

OCCURRENCE OF VESICULAR ARBUSCULAR MYCORRHIZAL FUNGI IN TROPICAL FOREST COMMUNITIES OF INDIA

TRIPATHI, P.* – KHARE, P.K.

*Department of Botany, Dr. H.S. Gour Central University,
Sagar (M.P.) 470003, India*

**Corresponding author
e-mail: priyanganaco@gmail.com*

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Abstract. The paper deals with the quantitative and qualitative assessment of vesicular arbuscular mycorrhizal fungi (AMF) associated with the rhizosphere of dominant plants of three vegetation viz., Tropical Evergreen Forest, Gibbon Wild-life sanctuary, Jorhat, Assam, Tropical Moist Forest, Kanha National Park, Mandla, M.P., and Tropical Dry Deciduous Forest, Sagar, M.P. Root and rhizosphere soil samples, collected near different trees species were analyzed for root colonization, spore count, extramatrical hyphae, vesicles, Paris coil and intraradical spores. VAM spores were isolated by wet sieving and decanting method and estimation of spores was carried out. In evergreen forest vegetation (Site-I), maximum root colonization (33 ± 0.35 %) was observed in *Terminalia myrocarpa* Henrck & Mull. and minimum in *Walsura robusta* Roxb. (19 ± 0.61 %). VAM spore count per 25g of soil was also highest (133 ± 0.41) in *Terminalia myrocarpa* Henrck & Mull. while it was low in *Mesua ferrea* Linn. (48 ± 0.21). In tropical moist forest (Site-II), *Bridelia retusa* Spreng. was observed to support least root colonization (33 ± 0.4 per cent) and spore density ($43 \pm 0.25/25$ g soil). Maximum root colonization (64 ± 0.49 %) and spores density (167 ± 0.46) were found in *Lagerstroemia parviflora* Roxb. In tropical dry deciduous forest (Site-III), root colonization (62 ± 0.62 %) and spores density (121 ± 0.31) were maximum in *Albizia lebbek* Benth.-highest among all the three sites. Interestingly, these attributes were also lowest at this site in *Wrightia tinctoria* Linn. In all, 25 VAM species were identified from all the three forest sites. Out of these, 5 VAM species were common to three forest sites. More number of exclusive VAM species (6) were recorded at site-II. *Glomus* spp. was found as dominant VAM at all the three forest sites. Besides, the other morphological details of different VAM fungi, the results of present investigation envisage that diverse conditions of the habitat i.e. Tropical Moist Forest, support more VAM species not only in terms of numbers but also as the root association and potential propagules in the form of spores.

Keywords: VAM, Root colonization, Qualitative and Quantitative characters

Introduction

The Vesicular Arbuscular Mycorrhizal (VAM) association with most of the terrestrial plants is perhaps as old as the evolution of terrestrial flora (Taylor et al., 1995). They are ubiquitous in distribution from tropical (Chaurasia and Khare, 2005), temperate (Vestberg, 1995) and arctic (Dalpe and Aiken, 1998) regions. They are responsible for enhancing the growth and improving the health of plants and soils (Hodge et al., 2001; Rausch et al., 2001) and play a crucial role in mineral nutrition of forest trees thus becoming important nutrient acquiring mechanisms (Pate, 1994).

Although the occurrence of VAM species is non specific, they exhibit variation in both qualitative and quantitative attributes. Variation may be found due to climatic, edaphic and composition of vegetation types.

The present paper deals with the occurrence and population variations of arbuscular mycorrhizal fungi in three different tropical forests of India namely tropical evergreen forest, tropical moist deciduous forest and tropical dry deciduous forest. Root

colonization, spore population and qualitative characteristics like external hyphae, vesicle, Paris coil, intraradical spores were determined and discussed.

