis lawii</i> (Hook.f.) Gamble.	Herb	AM	<i>Intermediate</i>
<i>Ageratum conyzoides</i> L.,	Herb	AM	<i>Arum</i>
<b>Balsamiaceae</b>			
<i>Impatiens campanulata</i> Wight,	Herb	AM	<i>Intermediate</i>
<b>Begoniaceae</b>			
<i>Begonia malabarica</i> Buch	Herb	AM	<i>Paris</i>
<b>Caesalpinaceae</b>			
<i>Cassia fistula</i> L.,	Tree	AM	<i>Intermediate</i>
<i>Delonix regia</i> (Bojer) Raf.,	Tree	AM	<i>Arum</i>
<b>Cleomaceae</b>			
<i>Cleome gynandra</i> L.	Herb	AM	<i>Intermediate</i>
<b>Combretaceae</b>			
<i>Terminalia arjuna</i> (Roxb. Ex DC.) Wight & Arn.	Tree	AM	<i>Intermediate</i>

<b>Commelinaceae</b>			
<i>Commelina benghalensis</i> Wall.,	Herb	AM	Arum
<b>Convolvulaceae</b>			
<i>Evolvulus alsinoides</i> (L.) L	Herb	AM	Arum
<i>Ipomoea batatas</i> L.	Shrub	AM	Arum
<b>Cucurbitaceae</b>			
<i>Mukia leiosperma</i>	Shrub	AM	Intermediate
<b>Euphorbiaceae</b>			
<i>Acalypha indica</i> Vell.,	Herb	AM	Intermediate
<i>Euphorbia hirta</i> L.	Herb	AM	Intermediate
<i>Jatropha gossypifolia</i> L.	Shrub	AM	Arum
<i>Phyllanthus amarus</i> Schumach. & Thonn.	Herb	AM	Intermediate
<i>Phyllanthus maderaspatensis</i> Thouars ex Baill.	Herb	AM	Intermediate
<b>Labiatae</b>			
<i>Leonotis nepetifolia</i> (L.) R. Br.,	Shrub	AM	Paris
<i>Leucas aspera</i> Link,	Herb	AM	Paris
<i>Plectranthus caninus</i> Roth	Herb	AM	Arum
<b>Malvaceae</b>			
<i>Abutilon indicum</i> (L.) Sweet	Shrub	AM	Arum
<i>Sida acuta</i> Burm. F.	Herb	AM	Intermediate
<i>Sida cordifolia</i> Forssk.,	Herb	AM	Arum
<b>Mimosaceae</b>			
<i>Mimosa pudica</i> L.	Herb	AM	Arum
<i>Prosopis cineraria</i>	Tree	AM	Arum
<i>Acacia pinnata</i>	Tree	AM	Arum
<b>Myrtaceae</b>			
<i>Syzygium cumini</i> (L.) Skeels,	Tree	AM	Arum
<b>Nyctaginaceae</b>			
<i>Boerhavia diffusa</i>	Herb	AM	Arum
<b>Oxalidaceae</b>			
<i>Biophytum intermedium</i> var. <i>pulneyensis</i>	Herb	AM	Intermediate
<i>Oxalis ausensis</i> R. Knuth	Herb	AM	Arum
<b>Papilionaceae</b>			
<i>Desmodium triflorum</i> (L.) DC.,	Herb	AM	Arum
<i>Indigofera tinctoria</i> Chapm.	Shrub	AM	Arum
<i>Acacia melanoxylon</i> R. Br.	Tree	AM	Paris
<b>Passifloraceae</b>			
<i>Passiflora leschenaultia</i> Dc.,	Shrub	AM	Arum
<b>Poaceae</b>			
<i>Bambusa bambos</i>	Herb	AM	Paris
<i>Cynodon dactylon</i> (L.) Pers.	Herb	AM	Arum
<i>Echinochloa colona</i> (L.) Link	Herb	AM	Arum
<i>Setaria verticillata</i> (L.) P. Beauv.	Herb	AM	Intermediate

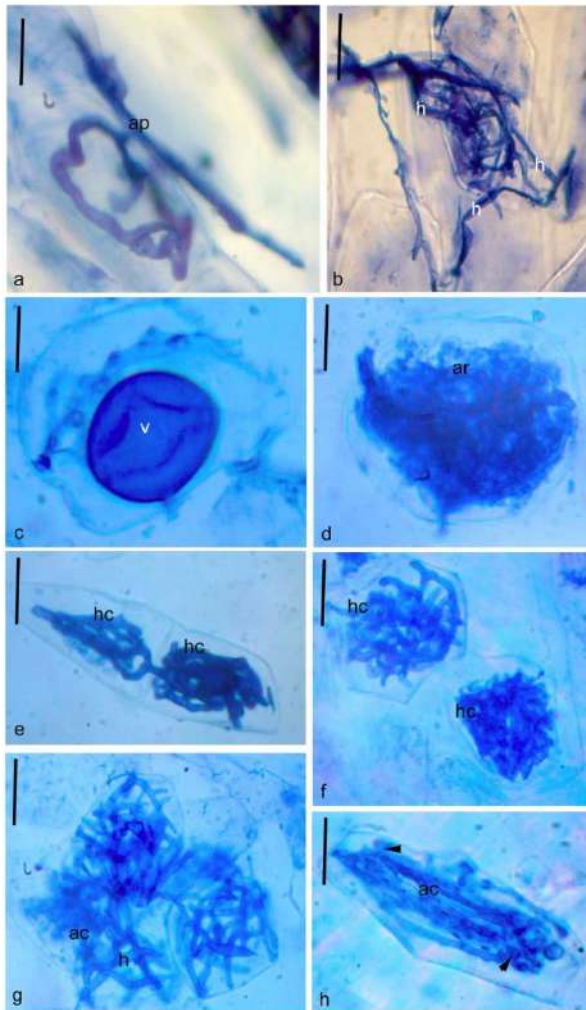
<i>Eragrostis nigra</i> Nees ex Steud	Herb	AM	Paris
<b>Polygonaceae</b>			
<i>Polygonum glabrum</i> Willd.	Herb	AM	Paris
<b>Rubiaceae</b>			
<i>Hedyotis puberula</i>	Herb	AM	Arum
<i>Lacianthus acminatus</i>	Shrub	AM	Intermediate
<i>Psychotria octosulcata</i>	Shrub	NAM	
<i>Psychotria nilgiriensis</i> var. <i>astephana</i>	Tree	AM	Intermediate
<i>Psychotria nilgiriensis</i> Deb & M. Gangop.,	Tree	NAM	
<i>Lasianthus attenuates</i> Jack	Shrub	AM	Intermediate
<i>Morinda pubescens</i> Sm.	Shrub	AM	Arum
<b>Rutaceae</b>			
<i>Toddalia asiatica</i> (L.) Lam.,	Shrub	AM	Intermediate
<b>Sapindaceae</b>			
<i>Cardiospermam helicacabum</i>	Herb	AM	Paris
<b>Solanaceae</b>			
<i>Solanum pubescens</i> Roxb.	Shrub	AM	Intermediate
<i>Solanum giganteum</i> Jacq	Shrub	AM	Arum
<b>Urticaceae</b>			
<i>Elatostema sessile</i>	Herb	AM	Intermediate
<b>Verbenaceae</b>			
<i>Clerodendrum phlomides</i>	Herb	AM	Arum
<i>Lantana camara</i> L.	Herb	AM	Intermediate
<i>Lippia javanica</i> (Burm.f.) Spreng.	Herb	AM	Arum
<b>Violaceae</b>			
<i>Hybanthus enneaspermus</i> (L.) F. Muell	Herb	AM	Paris

