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Exclosures restored the density and root colonization of arbuscular mycorrhizal fungi in Tigray, Northern Ethiopia

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Abstract

Introduction: Exclusion of grazing animals and tree plantations were among the methods used for the restoration of degraded lands in tropical semiarid areas. Exclosures can foster secondary forest succession by improving soil conditions and modifying microclimate for understory growth. This paper compared the arbuscular mycorrhizal fungi (AMF) spore density, root colonization of woody plants and soil chemical properties under exclosure with increasing age, and grazing land at different slope positions.

Methods: The study was conducted in northern Ethiopia from 12 exclosure sites paired each with adjacent grazing land in total from 24 sites with four treatments replicated three times. In the entire study, 216 plots were examined of which 108 were in exclosures and 108 in communal grazing lands. There were four age classes and three slope positions in each of the land uses. Composite soil and root samples were collected using nested plots measuring 100m² from four sides of 1763 plants for spore enumeration and root colonization. Soils for chemical properties were collected from the four corners and center of 5 m × 5 m plots.

Results: All the 61 woody plant species that belong to 41 families were colonized by AMF. Spore density and root colonization were significantly higher in exclosures as compared to grazing land and increased with increasing ages of exclosures. Foot slope had significantly higher spore density and root colonization than middle and upper slopes. Soil chemical properties were significantly higher in the exclosure, oldest age of exclosure, and foot slope position (except P). AMF spore density and root colonization were significantly positively correlated with soil chemical properties.

Conclusions: Exclosures are helpful to restore the AMF spore density and root colonization of woody species and soil fertility.

Keywords: Spore density, Root colonization, Age of exclosure, Slope position

Introduction

Exclosures are areas closed off from the interference of human and domestic animals with the goal of promoting natural regeneration of plants and reducing land degradation of formerly degraded communal grazing lands (Nedessa et al. 2005). They are usually established in steep, eroded, and degraded areas that have been used for

grazing in the past (Descheemaeker et al. 2006a, b). In Ethiopia, the inception of exclosures dates back to the 1980s and coincided with the introduction of large-scale land rehabilitation and soil and water conservation program (Nedessa et al. 2005). The main objective of establishing exclosures on communal grazing lands is to improve the overall ecological conditions of the degraded areas so that they can provide better environmental and socioeconomic benefits to the local communities. Several case studies have reported beneficial effects of exclosures: natural vegetation has regenerated, runoff, and sheet and rill erosion are reduced, the land is stabilized, water availability has increased, and soil physical and chemical

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properties are improved (Girmay et al. 2009; Mekuria et al. 2009; Mekuria et al. 2007; Yami et al. 2007; Descheemaeker et al. 2006a, b). These positive consequences are also noticed and appreciated by the local communities (Mekuria et al. 2009).

The impact of grazing on soil properties depends on grazing management, the non-grazed control site provided higher soil pH, available P, and Mg compared with moderately grazed site of 33 years old (Ayorlo et al. 2011). Available phosphorus (P), total nitrogen (N), calcium (Ca), magnesium (Mg), and potassium (K) decreased after 1.5 years of open grazing compared with an ungrazed control in a tropical pasture (Ayorlo et al. 2011). In addition, open grazing resulted in lower water infiltration (Hiernaux et al. 1999) and higher soil loss (Tadesse and Penden 2002) compared with moderately grazed sites. Exclosures with 12 years old enhance the total plant cover, dry matter yield, species richness, and contents of soil organic matter, total nitrogen, and water infiltration rate compared with continually grazed area (Jeddi and Chaieb 2010). Exclusion of livestock grazing for 20 years significantly increased aboveground and belowground biomass and species richness for five different communities compared with non exclosed sites in Loess Plateau, northwest China (Cheng et al. 2011). The exclosures had twice more plant species richness and diversity value as compared with communal grazing lands after 22 years of exclosure establishment in the central and northern highlands of Ethiopia (Mengistu et al. 2005). There were increases in the number of woody species richness after 8 years of exclosure establishment (Birhane et al. 2006), 1.1% increase in soil organic matter, 0.1% in total N, 1.8 mg kg⁻¹ in available P increase after 10 years of exclosure establishment (Mekuria et al. 2007). A considerable decrease in soil loss was reported after the establishment of exclosures on communal grazing lands (Girmay et al. 2009; Mekuria et al. 2009; Descheemaeker et al. 2006a, b). Exclosures have been reported to positively contribute to restore soil biological properties such as arbuscular mycorrhizal fungi (Birhane et al. 2010).