Study area

Site-I The site belongs to a tropical evergreen forest within Gibbon National Park near Jorhat, Assam, about 30 km West of Jorhat, situated at 26°46'N latitude and 96°16'E longitude. Climate is a seasonal with average monthly minimum and maximum temperature of 19°C and 30°C in January and May respectively. Annual rainfall is more than 300 cm received throughout the year. The soil is clay to sandy clay with pH ranging from 4.3 to 4.9. The forest types as classified by (Champion and Seth, 1968) is semievergreen to evergreen. Dominant forest tree species are *Dipterocarpus myrocarpa* Vesque, *Walsura robusta* Roxb., *Dillenia indica* Linn.

Site-II The site, tropical moist forest, is a part of Kanha National Park, a protected area dedicated to Project Tiger in the state of Madhya Pradesh in Central India. It is located in between 22°1' and 22°27'N, and 80°26' and 81°4'E longitude at an altitude of 800-900 msl. The climate is distinctly seasonal with three well marked seasons viz., rainy, winter and summer. Maximum temperature of 40°C reaches in summer and minimum as 0.0°C during the month of December in winter season. The total annual rainfall is 140 cm principally received during the months of July to September. Edaphically it constitutes the Central Indian highlands that forms main peninsula of India. The terrain is undulating with clayey laterite rich in Bauxite on tops and plains, and valley with granite gneisses. The soil is near neutral or alkaline type. Forests are tropical moist deciduous types dominated chiefly by *Shorea robusta* Gaertner F.

Site-III This site is a part of tropical dry deciduous forests in Central India, located in between 23°5' – 24°25' N and 78°10' – 79°15' E longitude at an elevation of 583 msl near the town of Sagar in the state of Madhya Pradesh. The climate is distinctly seasonal with summer, rainy and winter seasons. Total annual rainfall is 120 cm, nearly 90% of which is received in rainy season. Summer is hot with maximum temperature touching 47°C and winter is pleasant with minimum temperature of 8-9°C. A long dry season of about 7-8 months is responsible for the occurrence of tropical dry deciduous forests principally dominated by *Tectona grandis* with associates like *Terminalia tomentosa* W. & A., *Diospyros melanoxylon* Roxb. and *Butea monosperma* Lamk. Taub. Soil is thin, rich in calcium with pH ranging from 6.5 to 8.5.

Materials and methods

Sample collection

Soil and root samples were collected during the month June (Late Summer) from selected dominant and codominant forest tree species of all the three sites. Individual non-adhering rhizosphere soil samples were taken in triplicate from the depth of 10-15 cm, after removing upper detritus. Fine rootlets (feeder roots) were collected from the dominant species of each site. Samples were air dried, sieved and stored at 4°C until further processing.

Processing of root samples

Root samples were rinsed with tap water and cut into 1 cm pieces, and the staining was done with trypan blue (Phillips and Haymann, 1970; Kormainik et al., 1980). One hundred root pieces were randomly picked and mounted on glass slides in lactophenol and examined under compound microscope. The colonization was calculated using the following formula

$$\% \text{ colonization} = \frac{\text{no. of root segments colonized}}{\text{Total no. of root segments studied}} \times 100 \quad (\text{Eq.1})$$

Qualitative characteristics of mycorrhizal fungi i.e. external hyphae, presence of vesicles, Paris coil and occurrence of intraradical spores, were observed in stained root samples.

Extraction of spores and their identification

Spores from the soil samples (25g) were extracted by wet sieving and decanting method (Gerdemann and Nicolson, 1963). For the estimation of VAM species, a modified method (Gour and Adholeya, 1994) was followed and identification was done on current descriptions and identification manuals (Schenk and Perez, 1990; Mehrotra and Baizal, 1994; www.invam.caf.wvu.edu/my-info/Taxonomy/species.htm). Species name code of VAM fungi were followed (Perez and Schenck, 1990).