### Extent of AM association

There were large differences in the extent of AM colonization and root lengths with AM fungal structure between plant species. Total root length colonization (%RLTC) ranged from 25.84 % (*Commelina benghalensis*, Commelinaceae) to 95.14% (*Impatiens campanulata*, Balsamiaceae) and varied significantly among plant species ( $F_{70,213} = 44.49$ ;  $P < 0.001$ ) (Figure 3). The percentage root length with inter or intracellular hyphae (%RLH) ranged from 3.96% (*Oxalis ausensis*, Oxalidaceae) to 43.23% (*Halorrhena antidysenterica*, Apocynaceae) and varied significantly among plant species ( $F_{410, 213} = 55.38$ ;  $P < 0.001$ ). Similarly percentage root length with hyphal coils (%RLHC) ranged from 1.13 % (*Justicia tranquebariensis*, Acanthaceae) to 33.55% (*Eragrostis nigra*, Poaceae) and varied significantly among plant species ( $F_{70,213} = 68.86$ ;  $P < 0.001$ ). In colonized plants, percentage root length with arbuscules (%RLA) ranged from 1.67% (*Hybanthus enneaspermus*,

Violaceae) to 24.08% (*Curculigo orchioides*, Amaryllidaceae) and varied significantly among plant species ( $F_{70, 213} = 35.55$ ;  $P < 0.001$ ). The percentage root length with vesicles (%RLV) ranged from 0.42% (*Anaphalis lawii*, Asteraceae) to 26.81% (*Impatiens campanulata*, Balsamiaceae) and varied significantly among plant species ( $F_{70,213} = 54.15$ ;  $P < 0.001$ ). The percentage of root length with arbusculate Coils (%RLAC) ranged from 1.67% (*Echinocola colona*, Poaceae) to 22.67% (*Anaphalis lawii*, Acanthaceae) and varied significantly among plant species ( $F_{70,213} = 17.13$ ;  $P < 0.001$ ) (Table 2).

Plate - 1



**Table 2.** Extent of arbuscular Mycorrhizal (AM) fungal colonization and spore numbers in shola plant species of Kodaikanal.

Family/Plant species	% Colonization						Spore number (100g Soil)
	RLH	RLV	RLA	RLAC	RLHC	RLTC	
<b>Acanthaceae</b>							
<i>Justicia adhatoda</i>	25.15 ± 1.66	20.13 ± 1.26	15.80 ± 0.59	6.92 ± 1.26	2.52 ± 0.63	50.60 ± 4.00	3.14±0.60
<i>Justicia txanquebariensis</i>	19.77 ± 1.13	14.12 ± 0.56	12.43 ± 1.49	1.69 ± 0.98	1.13 ± 1.13	35.40 ± 4.34	
<i>Rungia repens</i>	27.83 ± 0.16	19.26 ± 0.54	12.47± 0.77	3.89 ± 1.31	2.75 ± 0.97	47.26 ± 2.18	5.33±0.56
<i>Thunbergia fragrans</i>	8.52 ± 1.19	2.92 ± 0.97	11.27 ± 0.64	6.76 ± 1.21	22.94 ± 1.12	52.41 ± 2.49	6.53±0.51
<i>Acorus calamus</i>	7.64 ± 2.67	0.00 ± 0.00	2.31±0.55	4.32 ± 0.62	25.34 ± 1.82	39.61 ± 2.84	5.88±3.13
<b>Amaranthaceae</b>							
<i>Achyranthus aspera</i>	26.54 ± 0.70	20.77 ± 0.16	18.46 ± 0.68	11.53 ± 0.09	10.38 ± 0.08	66.94 ± 0.52	3.97±1.15
<i>Aerva lanata</i>	26.94 ± 1.29	18.64 ± 1.40	12.30 ± 0.61	6.08 ± 1.30	3.84 ± 1.00	49.50 ± 3.51	5.31±0.46
<b>Amaryllidaceae</b>							
<i>Curculigo orchioides</i>	12.45 ± 1.22	0.00 ± 0.00	24.08 ± 0.84	8.64 ± 0.97	2.46 ± 0.61	47.63 ± 2.00	7.71±0.69
<b>Annonaceae</b>							
<i>Annona squamosa</i>	28.27 ± 1.84	20.25 ± 1.26	8.86 ± 0.73	7.60 ± 1.27	8.02 ± 1.12	53.12 ± 1.50	7.56±0.66
<b>Apiaceae</b>							
<i>Centella asiatica</i>	3.48 ± 0.59	4.86 ± 1.48	15.68 ± 1.94	6.92 ± 2.18	21.31 ± 1.47	52.25 ± 4.59	6.02±1.55
<b>Apocynaceae</b>							
<i>Cascable thevetia</i>	36.21 ± 4.34	19.54 ± 1.15	10.92 ± 0.58	3.45 ± 1.99	1.72 ± 0.00	52.29 ± 4.49	3.49±1.13
<i>Holarrhena antidysenterica</i>	43.23 ± 1.38	10.42 ± 2.27	8.34 ± 2.60	6.25 ± 1.80	7.81 ± 3.25	66.71 ± 4.58	3.42±0.91
<b>Aristolochiaceae</b>							
<i>Aristolochia bracteolata</i>	27.68 ±0.23	19.16 ± 0.53	12.42 ± 0.90	5.02 ± 0.73	2.72 ± 0.94	48.15 ± 2.27	9.20±0.57
<i>Aristolochia indica</i>	31.22 ± 0.25	22.77 ± 0.57	12.66 ± 0.79	9.69 ± 0.78	8.02 ± 0.48	61.76 ± 0.90	4.96±0.95
<b>Asclepiadaceae</b>							
<i>Gymnema sylvestre</i>	31.75 ± 1.19	18.31 ± 0.85	12.18 ± 0.55	10.15 ± 0.72	6.52 ± 0.50	60.76 ± 1.01	4.24±0.65
<b>Asteraceae</b>							
<i>Anaphalis lawii</i>	11.10 ± 0.74	0.42 ± 0.42	15.78 ± 1.69	22.76 ± 2.49	25.41 ± 2.14	75.47 ± 0.49	5.16 ±0.58

