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

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(Received 16th November 2009; accepted 5th January 2012)

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 Wildlife 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 ± 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 Albizzia 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 reason 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 rooflets (feedar roots) were collected from the dominant species of each sites. 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* Roxb. (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.), *Mesua 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>
<thead>
<tr>
<th>S. No.</th>
<th>Plant</th>
<th>Quantitative Characters</th>
<th>Qualitative Characters</th>
<th>VAM Species</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>% Root colonisation</td>
<td>No. of spores in 25g soil</td>
<td>Exhamatric Hyphae</td>
</tr>
<tr>
<td>1</td>
<td>Dipterocarpus macrocarpus Vesque</td>
<td>29</td>
<td>SD±0.30 SE±0.15</td>
<td>85</td>
</tr>
<tr>
<td>2</td>
<td>Castanopsis spp. Blume</td>
<td>23</td>
<td>SD±0.45 SE±0.23</td>
<td>66</td>
</tr>
<tr>
<td>3</td>
<td>Dillenia indica Linn.</td>
<td>32</td>
<td>SD±0.38 SE±0.19</td>
<td>92</td>
</tr>
<tr>
<td>4</td>
<td>Lagerstromia spectrosa (L.)</td>
<td>22</td>
<td>SD±0.31 SE±0.15</td>
<td>105</td>
</tr>
<tr>
<td>5</td>
<td>Tetrameles nodifloras (R. Ar.)</td>
<td>32</td>
<td>SD±0.64 SE±0.32</td>
<td>94</td>
</tr>
<tr>
<td>6</td>
<td>Chukrasia tabularis (A. Juss.)</td>
<td>20</td>
<td>SD±0.42 SE±0.21</td>
<td>102</td>
</tr>
<tr>
<td>7</td>
<td>Amoora wallichii Benth.</td>
<td>28</td>
<td>SD±0.56 SE±0.28</td>
<td>99</td>
</tr>
<tr>
<td>8</td>
<td>Terminalia myrocarpa Henrick&amp;Mull.</td>
<td>33</td>
<td>SD±0.35 SE±0.18</td>
<td>133</td>
</tr>
<tr>
<td>9</td>
<td>Mesua ferrea Linnaeus</td>
<td>27</td>
<td>SD±0.95 SE±0.48</td>
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<tr>
<td>10</td>
<td>Artocarpus lakoocha Roxb.</td>
<td>27</td>
<td>SD±0.78 SE±0.39</td>
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<tr>
<td>11</td>
<td>Artocarpus chama Buch, Ham</td>
<td>30</td>
<td>SD±0.59 SE±0.31</td>
<td>122</td>
</tr>
<tr>
<td>12</td>
<td>Walsura robusta Roxburgh</td>
<td>19</td>
<td>SD±0.61 SE±0.31</td>
<td>57</td>
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</tbody>
</table>

Table 1. Quantitative and Qualitative characters of VAM fungi associated with some important forest tree species at Site-I (Jorhat).
Table 2: Quantitative and Qualitative characters of VAM fungi associated with some important forest tree species at Site-II (Mandla)