Results

Data of rhizospheric soil analysis of three study sites is presented in the table 1 to 4. Twelve tree species were analyzed for quantitative and qualitative characters of VAM fungi from site I. The lowest root colonization was observed in *Walsura robusta* Roxburgh (19%) and highest in *Terminalia myrocarpa* Henrck & Mull. (33%) followed by *Terminalia nodiflora* (R. Ar.) (32%) and *Dillenia indica* Linn. (32%). VAM fungal spore population per 25 g of rhizospheric soil of 12 tree species from site I is presented in the *Table 1*. It is evident that the lowest spore population (48/25 gm) was observed in rhizosphere soil of *Mesua ferrea* Linn. and highest spore population (133/25 g) in *Terminalia myrocarpa* Henrck & Mull. Qualitative characters of VAM fungi viz., external hyphae, vesicle, Paris coil, intraradical spores are given in *Tables 1 to 3*. The presence of extrametrical hyphae were recorded in *Terminalia myrocarpa* Henrck & Mull., *Chukrasia tabularis* (A. Juss.), *Tetrameles nodiflora* (R. Ar.), *Meusa ferrea* Linn., *Artocarpus lakoocha* Roxb. and *Walsura robusta* Roxb.. Vesicles were less common although observed in *Dillenia indica* Linn., *Mesua ferrea* Linn., *Artocarpus lakoocha* Roxb. and *Walsura robusta* Roxb. Paris type coils were observed in root pieces of *Dipterocarpus macrocarpus* Vesque, *Castanopsis spp* Blume., *Chakrasia tabularis* (A. Juss.), *Terminalia myrocarpa* Henrck & Mull. and *Artocarpus chama* Buch.Ham. Intraradical spores were abundant with the exception of *Dipterocarpus macrocarpus* Vesque, *Dillenia indica* Linn., *Artocarpus lakoocha* Roxb. and *Artocarpus chama* Buch.Ham. Total 15 VAM species were recovered from this site, out of which 5 VAM species were exclusive, i.e. restricted to this site only.

Table 1. Quantitative and Qualitative characters of VAM fungi associated with some important forest tree species at Site-I (Jorhat)

S. No.	Plant	Quantitative Characters			Qualitative Characters		Vesicle	Paciscoil	Intraradical	VAM Species
		% Root colonization	No. of spores in 25g soil	Exhamatrica Hyphae						
1	<i>Dipterocarpus macrocarpus Vesque</i>	29 SD±0.30 SE±0.15	85 SD±0.45 SE±0.26	-	-	-	+	-	ARHM, LFSC, SPKS, LABS, LMSS	
2	<i>Castanopsis spp. Blume</i>	23 SD±0.45 SE±0.23	66 SD±0.25 SE±0.14	-	-	-	+	+	AUDL, LFSC, LAGR, LABS, LMSS	
3	<i>Dillenia indica Linn.</i>	32 SD±0.38 SE±0.19	92 SD±0.42 SE±0.24	-	-	++	-	-	ASP, LFSC, LAGR, LABS, LHTS	
4	<i>Lagerostromia speritosa (L.)</i>	22 SD±0.31 SE±0.15	105 SD±0.51	-	-	-	-	+	CPLC, LFSC, LAGR, LOCT, LMSS, LHTS	
5	<i>Tetrameles nodiflora (R. Ar.)</i>	32 SD±0.64 SE±0.32	94 SD±0.45 SE±0.26	+	+	-	-	+	ASP, ARHM, LPKS, LAGR, LHTS	
6	<i>Chukrasia tabularis (A. Juss.)</i>	20 SD±0.42 SE±0.21	102 SD±0.83 SE±0.48	+	+	-	+	+	AUDL, LFSC, LAGR, LMSS, LABS	
7	<i>Amoora wallichii Benth.</i>	28 SD±0.56 SE±0.28	99 SD±0.58 SE±0.33	-	-	-	-	+	LMSS, GRSA, LPKS, SSNS, LMSS	
8	<i>Terminalia myrocarpa Henck & Mull.</i>	33 SD±0.35 SE±0.18	133 SD±0.41 SE±0.21	+	+	-	+++	+	ASP, LFSC, SPKS, LOCT, LMSS, LHTS	
9	<i>Mesua ferrea Linnaeus</i>	27 SD±0.95 SE±0.48	48 SD±1.21 SE±0.70	+	+	+	-	+	GRSA, ARHM, LAGR, LABS	
10	<i>Artocarpus lakoocha Roxb.</i>	27 SD±0.78 SE±0.39	79 SD±0.39 SE±0.22	+	+	+	-	-	GRSA, LFSC, LAGR, LOCT, LHTS	
11	<i>Artocarpus chama Buch. Ham</i>	30 SD±0.59 SE±0.31	122 SD±0.47 SE±0.27	-	-	-	+	-	ATBC, LFSC, SPKS, LOCT, LHTS	
12	<i>Walsura robusta Roxburgh</i>	19 SD±0.61 SE±0.31	57 SD±0.53 SE±0.30	+	+	+	-	+	ATBC, ARHM, LAGR	