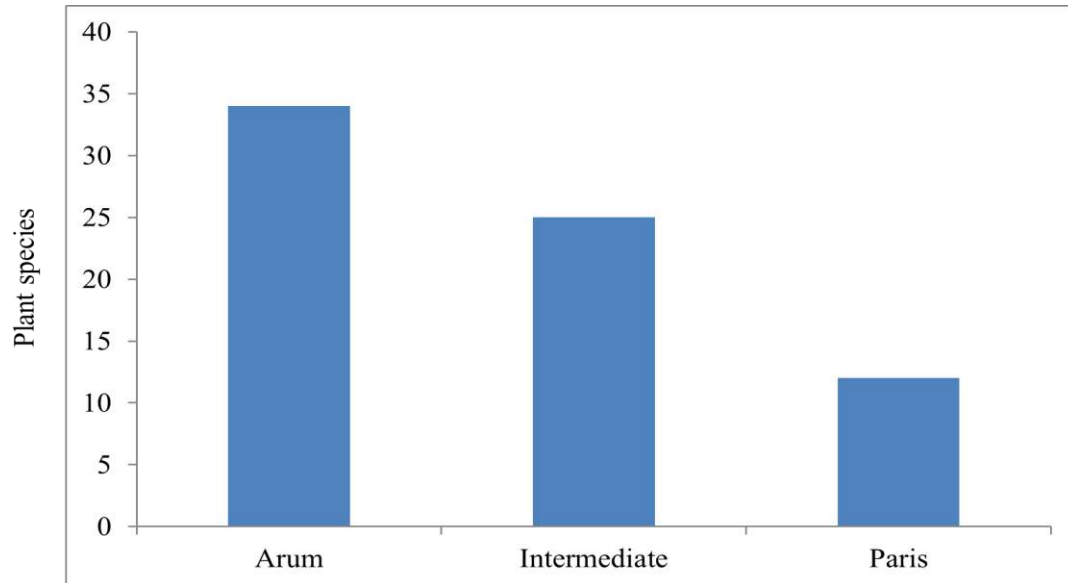
<i>Ageratum conyzoides</i>	8.64 ± 0.50	6.05 ± 0.95	21.29 ± 3.09	8.18 ± 0.31	4.83 ± 1.45	48.99 ± 2.07	7.60±0.99
<b>Balsamiaceae</b>							
<i>Impatiens campanulata</i>	38.23 ± 0.47	26.81 ± 0.93	9.27 ± 0.43	2.92 ± 0.21	17.91 ± 0.44	95.14 ± 0.39	4.35±1.10
<b>Begoniaceae</b>							
<i>Begonia malabarica</i>	8.52 ± 0.59	5.29 ± 0.55	4.36 ± 0.59	5.76 ± 1.84	29.26 ± 1.25	53.18 ± 3.65	13.38±0.62
<b>Caesalpinaceae</b>							
<i>Cassia fistula</i>	30.80 ± 0.42	22.78 ± 0.73	13.08 ± 0.42	9.70 ± 0.84	8.01 ± 0.42	61.74 ± 1.43	5.24±1.05
<i>Delonix regia</i>	25.15 ± 1.66	20.13 ± 1.26	15.80 ± 0.59	6.92 ± 1.26	2.52 ± 0.63	50.60 ± 4.00	4.44±0.72
<b>Cleomaceae</b>							
<i>Cleomy gynandra</i>	26.65 ± 1.63	21.42 ± 1.08	17.28 ± 0.15	10.84 ± 0.07	7.51 ± 1.10	62.65 ± 0.69	13.30±0.69
<b>Combretaceae</b>							
<i>Terminalia arjuna</i>	21.82 ± 0.34	9.86 ± 0.70	7.02 ± 0.57	3.50 ± 0.67	3.65 ± 0.76	36.25 ± 1.87	5.10±0.53
<b>Commelinaceae</b>							
<i>Commelina benghalensis</i>	15.34 ± 2.26	9.65 ± 0.52	6.31 ± 1.06	2.08 ± 0.10	2.08 ± 0.10	25.84 ± 2.00	6.31±0.52
<b>Convolvulaceae</b>							
<i>Evolvulus alsinoides</i>	27.95 ± 1.71	14.63 ± 1.01	10.35 ± 0.75	5.33 ± 1.09	2.75 ± 0.57	46.56 ± 1.70	3.60±0.84
<i>Ipomoea batatas</i>	5.15 ± 1.19	0.00 ± 0.00	22.38 ± 0.45	8.18 ± 0.32	3.80 ± 1.37	39.51 ± 1.30	5.04±0.91
<b>Cucurbitaceae</b>							
<i>Mukia leiosperma</i>	14.97 ± 1.58	2.89 ± 0.94	19.51 ± 0.53	6.31 ± 0.89	8.53 ± 0.55	52.21 ± 2.81	11.00±0.96
<b>Euphorbiaceae</b>							
<i>Acalypha indica</i>	26.38 ± 0.49	22.18 ± 0.04	17.62 ± 0.94	12.64 ± 0.52	10.21 ± 0.16	66.90 ± 0.40	4.53±0.35
<i>Euphorbia hirta</i>	29.88 ± 2.97	22.43 ± 0.13	12.92 ± 0.95	10.22 ± 1.70	8.07 ± 0.63	61.30 ± 0.81	9.30±0.49
<i>Jatropha gossypifolia</i>	25.18 ± 0.88	15.24 ± 0.78	11.89 ± 1.02	3.29 ± 0.63	3.32 ± 0.68	43.92 ± 0.93	3.55±0.35
<i>Phyllanthus amarus</i>	27.41 ± 0.95	20.77 ± 0.16	18.46 ± 0.68	11.48 ± 0.08	10.38 ± 0.08	67.76 ± 0.38	2.65±0.87
<i>Phyllanthus maderaspatensis</i>	28.25 ± 1.61	23.26 ± 2.81	8.85 ± 0.67	7.16 ± 1.51	8.04 ± 1.21	52.71 ± 1.67	9.80±0.65
<b>Labiatae</b>							
<i>Leonotis nepetiifolia</i>	29.33 ± 0.77	13.33 ± 0.77	12.00 ± 0.77	12.00 ± 0.77	11.11 ± 1.18	64.84 ± 1.63	5.11±0.88
<i>Leucas aspera</i>	29.21 ± 0.78	14.60 ± 0.71	11.94 ± 0.72	11.51 ± 0.91	10.61 ± 0.73	63.52 ± 0.74	5.31±0.46
<i>Plectranthus caninus</i>	4.11 ± 1.07	0.00 ± 0.00	30.89 ± 1.68	13.46 ± 0.59	4.81 ± 1.12	53.26 ± 1.85	4.23±0.60