<table>
<thead>
<tr>
<th>S. No.</th>
<th>Plant Name</th>
<th>Quantitative Characters</th>
<th>Qualitative Characters</th>
<th>VAM Species</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>% Root colonization</td>
<td>No. of spores in 25g soil</td>
<td>Exhamatrica Hyphae</td>
</tr>
<tr>
<td>1</td>
<td>Bridelia retusa Spreng</td>
<td>33</td>
<td>SD±0.47 SE±0.24</td>
<td>43</td>
</tr>
<tr>
<td>2</td>
<td>Syzygium cumini (L.) Skeels</td>
<td>48</td>
<td>SD±1.28 SE±0.64</td>
<td>96</td>
</tr>
<tr>
<td>3</td>
<td>Diospyros melanoxylon (Roxb.)</td>
<td>43</td>
<td>SD±0.45 SE±0.23</td>
<td>84</td>
</tr>
<tr>
<td>4</td>
<td>Cassia fistula Linn.</td>
<td>41</td>
<td>SD±0.58 SE±0.29</td>
<td>103</td>
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<tr>
<td>5</td>
<td>Careya arborea (Roxb.)</td>
<td>52</td>
<td>SD±0.69 SE±0.35</td>
<td>91</td>
</tr>
<tr>
<td>6</td>
<td>Terminalia tomentosa W. &amp; A.</td>
<td>55</td>
<td>SD±0.39 SE±0.21</td>
<td>127</td>
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<tr>
<td>7</td>
<td>Shorea robusta Gaertner F.</td>
<td>60</td>
<td>SD±0.31 SE±0.15</td>
<td>157</td>
</tr>
<tr>
<td>8</td>
<td>Schleichera oleosa (Lour) Oken</td>
<td>46</td>
<td>SD±0.42 SE±0.21</td>
<td>83</td>
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<tr>
<td>9</td>
<td>Terminalia chebula Retz.</td>
<td>57</td>
<td>SD±0.77 SE±0.38</td>
<td>133</td>
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<tr>
<td>10</td>
<td>Adina cordifolia Hook.F.</td>
<td>39</td>
<td>SD±0.50 SE±0.25</td>
<td>139</td>
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<tr>
<td>11</td>
<td>Bombax ceiba Auct.</td>
<td>40</td>
<td>SD±0.50 SE±0.25</td>
<td>123</td>
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<tr>
<td>12</td>
<td>Lagerstroemia parviflora Roxb.</td>
<td>64</td>
<td>SD±0.49 SE±0.24</td>
<td>167</td>
</tr>
<tr>
<td>13</td>
<td>Embellica officinalis Gaerth.</td>
<td>53</td>
<td>SD±0.19 SE±0.11</td>
<td>91</td>
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<tr>
<td>14</td>
<td>Mitragyna parviflora Korth.</td>
<td>54</td>
<td>SD±0.66 SE±0.33</td>
<td>100</td>
</tr>
<tr>
<td>S. No.</td>
<td>Plant</td>
<td>Quantitative Characters</td>
<td>Qualitative Characters</td>
<td>Vesicle</td>
</tr>
<tr>
<td>-------</td>
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<td>------------------------</td>
<td>---------</td>
</tr>
<tr>
<td></td>
<td></td>
<td>% Root colonization</td>
<td>No. of spores in 25g soil</td>
<td>Exhamatrica Hyphae</td>
</tr>
<tr>
<td>1</td>
<td>Holoptelia integrifolia Planch</td>
<td>16 SD±0.89</td>
<td>47 SD±0.43</td>
<td>+</td>
</tr>
<tr>
<td>2</td>
<td>Flacourtia indica (Burn F) Merr.</td>
<td>29 SD±1.06</td>
<td>64 SD±0.90</td>
<td>-</td>
</tr>
<tr>
<td>3</td>
<td>Wrightia tinctoria Linn.</td>
<td>14 SD±0.85</td>
<td>29 SD±0.67</td>
<td>-</td>
</tr>
<tr>
<td>4</td>
<td>Acacia catechu Wild.</td>
<td>21 SD±0.60</td>
<td>33 SD±1.20</td>
<td>-</td>
</tr>
<tr>
<td>5</td>
<td>Butea monosperma (Lamk. Taub)</td>
<td>38 SD±0.49</td>
<td>99 SD±0.29</td>
<td>-</td>
</tr>
<tr>
<td>6</td>
<td>Anogeissus latifolia Wall</td>
<td>54 SD±1.08</td>
<td>103 SD±0.74</td>
<td>+</td>
</tr>
<tr>
<td>7</td>
<td>Buchanania lanzan Spreng</td>
<td>35 SD±0.71</td>
<td>67 SD±0.17</td>
<td>-</td>
</tr>
<tr>
<td>8</td>
<td>Lannea coromandelica (Houtt.) Merr.</td>
<td>58 SD±0.76</td>
<td>111 SD±1.13</td>
<td>-</td>
</tr>
<tr>
<td>9</td>
<td>Aegle mormelos Correa.</td>
<td>33 SD±0.25</td>
<td>86 SD±1.09</td>
<td>+</td>
</tr>
<tr>
<td>10</td>
<td>Albizia lebbek Benth.</td>
<td>62 SD±0.62</td>
<td>121 SD±0.31</td>
<td>+</td>
</tr>
<tr>
<td>11</td>
<td>Milicia tomentosa (Roxb.) J. sinclair</td>
<td>41 SD±0.72</td>
<td>52 SD±0.27</td>
<td>+</td>
</tr>
<tr>
<td>12</td>
<td>Bauhinia retusa Ham.</td>
<td>53 SD±1.29</td>
<td>97 SD±0.55</td>
<td>-</td>
</tr>
</tbody>
</table>
Table 4. Occurrence of different VAM Species at three different forest vegetation

<table>
<thead>
<tr>
<th>Jorhat (Site-I)</th>
<th>Manda (Site-II)</th>
<th>Sagar (Site-III)</th>
</tr>
</thead>
<tbody>
<tr>
<td>LFSC*, LABS*, LMSS*,</td>
<td>LFSC*, LCVL, LGSP,</td>
<td>LAGR*, LFSC*, LMSS*,</td>
</tr>
<tr>
<td>LHTS*</td>
<td>LAGR*</td>
<td>LABS*</td>
</tr>
<tr>
<td>LAGR*, LOCT, LMST,</td>
<td>LETC, LMSS*, LDST, LAST</td>
<td>LTNB, LHTS*, LMST, LOCT</td>
</tr>
<tr>
<td>ARHM</td>
<td></td>
<td></td>
</tr>
<tr>
<td>AUDL, ASPN, ATBC, SPKS</td>
<td></td>
<td></td>
</tr>
<tr>
<td>SSNS, CPLC, GRSC</td>
<td>CFLG, CPLC, ATBC, LFLV</td>
<td>SPKS, SSNS</td>
</tr>
</tbody>
</table>

* = common species, # = exclusive species

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

LABS = G. ambisporum smith and schenck.
LTNB = G. tenebrous(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, Bless and Manga.
LAST = G. australe (Berk) Berch.
LMCC = G. macrocarpum Tulasne and Tulasne.
LPST = G. postulatum Koske, friese. walker and Dalpe.
LPC = G. multicaule Gerdemann and Bakshi.
ASCB = Acaulospora Scrobiculata Trappe.
ARM = A. rehmi 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.

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 Diospyros melanoxylon (Roxb.), Shorea robusta Gaertner F. and Mitragyna parviflora Korth.. Intraradical spores were found in Bridelia retusa Spreng, Syzygium cumini (L.)Skeels, Diospyros 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 Albizia 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 *Albizia lebbeck* 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 coronandra* (Houit.) Merr.. Intraradical spores were found within most of the tree species except in *Wrightia tinctoria* Linn., *Acacia catechu* Willd, *Aegle marmelos* Correa, *Albizia lebbeck* 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 biotropically 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 Mukerji, 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.

Acknowledgements. Author wish to thank Dr. P. K. Khatri, Tropical Rain Forest Institute, and Dr. R.K. Shukla, Kanha Tiger reserve, Mandla, for providing field facility. One of the authors (Priyangana Tripathi) is thankful to University Grants Commission, New Delhi, for the award of fellowship and financial assistance.

REFERENCES


[33] See www.invam.caf.wvu.edu/my-info/Taxonomy/species.htm/.