Table 2. Quantitative and Qualitative characters of VAM fungi associated with some important forest tree species at Site-II (Mandla)

S. No.	Plant	Quantitative Characters			Qualitative Characters		Vesicle	Paciscoil	Intraradical	VAM Species
		% Root colonization	No. of spores in 25g soil	Exhamatrica Hyphae	Exhamatrica Hyphae					
1	<i>Bridelia retusa</i> Spreng	33 SD±0.47 SE±0.24	43 SD±0.25 SE±0.15	+	+	-	+	++	LGSP, LETC	
2	<i>Syzygium cumini</i> (L.) skeels	48 SD±1.28 SE±0.64	96 SD±0.25 SE±0.14	+	+	+	+	+	LAGR, LFSC, LCVL, LDST	
3	<i>Diospyros melanoxylon</i> (Roxb.)	43 SD±0.45 SE±0.23	84 SD±0.36 SE±0.41	+	+	+	-	+	LMSS, LFSC, LDST	
4	<i>Cassia fistula</i> Linn.	41 SD±0.58 SE±0.29	103 SD±0.08 SE±0.	+	+	-	+	-	LAST, LETC, LDST, LHTS	
5	<i>Careya arborea</i> (Roxb.)	52 SD±0.69 SE±0.35	91 SD±0.94 SE±0.55	+	+	+	+	-	LABS, LFSC, LCVL	
6	<i>Terminalia tomentosa</i> W. & A.	55 SD±0.39 SE±0.21	127 SD±0.08 SE±0.05	+	+	+	+	-	CFLG, LMSS, LETC, LHTS	
7	<i>Shorea robusta</i> Gaertner F.	60 SD±0.31 SE±0.15	157 SD±0.52 SE±0.31	-	+	+	-	+	LGSP, LFSC, LDST, LHTS	
8	<i>Schleichera oleosa</i> (Lour) Oken	46 SD±0.42 SE±0.21	83 SD±0.43 SE±0.25	+	+	-	+	+	LHTS, LETC, LDST	
9	<i>Terminalia chebula</i> Retz.	57 SD±0.77 SE±0.38	133 SD±0.42 SE±0.25	-	+	+	+	-	LGSP, LFSC, LETC	
10	<i>Adina cordifolia</i> Hook.F.	39 SD±0.50 SE±0.25	139 SD±0.08 SE±0.05	-	+	-	+	+	LHTS, LFSC, LCVL	
11	<i>Bombox ceiba</i> Auct.	40 SD±0.50 SE±0.25	123 SD±0.53 SE±0.30	-	+	+	+	-	LGSP, LHTS, LETC, LFLV	
12	<i>Lagerstroemia parviflora</i> Roxb.	64 SD±0.49 SE±0.24	167 SD±0.45 SE±0.26	+++	+	+	+	-	ATBC, LAST, CPLC, LTNB, LMCC	
13	<i>Embellica officinalis</i> Gaerth.	53 SD±0.19 SE±0.11	91 SD±0.46 SE±0.26	+	+	+	+	+	LGSP, LFSC, LDST, LHTS	
14	<i>Mitragyna parviflora</i> Korth.	54 SD±0.66 SE±0.33	100 SD±0.65 SE±0.37	-	+	+	-	-	LGSP, LHTS, LDST	