<b>Malvaceae</b>							
<i>Abutilon indicum</i>	28.28 ± 0.51	18.18 ± 1.51	14.65 ± 0.50	5.56 ± 1.01	3.54 ± 1.33	52.47 ± 2.68	6.55±0.55
<i>Sida acuta</i>	33.41 ± 2.26	19.21 ± 0.39	12.55 ± 0.39	10.20 ± 0.39	7.06 ± 0.68	63.44 ± 1.71	5.52±0.55
<i>Sida cordifolia</i>	28.07 ± 1.68	12.17 ± 0.27	10.42 ± 0.38	3.43 ± 0.94	2.86 ± 0.46	44.93 ± 0.33	2.75±0.35
<b>Mimosaceae</b>							
<i>Mimosa pudica</i>	28.87 ± 0.73	17.61 ± 1.35	14.45 ± 1.03	6.44 ± 1.01	4.29 ± 0.59	54.26 ± 2.38	5.41±0.69
<i>Prosopis cineraria</i>	25.45 ± 1.05	12.73 ± 1.82	6.66 ± 0.61	3.03 ± 0.61	1.21 ± 0.61	36.57 ± 1.07	11.97±1.50
<i>Acacia pinnata</i>	27.10 ± 1.77	19.83 ± 1.50	12.40 ± 0.94	4.50 ± 0.50	4.52 ± 1.13	48.90 ± 3.99	5.45±0.59
<b>Myrtaceae</b>							
<i>Syzygium cumini</i>	25.44± 0.69	13.30 ± 1.43	7.61 ± 0.95	2.38 ± 0.40	1.92 ± 0.09	37.37 ± 0.99	7.22±0.58
<b>Nyctaginaceae</b>							
<i>Boerhavia diffusa</i>	24.71 ± 0.15	12.26 ± 1.26	6.49 ± 0.63	2.91 ± 0.52	1.77 ± 0.07	35.90 ± 0.09	7.79±0.70
<b>Oxalidaceae</b>							
<i>Biophytum intermedium</i>							5.35±0.45
<i>var. pulneyensis</i>	43.72± 2.04	9.21 ± 0.62	0.64 ± 0.04	5.09 ± 0.17	8.43 ± 0.29	67.09 ± 2.48	
<i>Oxalis ausensis</i>	3.96 ± 0.94	5.86 ± 1.76	34.75 ± 2.76	5.51 ± 1.01	5.44 ± 0.63	55.51 ± 5.62	4.64±0.60
<b>Papilionaceae</b>							
<i>Desmodium triflorum</i>	27.83 ± 0.16	19.20 ± 0.48	12.47 ± 0.77	3.89 ± 1.31	2.75 ± 0.97	47.26 ± 2.18	2.80±0.32
<i>Indigofera tinctoria</i>	25.29 ± 1.36	13.87 ± 2.33	8.83 ± 1.59	3.16 ± 0.62	1.90 ± 0.02	39.18 ± 3.45	11.63±0.84
<i>Acacia melanoxylon</i>	10.50 ± 1.01	3.38 ± 0.55	1.94 ± 1.36	4.65 ± 1.19	38.49 ± 2.25	58.96 ± 3.94	4.14±0.58
<b>Passifloraceae</b>							
<i>Passiflora leschenaulti</i>	22.63 ± 4.03	14.76 ± 1.39	17.60 ± 2.17	16.87± 1.20	8.81 ± 0.86	80.67 ± 3.78	7.66±0.73
<b>Poaceae</b>							
<i>Bambusa bambos</i>	28.95 ± 0.76	13.16 ± 0.76	11.84 ± 0.76	11.74 ± 0.85	10.87 ± 1.18	63.79 ± 1.99	9.52±0.43
<i>Cynodon dactylon</i>	27.26 ± 0.63	16.17 ± 1.31	10.37 ± 0.37	3.90 ± 0.10	2.57 ± 0.57	44.29 ± 1.08	12.01±0.95
<i>Echinochloa colona</i>	19.44 ± 0.56	15.55 ± 1.47	14.44 ± 2.42	1.67 ± 0.96	1.67 ± 0.96	37.54 ± 2.95	5.71±0.58
<i>Setaria verticillata</i>	32.54 ± 0.40	21.72 ± 3.28	13.09 ± 0.69	9.52 ± 0.69	7.54 ± 0.79	62.94 ± 1.95	2.41±0.42
<i>Eragrostis nigra</i>	2.50 ± 0.66	1.85 ± 0.33	2.63 ± 0.62	2.31 ± 0.32	33.55 ± 2.15	42.84 ± 3.04	7.75±0.75
<b>Polygonaceae</b>							
<i>Polygonum glabrum</i>	7.49 ± 1.08	0.00 ± 0.00	1.78 ± 0.34	4.49 ± 1.07	31.13 ± 2.19	44.89 ± 3.22	5.95±0.25

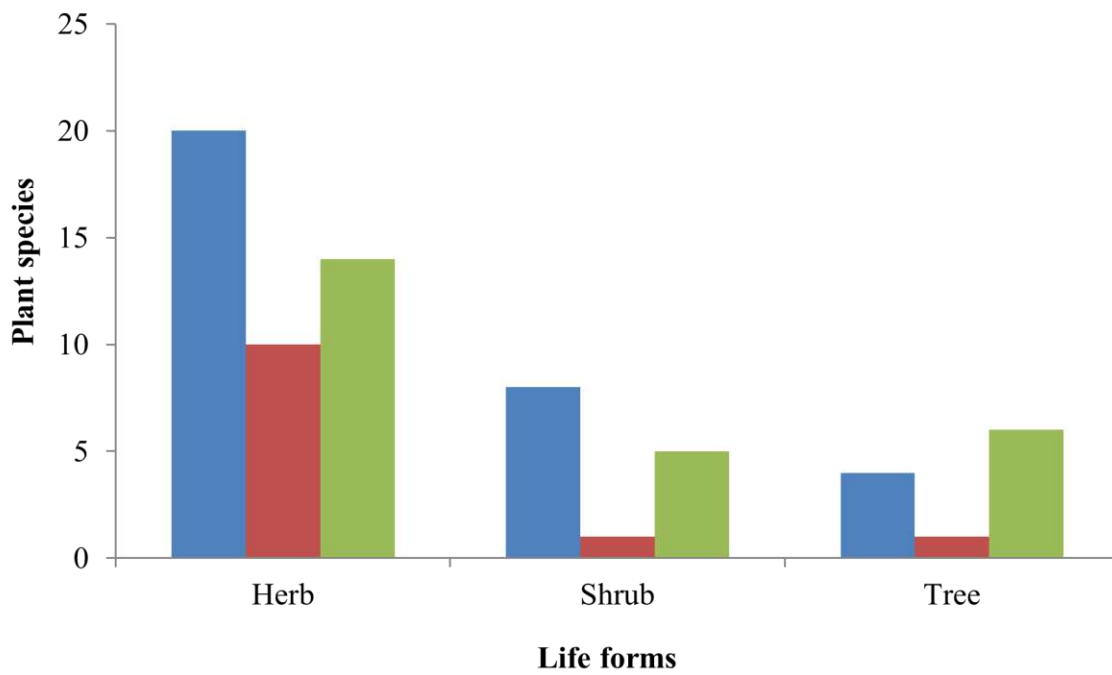
<b>Rubiaceae</b>							
<i>Hedyotis puberula</i>	27.41 ± 0.95	20.77 ± 0.16	18.46 ± 0.68	11.54 ± 0.09	10.38 ± 0.08	67.82 ± 0.35	12.57±0.91
<i>Lacianthus acminatus</i>	42.60 ± 1.10	3.12 ± 0.11	6.22± 0.25	10.32 ± 0.45	9.43 ± 0.52	71.69 ± 0.51	5.95±0.43
<i>Psychotria octosulcata</i>	13.33 ± 0.48	0.00 ± 0.00	0.00 ± 0.00	0.00 ± 0.00	0.00 ± 0.00	13.33 ± 0.48	8.88±0.93
<i>Psychotria nilgiriensis</i> <i>var. astephana</i>	23.21 ± 0.26	2.63 ± 0.20	1.92 ± 0.11	0.00 ± 0.00	4.44 ± 0.47	32.20 ± 0.69	7.32±0.64
<i>Psychotria nilgiriensis</i>	8.67 ± 0.25	0.00 ± 0.00	0.00 ± 0.00	0.00 ± 0.00	0.00 ± 0.00	8.67 ± 0.25	5.25±0.57
<i>Lasianthus attenuatus</i>	23.04 ± 1.10	2.64 ± 0.12	0.00 ± 0.00	0.65 ± 0.07	1.31 ± 0.09	27.64 ± 1.05	5.24±0.51
<i>Morinda tinctoria</i>	29.66 ± 0.43	19.55 ± 0.45	14.88 ± 1.03	5.28 ± 1.33	4.17 ± 0.99	54.33 ± 2.54	3.37±0.53
<b>Rutaceae</b>							
<i>Toddalia asiatica</i>	10.27 ± 1.33	0.00 ± 0.00	22.53 ± 4.16	17.94 ± 2.48	22.76 ± 2.30	73.51 ± 6.99	5.42±0.52
<b>Sapindaceae</b>							
<i>Cardiospermam</i> <i>helicacabum</i>	7.27 ± 0.42	3.51 ± 0.72	2.16 ± 0.55	8.31 ± 1.29	26.60 ± 1.28	47.84 ± 1.21	3.12±0.39
<b>Solanaceae</b>							
<i>Solanum pubescens</i>	30.87 ± 0.33	22.06 ± 0.73	16.20 ± 1.00	9.11 ± 0.99	6.36 ± 0.39	62.67 ± 1.48	8.67±1.44
<i>Solanum giganteum</i>	13.88 ± 1.66	8.87 ± 2.17	23.40 ± 1.52	6.42 ± 0.67	1.19 ± 0.14	53.75 ± 2.07	5.44±0.46
<b>Urticaceae</b>							
<i>Elatostema sessile</i>	26.67 ± 0.54	0.67 ± 0.09	0.00 ± 0.00	0.00 ± 0.00	6.00 ± 0.79	33.34 ± 1.03	6.22±0.94
<b>Verbenaceae</b>							
<i>Clerodendrum phlomides</i>	21.10 ± 0.59	15.16 ± 1.03	13.16 ± 0.82	2.97 ± 0.50	2.36 ± 0.46	39.75 ± 1.44	6.04±0.90
<i>Lantana camara</i>	28.27 ± 1.84	20.25 ± 1.26	8.86 ± 0.73	7.60 ± 1.27	8.02 ± 1.12	53.12 ± 2.15	10.12±1.20
<i>Lippia javanica</i>	10.71 ± 0.88	4.68 ± 1.39	13.61 ± 1.91	5.81 ± 1.90	1.76 ± 0.23	36.57 ± 1.54	8.31±0.57
<b>Violaceae</b>							
<i>Hybanthus enneaspermus</i>	6.20 ± 0.88	3.41 ± 0.60	1.67 ± 0.39	2.98 ± 0.99	29.10 ± 1.12	43.36 ± 0.71	5.21±0.54