Arbuscular mycorrhizal fungi (AMF) colonize about 80% of plants (Wang and Zao 2008) and they are regular component of rhizosphere micro-flora in natural ecosystem (Sharma et al. 2009). AMF form obligate symbiotic association with many tropical plants (Manoharachary et al. 2005). The function of AMF in exclusion process and knowledge of AMF diversity in a specific area is essential to utilize AMF in any application (Wang and Zao 2008). The very few conditions where infective AMF are low in density and diversity is when the soil erodes, is disturbed, and is devoid of vegetation cover (Asmelash et al. 2016). The presence of enough AMF propagules in a restored site facilitates the establishment

of plant communities through increased availability of nutrients and moisture to plant roots (Asmelash et al. 2016). Though exclosures positively contributed to land rehabilitation, there are limited studies on the role of exclosures to restore arbuscular mycorrhizal fungi. There is a limited study that attempted to understand the role of exclosure in the restoration of AMF with increasing age of exclosures at different slope positions during the rainy season. The slope position significantly affects the density of spores found in a given landscape. The spore density was found to be higher from exclosures of middle slope position in the dry season (Birhane et al. 2017). We investigated the spore density in the soils and woody plant root colonization of AMF from paired exclosure and adjacent grazing land with increasing age of exclosures at the foot, middle, and upper slope hillsides. The correlation between the availability of soil nutrients with AMF spore density and root colonization was assessed. We aimed to answer the following questions: What is the dynamics of AMF with increasing age exclosures at different slope positions? Does the availability of nutrients correlated with AMF spore density and root colonization? And does the increase in the age of exclosures increased the availability of nutrients at different slope positions?

Methods

Study area

The study was conducted in the highlands of Tigray region, Northern Ethiopia in 12 sites representing four zones and four districts (Table 1). All sites have tropical semi-arid climate. The altitude of the study sites ranged from 2232 to 2937 m.a.s.l. The rainy season usually occurs between June and September (Fig. 1), and the growing season varies between 90 and 120 days. The highest rainfall is in July and August which ranges between 162 and 228 mm. The temperature of the study sites ranges from 19 to 30 °C, the maximum is recorded in the months of May and June (Fig. 1).

Luvisols (Alfisols), Regosols (Entisols), Cambisols (Inceptisols), and Calcisols (Aridisols) were major soil groups in the study area (WRB 2006). The 24 study sites were dominated by Luvisols (Alfisols) and Cambisols (Inceptisols). *Acacia etbaica*, *Acacia seyal* (Del.), *Becium grandiflorum* (Lam.) Pichi-Serm., *Euclea racemosa subsp. schimperi* (A. DC.) Dandy, and *Maytenus arbutifolia* (Hochst. ex. A. Rich) Wilczek were the woody vegetation species common in exclosures and adjacent grazing lands (Mekuria 2010).

Mixed farming system that integrates crop and livestock was the main means of livelihood. The major land uses were cultivated land, forest land, exclosure, and communal grazing land (Mekuria 2010). The main crops cultivated were Teff (*Eragrostis teff* (Zucc.) Trotter),