Table 3. Quantitative and Qualitative characters of VAM fungi associated with some important forest tree species at Site-III (Sagar)

S. No.	Plant	Quantitative Characters			Qualitative Characters Exhamatrica Hyphae	Vesicle	Paciscoil	Intraradical	VAM Species
		% Root colonization	No. of spores in 25g soil						
1	<i>Holoptelia integrefolia</i> Planch	16 SD±0.89 SE±0.44	47 SD±0.43 SE±0.25		+	+	+	LAGR, LFSC, ASCB, LMSS	
2	<i>Flacourtia indica</i> (Burm F) Merr.	29 SD±1.06 SE±0.27	64 SD±0.90 SE±0.52		-	+	+	CFLG, ASCB, LAGR, LMSS	
3	<i>Wrightia tinctoria</i> Linn.	14 SD±0.85 SE±0.42	29 SD±0.67 SE±0.39		-	+	-	LFSC, ASCB, LAGR, LMSS	
4	<i>Acacia catechu</i> Willd.	21 SD±0.60 SE±0.30	33 SD±1.20 SE±0.71		-	+	-	SPKS, SSNS, LFSC, LAGR	
5	<i>Butea monosperma</i> (Lamk. Taub)	38 SD±0.49 SE±0.25	99 SD±0.29 SE±0.17		-	++	+	LAGR, LTNB, ARHM, SPKS, SSNS, ASCB	
6	<i>Anogeissus latifolia</i> Wall	54 SD±1.08 SE±0.54	103 SD±0.74 SE±0.43		+	+	+	LHTS, LABS, SPKS, SSNS, LMTC	
7	<i>Buchanania lanzan</i> Spreng	35 SD±0.71 SE±0.36	67 SD±0.17 SE±0.10		-	-	+	LFSC, LAGR, ASCB, LMSS	
8	<i>Lannea coromandelica</i> (Houtt.) Merr.	58 SD±0.76 SE±0.38	111 SD±1.13 SE±0.57		-	++	+	ASCB, LAGR, SSNS, LMSS, LOCT	
9	<i>Aegle mormelos</i> Correa.	33 SD±0.25 SE±0.15	86 SD±1.09 SE±0.55		+	-	-	LFST, LAGR, ASCB, LABS	
10	<i>Albizia lebbek</i> Benth.	62 SD±0.62 SE±0.36	121 SD±0.31 SE±0.18		+	-	-	ASCB, LAGR, SSNS, LMSS, LOCT	
11	<i>Milusa tomentosa</i> (Roxb.) J. sinclair	41 SD±0.72 SE±0.36	52 SD±0.27 SE±0.15		++	-	-	LETC, LFSC, LAGR, LMSS	
12	<i>Bauhinia retusa</i> Ham.	53 SD±1.29 SE±0.64	97 SD±0.55 SE±0.32		-	-	+	LFSC, LAGR, ASCB	

Table 4. Occurrence of different VAM Species at three different forest vegetation

<i>Jorhat (site-I)</i>	<i>Mandla (Site-II)</i>	<i>Sagar (Site-III)</i>
LFSC*, LABS*, LMSS*, LHTS*	LFSC*, LCVL, LGSP, LAGR*	LAGR*, LFSC*, LMSS*, LABS*
LAGR*, LOCT, LMST, ARHM	LETC, LMSS*, LDST, LAST	LTNB, LHTS*, LMST, LOCT
AUDL, ASPN, ATBC, SPKS	LHTS*, LABS*, LTNB, LMCC	LETC, ASCB, ARHM, CFLG
SSNS, CPLC, GRSC	CFLG, CPLC, ATBC, LFLV	SPKS, SSNS

*= common species, # =exclusive species

- species code of VAM fungi were followed after Perez and Schenk (1990)

LABS= *G. ambisporum* smith and schenck.

LTNB= *G. tenebrosus*(Thaxter) Berch.

LHTS= *G. heterosporum* Smith and shenck.

LMST= *G. multisubtensum* mukerji,

Bhattacharjee and Tiwari.