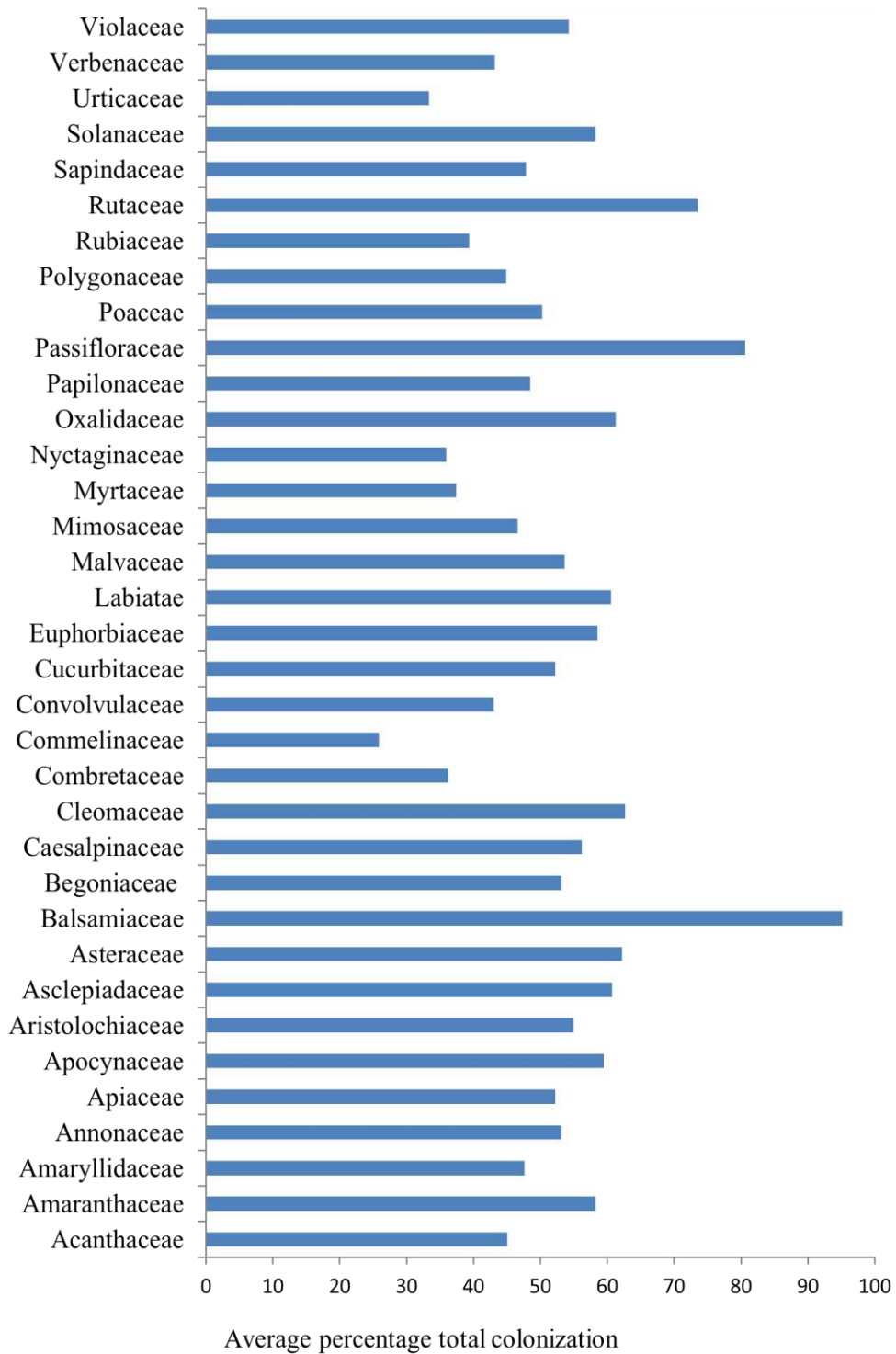
**Figure 1. Arbuscular mycorrhizal fungal morphology in shoal plant species of Kodaikanal**