Table 1 Specific study site age, altitude and geographic location

Specific site	Zone	District	Age (year)	Altitude (m)	Geographic location
Adihintaweinai	East	Atsbi wonberta	< 10	2201–2312	39° 38' 60"–39° 50' 52" East 13° 12' 75"–14° 4' 41" North
Halla	Southeast	Degua Tembien	< 10	2232–2937	38° 30' 17"–38° 40' 57" East 13° 54' 3"–14° 20' 30" North
Melgim	East	Atsbi wonberta	< 10	2264–2343	39° 38' 60"–39° 50' 52" East 13° 12' 75"–14° 4' 41" North
Endagebriel	East	Atsbi wonberta	10–20	2248–2351	39° 38' 60"–39° 50' 52" East 13° 12' 75"–14° 4' 41" North
Gurzoemni	Central	Tahtay Maichew	10–20	2244–2322	38° 30' 17"–38° 40' 57" East 13° 54' 3"–14° 20' 30" North
Mezewle	East	Atsbi wonberta	10–20	2325–2411	39° 38' 60"–39° 50' 52" East 13° 54' 3"–14° 20' 30" North
Adikolakul	Southeast	Degua Tembien	20–30	2180–2214	38° 30' 17"–38° 40' 57" East 13° 54' 3"–14° 20' 30" North
Suhulkoma	East	Atsbi wonberta	20–30	2295–2347	39° 38' 60"–39° 50' 52" East 13° 12' 75"–14° 4' 41" North
Wereriba	Southeast	Doguetembien	20–30	2200–2358	13° 16' 23"–13,047' 44" East 39° 3' 17"–39,024' 48" North
Gratselim	Southeast	Degua Tembien	30–40	2369–2458	38° 30' 17"–38° 40' 57" East 13° 54' 3"–14° 20' 30" North
Kerenadidemsash	South	Endamekoni	30–40	2314–2419	39° 16' 52"–39° 35' 31" East 12° 38' 4"–12° 51' 39" North
Maibiati	Southeast	Degua Tembien	30–40	2358–2429	38° 30' 17"–38° 40' 57" East 13° 54' 3"–14° 20' 30" North

Bread wheat (*Triticum aestivum*), Maize (*Zea mays L.*), Sorghum (*Sorghum bicolor*), Barley (*Hordeum vulgare*), and Faba bean (*Vicia faba*). The main animal which graze the communal grazing lands were cattle, goats, sheep, donkeys, camel, mule, and horse.

Experimental design

The age of restoration, slope position, and land use effect on AMF spore density, woody plant root colonization, and soil chemical properties were studied from 12

closure sites paired each with adjacent grazing land during the dry season. The distance between the enclosure and the adjacent grazing land was in the range of 50 to 100 m. There were four age classes and three slope positions in each of the land uses. The first age group had less than 10 years old enclosure with triplicate sites (Halla, Meligim, and Adihantaweynay), the second age group was 10 to 20 years old enclosure with triplicate sites (Gurzoemni, Mezewle, and Endagebriel), and the third age group had 20 to 30 years old enclosure with triplicates