LOCT= *G. occultum* Walker.

LETC= *G. etunicatum* Becker and Gerdemann.

LCVL= *G. convolutum* Gerdemann and Trappe.

LGSP= *G. geosporum* (Nical and Gerd.) walber.

LDST= *G. deserticola* Trappe, Bloss and Menga.

LAST= *G. australe* (Berk) Berch.

LMCC= *G. macrocarpum* Tulasne and Tulasne.

LPST= *G. postulatum* Koske, friese. walker and Dalpe.

LMTC = *G. multicaule* Gerdemann and Bakshi.

ASCB= *Acaulospora Scrobiculata* Trappe.

ARHM= *A. rehmii* sieverding and Toro.

LAGR= *Glomus aggregatum* schenck and smith emend. koske.

LFSC= *Glomus fasciculatum* (Thaxter) Gerdemann and Trappe emend. walker and koshe.

LMSS= *G. moseae*(Nicolson and Gerdemann) Gerdemann and Trappe.

ATBC= *A. tuberculata* Janos and Trappe.

AUDL= *A. undulata* Sieverding.

CFLG = *scutellespora Fulgida* Koske and Walker.

CPLC= *Scutellospora pellucida* (Nicol & Schenck) walker and Sander.

SSNS = *Sclerocystis sinosa* Gerdemann and Bakshi.

GRSA = *Gisaspora rosea* Nicolson and Schenk

SPKS = *Sclerocystis pakistanica* Iqbal and Bushra

At site II, 14 tree species were selected for analysis of different attributes. Lowest root colonization was observed in *Bridelia retusa* Spreng. (33%) and highest (64%) in *Lagerastroemia parviflora* Roxb. followed by *Terminalia chebula* Retz. (57%). VAM fungal spore population per 25 g of soil collected from the rhizosphere of 14 tree species (Table 2) showed wide variations from 43 in *Bridelia retusa* Spreng to 167 in *Lagerastroemia parviflora* Roxb.

Data indicate that extramatrical hyphae were present in all the tree species except in *Terminalia chebula* Retz., *Adina cordifolia* Hook.F., *Bombox ceiba* Auct. and *Mitragyna parviflora* Korth.. Occurrence of vesicles was common however, not observed in *Bridelia retusa* Spreng., *Cassia fistula* Linn., *Schliechera oleosa* (Lour) Oken. and *Adina cordifolia* Hook. F.. Paris coil type association was a common feature in all the tree species barring *Diospyros melanoxylon* (Roxb.), *Shorea robusta* Gaertner F. and *Mitragyna parviflora* Korth.. Intraradical spores were found in *Bridelia retusa* Spreng, *Syzygium cuminii* (L.)Skeels, *Disopyros melanoxylon* (Roxb.) *Shorea robusta* Gaertner F., *Schliechera oleosa*(Lour) Oken, *Adina cordifolia* Hook. F. and *Embellica officinalis* Gaerth.

Total 16 VAM species were found at the site II, 6 species were exclusive to this site.

From site III, twelve tree species were considered and root colonization was minimum in *Wrightia tinctoria* Linn. (14%) and maximum (62%) in *Albizzia lebbek* Benth. followed by *Lannea coromandelica* (Houtt.) Merr. (58%). VAM fungal spore population per 25 g of rhizospheric soil for 12 selected tree species (Table 3) & showed

highest spore population (121/25 g) in *Albizzia lebbek* Benth. and lowest (29/25 g) in *Wrightia tinctoria* Linn.

External Hyphae were present in *Holoptelia integrifolia* Planch and *Anogeissus latifolia* Wall. Vesicles were found in 10 tree species except in *Miliusa tomentosa* and *Bauhinia retusa*. Paris type coils were recorded in the *Holoptelia integrifolia* Planch, *Flacoutia indica* (Burm F) Merr., *Butea monosperma* (Lamk. Taub.), *Anogeissus latifolia* Wall, *Lannea coromandelica* (Houtt.) Merr.. Intraradical spores were found within most of the tree species except in *Wrightia tinctoria* Linn., *Acacia catechu* Willd, *Aegle marmelos* Correa, *Albizzia lebbek* Benth., *Miliusa tomentosa* (Roxb.) J. Sinclair.