**Figure 2. Arbuscular mycorrhizal fungal morphology in various life forms of shoal species in Kodaikanal**



**Figure 3. Average arbuscular mycorrhizal fungal colonization in plant species of shoal forests in Kodaikanal**



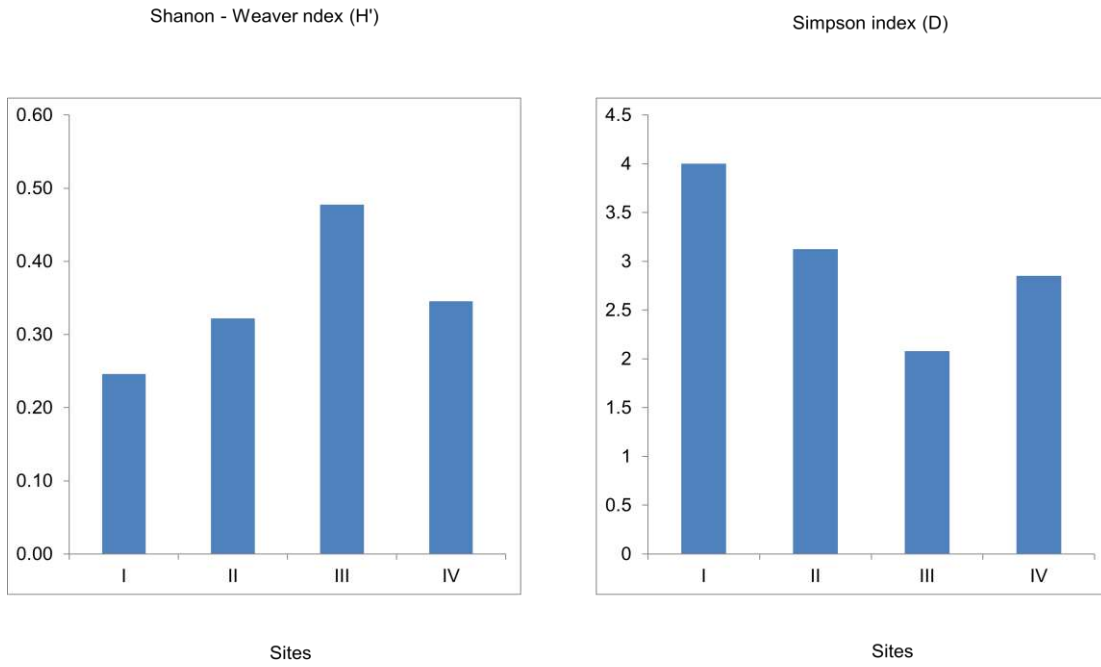
### AM fungal characteristics of the study sites

A total of Seven AM fungal morphotypes could be distinguished on the basis of spore morphology, to the species level (Plate 2; Table 3). These included one species in *Acaulospora scrobiculata* Trappe), *Scutellospora calospora* Walker and Sanders, *Funneliformis geosporum* (Nicol. and Gerd.) Walker, *Glomus aggregatum* Schenck and Sm. emend. Koske, *Glomus sinuosum* (Gerd. and Bakshi) Almeida and Schenk, *Glomus viscosum* Nicolson, and *Funneliformis mosseae* (Nicolson&Gerd.) C. Walker &A.Schubler. Distribution of AM fungal spores (Table. 5) ranged from 2 spores 100 g<sup>-1</sup>soil (*Setaria verticillata*, Poaceae) to 13 spores 100 g<sup>-1</sup> soil (*Cleome gynandra*, Cleomaceae)(Table 3). The spore numbers were not related to the extent of AM colonization ( $r = -0.10$ ;  $p >0.05$ ;  $n = 213$ ). The diversity indices like Shannon - Weaver index ( $H'$ ) ranged from 0.25(Site I) to 0.48 (Site III) and the Simpson index ( $D$ ) ranged from 2.08 (Site III) to 4 (Site I) (Figure 4). *Glomus aggregatum* was the most frequent species in shoal forest. In Seasonal pattern of AM fungal spore numbers shows that, 24 spore in the month of September and six spores in the month of July (Figure 5).

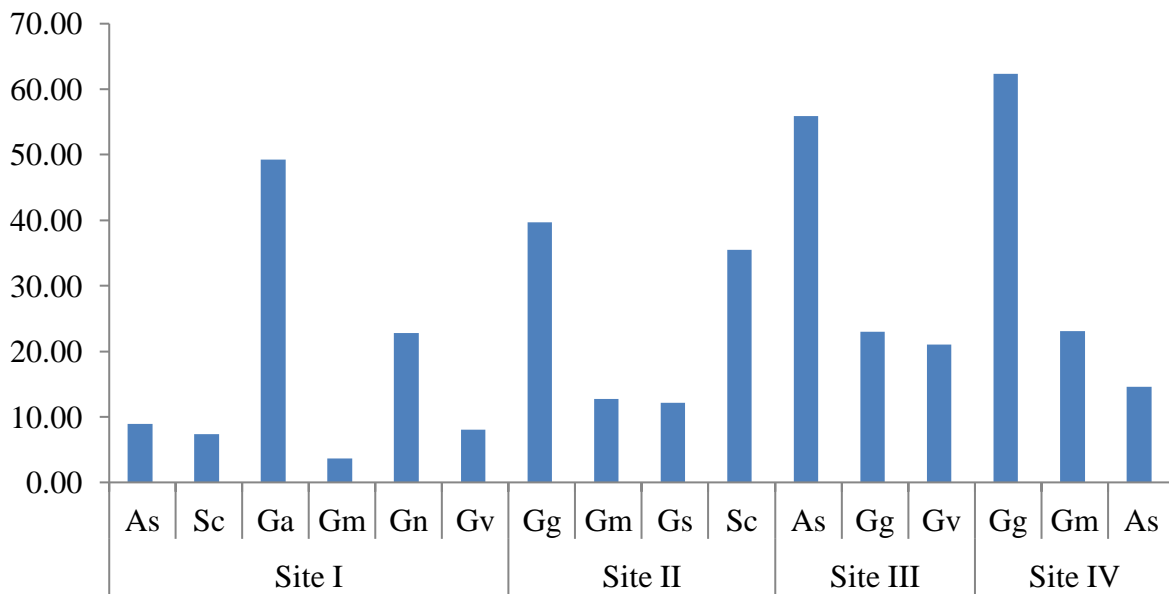
**Table 3.** Arbuscular mycorrhizal fungal spore morphotypes isolated from different sites in sholas at Kodaikanal. (X indicates the presence).

<b>Fungal species</b>	<b>Site I</b>	<b>Site II</b>	<b>Site III</b>	<b>Site IV</b>
<i>Acaulospora scrobiculata</i> Trappe	X		X	X
<i>Scutellospora calospora</i> Scutellospora calospora (T.H. Nicolson & Gerd) C. Walker & F.E. Sanders	X	X		
<i>Glomus aggregatum</i> N.C. Schenck & G.S. Sm	X			
<i>Funneliformis mosseae</i> (T.H. Nicolson & Gerd.) C. Walker & A. Schüßler	X	X		X
<i>Glomus sinuosum</i> T.H. Nicolson	X	X		
<i>Glomus viscosum</i>	X		X	
<i>Funneliformis geosporum</i> (T.H. Nicolson & Gerd.) C. Walker & A. Schüßler		X	X	X

**Figure 4: Arbuscular mycorrhizal diversity indices of shoal forests in Kodaikanal**



**Figure 5: Frequency of arbuscular mycorrhizal fungal species in the shoal forests of Kodaikanal**



**Figure 6: Arbuscular mycorrhizal fungal species dynamics in various seasons of Kodaikanal**

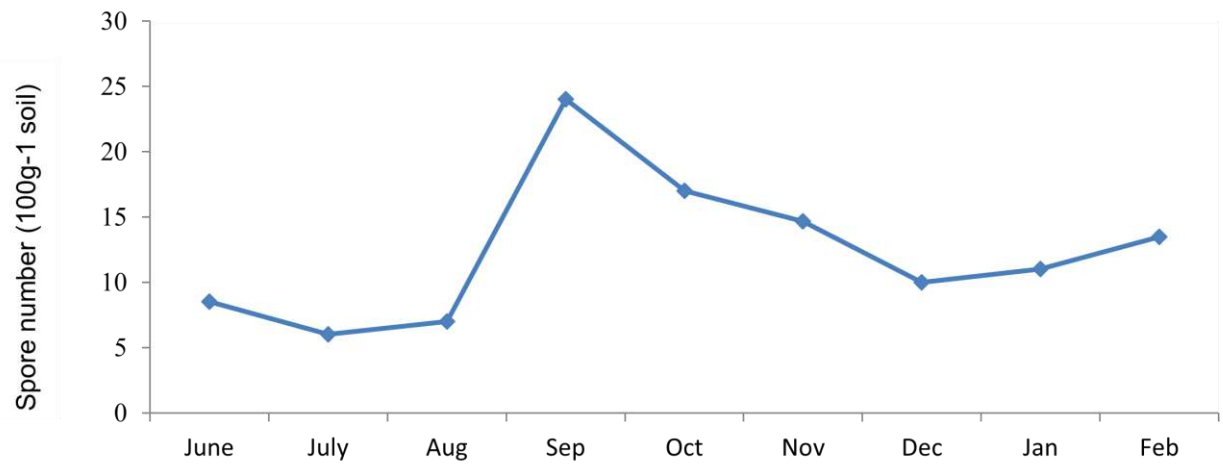
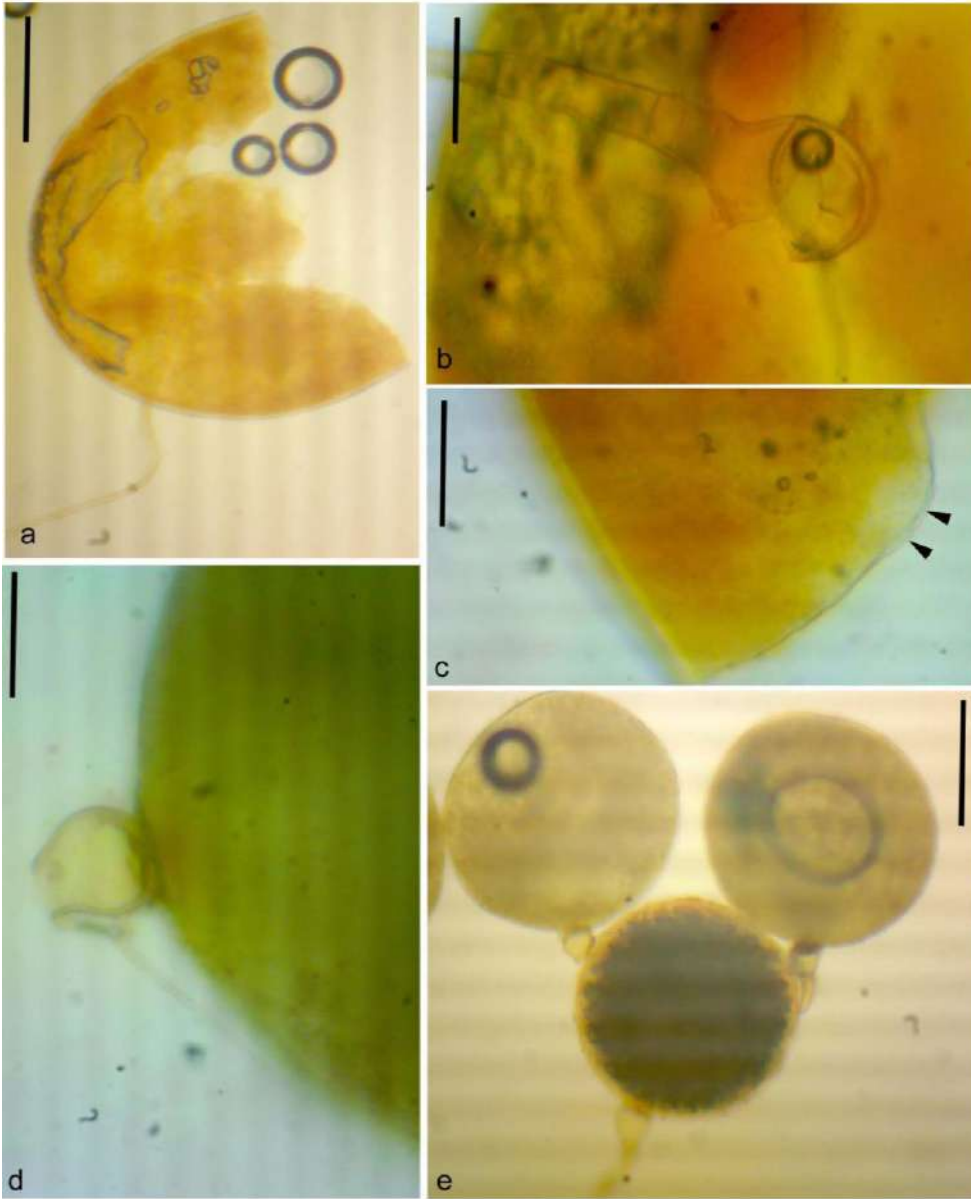


Plate - 2



## Discussion

AM fungi colonize the roots of most land plants, where they facilitate mineral nutrient uptake from the soil in exchange for plant assimilated carbon. Though about 80% of the land plants are assumed to form AM association only slightly over 10,000 plant species i.e. around 3% of the known plant species have been examined for AM association (Wang and Qui, 2006). In recent years more information has been gathered regarding the mycorrhizal status of plants in natural ecosystems (Muthukumar *et al.*, 2006; Radhika and Rodrigues, 2007; Gai *et al.*, 2006; Grippa *et al.*, 2007; Tawaraya *et al.*, 2003; Tanumi Fuman and Monoranjana Ghose, 2008). In particular, there is evidence that AM fungi are common in plant groups once considered as non-mycorrhizal (Muthukumar *et al.*, 2004; Radhika and Rodrigues, 2007). This study was intended to generate more information on the occurrence of AM fungal association in Shola plant species.

The incidence of mycorrhiza (97%) in plant species of Shola forest was higher than those reported for angiosperms by Trappe (1987) and Wang and Qui (2006). Seventy percent of the 6,500 angiosperms indexed by Trappe (1987) and 80% of the 2,964 angiosperms listed by Wang and Qui (2006) were mycorrhizal. The higher mycorrhizal incidence of angiosperms in the present study could be attributed to the low nutrient status of the soils along with high plant competition. Non-mycotrophy was low (3%) in this study compared with reports from other vegetation types worldwide (Ragupathy and Mahadevan, 1993; Muthukumar and Udaiyan, 2000; Zhao *et al.*, 2001; Muthukumar *et al.*, 2003). The phenomenon of non-mycotrophy is often associated with high levels of disturbance, or under extreme environmental conditions, the low incidence of non-mycotrophy in the present study is not surprising. However, plants that lacked mycorrhizae belonged to Commelinaceae, Cleomaceae and Convolvulaceae are reported as mycorrhizal plant families (Wang and Qui, 2006).