Total 14 VAM species were recorded from site III in which 2 VAM species were exclusive.

Total 25 VAM species were observed from all the three study sites. Out of which 15 were from site I, 16 from site II and 14 VAM species from site III (Table 4). 5 VAM species were common to all the three sites.

Discussion

Arbuscular mycorrhizal fungi are most widely distributed symbiotic organisms in nature. The soil borne mycorrhizal fungi colonize the root cortex biotrophically followed by the development of external mycelium which is a bridge connecting the root with the surrounding soil microhabitats. The VAM fungi have been reported from different plant communities of varied geographical regions such as – Tropical region (Chaurasia and Khare, 2000); Temperate region (Vestberg, 1995) and arctic region (Dalpe and Aiken, 1998).

Results of the present study indicate that all the tree species from tropical forests had VAM association with variation in the degree of infection. Such variability of colonization has been observed by number of workers in tropical forests and attributed to a number of local edaphic and climate factors (Safir and Dunisway, 1982; Thapar and Dunisway, 1992; Byra Reddy et al., 1994; Rahangdale and Gupta, 1998; Kalita et al., 2004).

Highest spore population was observed in rhizosphere soils of site II than at site I and site III. These differences can be attributed to the type of VAM species associated with their plants (Pringle and Bever, 2002). Seasonal variation in spores population in tropical forests depend upon the climatic factors, soil type, nutrients status growth and metabolic activities of plants, together with preferences of the fungi to host species and spatial distribution of host species (Louis and Lim, 1997; Harinikumar and Bagyaraj, 1988; Johnson and Wedin, 1997; Allen et al., 1998; Lovelock et al., 2003). Moreover, VAM spore densities vary greatly according to the physiological state of the host plant and relationship with the plant species. It is interesting that Arbuscules (Arum type) in VAM fungi were absent from all the three sites, although as a defining characteristic, arbuscules may be difficult to find under natural conditions due to their short life span (Smith and Read, 1997).

Paris type (Hyphal coil) association were more frequently observed. They were particularly prolific at the site II than in the other two sites. The dominance of Paris type association imply that coiled hyphae live longer and are more tolerant to stress conditions such as drought, suggesting that external factors determine the dominance of the type of endomycorrhizae (Paris Vs arum) over another.

The frequent occurrence of vesicles in most of the tree species from all the three study sites showed the presence of VAM fungi belonging to the Glomineae. The highest frequency of oval to elongate vesicles over the irregular and lobbed vesicles showed the dominance of species of the genus *Glomus* among other genera. Predominance of Glomineae has already been reported from tropical sites (Sieverding, 1991; Onguene and Kuyper, 2001). Results also showed the dominance of Glomineae over Gigasporineae. In the present study, the rhizosphere of different forest tree species from three forest sites showed common as well as variant VAM flora. Such variations are common features of VAM (Klironomos et al., 1993; Vishalakshi, 1997; Rawat and Mukerjee, 1998; Muthukumar and Udaiyan, 2000; Rama Bhat and Kaveriappa, 2001; Tamuli and Boruah, 2002).

A large number of VAM species belong to the genus *Glomus* in the present study. The genus *Glomus* has been reported to be the dominant VAM fungus in a number of forest communities (Sharma et al., 1986; Byra Reddy et al., 1994; Uniyal and Uniyal, 2000; Tamuli and Boruah, 2002; Mohan Kumar and Sivaswamy, 1992). This may be explained due to the wider adaptability of *Glomus* spp. as compared to other species (Sparling and Tinker, 1978; Sharma et al., 1986; Byra Reddy et al., 1994; Schenk and Kinloch, 1980).

Comparative assessment of occurrence of VAM fungi from three sites envisage that tropical moist forests (site II) harbour more VAM population in terms of richness and diversity than tropical evergreen and deciduous forest communities.

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