Surveys of earlier literature showed that *Paris* type colonization occurs more predominantly in wild angiosperms (Smith and Smith, 1996; 1997; Menoyo *et al.*, 2007). In the present study, 35% (25/71) of mycorrhizal plant species had Intermediate-type of AM morphology and typical *Paris*-type morphology occurred only in 17% (12/71) of the mycorrhizal species and 48% (34/71) of the shoal species had *Arum*- type AM morphology. It has been shown that host plants control the morphological types of AM. Gerdemann (1965) demonstrated that the same species of AM fungi formed the *Paris*-type in *Liriodendron* and *Arum*- type in maize, respectively. Likewise, Jacquelinet – Jeanmougin and Gianinazzi – Pearson (1983) showed that the *Paris*- type in *Gentiana* was formed by the same fungus which formed the *Arum*- type in *Allium*. Brundrett and Kendrick (1990) suggested that the types of AM are determined by the presence of continuous longitudinal air-spaces in the root cortex, i.e. the *Arum*- type is formed in their presence and the *Paris*-type is formed in their absence. However, even though the fungal identity could determine the morphological types of AM in some cases, it still seems likely that only a single type is found in a plant in most cases, which indicates the morphological types of AM depend on the characteristics of plants rather than those of Fungi (Yamato and Iwasaki, 2002).

The spore numbers of 2 to 13 spores per 100g soil is low compared to 14 to 93 per 100g soil reported by Muthukumar *et al* (2003) and 55 to 191 spores per 100g soil reported by Zhao *et al.* (2001) from Primary forest of Xishuangbanna, southwest China. The low density of AM fungal spores of the present study corroborates the reports from humid tropical forest where spore numbers tend to be low or infrequent (Janos, 1980; Fischer *et al.*, 1994). Generally, AM fungal spores in natural soils are dead or parasitized and are merely spore cases (Muthukumar and Udaiyan, 1999). The spore number reported in this study is intact healthy spores. In addition, a range of environmental, host and fungal factors influences AM fungal sporulation and spore numbers tend to decrease during root growth, but tend to increase during root inactivity or senescence (Brundrett, 1991). In undisturbed forest, spores may be relatively less important than other vegetative propagules, and primarily the soil hyphal networks initiate the colonization of new roots (Jasper *et al.*, 1989). As a result, forests with root growth throughout the year usually have small spore populations and high mycorrhizal colonization levels (Muthukumar *et al.*, 2003; Zandavalli *et al.*, 2008). In this study, AM fungal spores were present in the rhizosphere of two shoal plants species lacking



AM colonization. In natural soils, roots of adjacent plants often grow in close proximity and are interwoven, so spores in the rhizosphere of a host could be contributed by a companion plant species (Muthukumar and Udaiyan, 2000).

A total of seven AM fungal species were identified based on the morphological characters of the spores. This number is about half of those reported in semiarid Mediterranean ecosystems (Ferrol *et al.*, 2004) and semi- arid areas in Brazil (Silva *et al.*, 2005) where 23 and 21 AM fungal species were reported. However, a high AM fungal diversity has also been reported in other natural ecosystems. Muthukumar and Udaiyan (2000) reported 6–22 species per site from Western Ghats region of South India. Forty four AM fungal species were isolated from grasslands of Namibia (Uhlmann *et al.*, 2004), 43 from an arid steppe of inner Mongolia (Tian *et al.*, 2008) and 27 species from tropical rain forest of Xishuangnanna, southwest China (Zhao *et al.*, 2003). In this study, the rhizosphere soil samples of plant species contained spore morphotypes of 1 to 4 AM fungal species. This shows that several fungal species can colonize the roots of an individual plant in a natural ecosystem (Van Tuinen *et al.*, 1998; Helgason *et al.*, 1999), thus showing lack of host specificity (Smith and Read 1997). AM fungal spores belonging to *Glomus* predominated species diversity, which is in accordance with the observations that species of *Glomus* dominate tropical soils (Muthukumar and Udaiyan, 2000; Zaho *et al.*, 2001; 2003).

The lack of correlation between the AM fungal spore numbers and percent root length colonization is consistent with several previous reports in which the lack of demonstrable relationship was reported between these mycorrhizal variables (Brundrett, 1991; Zhaka *et al.*, 1995; Brundrett *et al.*, 1996). As a wide range of plant, fungal and environmental factors influence AM fungal colonization and sporulation the observed lack of relationship between these mycorrhizal variable is tenable (Helgason and Fitter, 2005).

In conclusion, AM fungal association was found to be wide spread in the plant species of shoal forests of Kodaikanal Hills, Western Ghats region. AM fungi can enhance root functions of native plants in natural ecosystems, where they are exposed to extreme competition. The future phase of this study is to entail experimental studies of these rare and economically important plant species to determine the effects of fungal inoculants on growth to restoration of shoal species in forestry.

## Summary

In natural ecosystems plant roots are colonized various microorganisms which affects plant distribution, survival and growth in different mechanisms. Most prevalent microorganism in most of the ecosystems is AM fungi. However, the prevalence of this association has been well reported from several natural ecosystems, information on AM fungal association and their abundance are unknown for shola ecosystems. In the present study, 71 plant species (in 35 families) examined, all the families were colonized by AM fungi except two species in a genus *Psychotria*. AM association was observed in members of supposedly non-mycorrhizal families Commelinaceae, Cleomaceae and Convolvulaceae. Only those species in which arbuscules or arbusculate coils were found were considered to have AM association. The fungal entry into roots was characterized by the presence of appressorium. Thirty four of the plant species had Arum-type morphology, 25 had Intermediate- type and 12 had typical Paris-type morphology. In herbs 20 species had Arum type morphology, 10 had Paris type morphology and 14 had Intermediate type morphology. In Shrubs, 8 species had Arum, one species had Paris and 5 species had Intermediate type morphology. In Tree species 4, 1 and 6 plant species had Arum, Paris and Intermediate type morphology respectively. There were large differences in the extent of AM colonization and root lengths with AM fungal structure between plant species. Total root length colonization (%RLTC) ranged from 25.84 % (*Commelina benghalensis*, Commelinaceae) to 95.14% (*Impatiens campanulata*, Balsamiaceae). The percentage root length with inter or intracellular hyphae (%RLH) ranged from 3.96% (*Oxalis ausensis*, Oxalidaceae) to 43.23% (*Halorrhena antidysenterica*, Apocynaceae). Similarly percentage root length with hyphal coils (%RLHC) ranged from 1.13 % (*Justicia tranquebariensis*, Acanthaceae) to 33.55% (*Eragrostis nigra*, Poaceae). In colonized plants, percentage root length with arbuscules (%RLA) ranged from 1.67% (*Hybanthus enneaspermus*, Violaceae) to 24.08% (*Curculigo orchioides*, Amaryllidaceae). The percentage root length with vesicles (%RLV) ranged from 0.42% (*Anaphalis lawii*, Asteraceae) to 26.81% (*Impatiens campanulata*, Balsamiaceae). The percentage of root length with arbusculate Coils (%RLAC) ranged from 1.67% (*Echinocola colona*, Poaceae) to 22.67% (*Anaphalis lawii*, Acanthaceae). A total of Seven AM fungal morphotypes could be distinguished; these included species in *Acaulospora*, *Scrobiculata*, *Scutellospora calospora* Walker and Sanders, *Funneliformis geosporum* (Nicol. and Gerd.) Walker, *Glonus*

*aggregatum* Schenck and Sm. emend. Koske, *Glomus sinuosum* (Gerd. and Bakshi) Almeida and Schenk, *Glomus viscosum* Nicolson, and *Funneliformis mosseae* (Nicolson&Gerd.) C. Walker &A.Schubler. Distribution of AM fungal spores ranged from 2 (*Setaria verticillata*, Poaceae) to 13 spores 100 g<sup>-1</sup> soil (*Cleome gynandra*, Cleomaceae). The spore numbers were not related to the extent of AM colonization. *Glomus aggregatum* was the most frequent species in shoal forest.

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