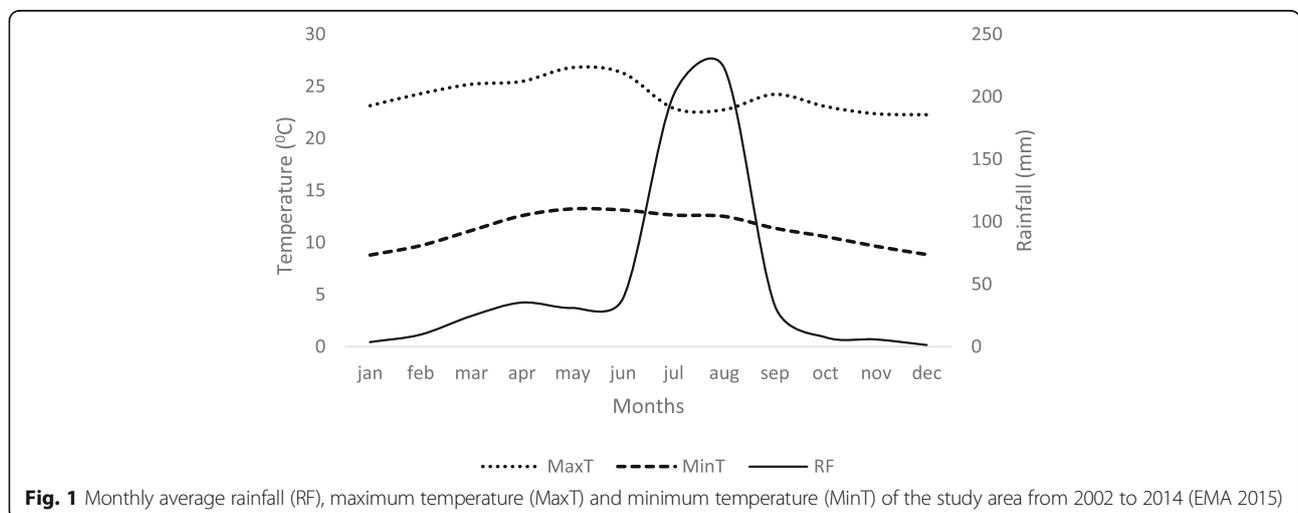


Fig. 1 Monthly average rainfall (RF), maximum temperature (MaxT) and minimum temperature (MinT) of the study area from 2002 to 2014 (EMA 2015)

sites (Addikolakul, Wereriba, and Shul-koma). The fourth group had greater than 30 years old enclosure with triplicate sites (Maybe'ati, Kerenadidemsash, and Gratselim). The assumption were prior to establishment, enclosures and the adjacent grazing lands had similar biophysical, socioeconomic, and management conditions as the enclosures were established on the same communal grazing lands which were used for livestock grazing (Mekuria 2010). The experiment was composed of 12 experimental units with four treatments replicated three times. Site had two levels (closed and open), age with four levels (<10, 10–20, 20–30, and >30 years old) with three slope positions (upper, middle, and foot slope). In the entire study, 216 plots (12 × 3 small plots × 3 slope positions × 2 pair enclosure and adjacent grazing land) were examined of which 108 were in enclosures and 108 in communal grazing lands. The small plots used to collect soil samples. In each enclosure and grazing land randomly established three transects spaced at a minimum distance of 75 m (Mekuria 2010).

The number of transects were based on vegetation density, spatial heterogeneity of vegetation, and area of the site. The area of the selected enclosures ranged from 7 to 85 ha, whereas that of the adjacent communal grazing lands ranged from 2 to 38 ha. To avoid edge effects, the first transect was laid 30–50 m inside the enclosures and grazing lands. Transects were parallel to each other and to the topography of the landscape. In each transect, three slope positions (upper slope (US), mid slope (MS), and foot slope (FS)) were delineated and in each landscape position, a sampling plot measuring 10 m × 10 m was established. In each plot, 5 m × 5 m subplots for soil physical and chemical analysis were developed. The US position is the uppermost portion of each study site, and it can receive little or no overland flow but may contribute runoff to down slope areas. The MS position receives overland flow from the upper slope and contributes runoff to the FS. The FS represents the lowest part of each study site and receives overland flow from both mid and upper slopes.

Plant and soil sampling methods

The plant roots and soils were sampled during the dry season from October 2013 to January 2014 for a total of 4 months from enclosures and adjacent surrounding grazing lands. All woody plants were sampled from the 10 m × 10 m quadrates (100 m²) for trees and shrubs from both land uses. Woody plants were replicated three times in each plot. Composite soil and root samples were collected for spore density and root colonization from a total of 1763 samples from 4 sides of each plant found in each plot from the enclosures and adjacent grazing lands. The identity of the plant was identified at the field and verified using reference books such as

Bekele (2007), Inga et al. (2003), Sue et al. (1995, 2000), and Inga and Sue (1989).

Soil samples were collected from 5 m × 5 m subplots nested within the 10 m × 10 m at the center of the main plots. A total of 216 soil samples were collected. The soil samples for one plot was taken from the four corners and the center of a square plot at 0–50 cm soil depths following an “X” pattern of the main plot to form one composite sample in order to determine texture, soil organic carbon, pH, EC, N, K, and P in enclosures and adjacent grazing lands. The five soil samples measured 300 g each were mixed and form 1500 g in total and make single composite sample to represent the sample plot. From the composite sample, 1 kg soil was taken and was put into plastic bags, secured, labeled and brought to the soil laboratory. Soil samples were analyzed for pH and electrical conductivity on 1:2.5, soil: water suspension method. Organic matter was analyzed using the Walkley-Black method (Van Ranset et al. 1999), total nitrogen content by the Kjeldahl method (Bremmery and Mulvaney 1982). Available potassium and texture were analyzed with flame spectrophotometer and hydrometer method (Gee and Bauder 1982). Available P determined using the Olson method (Olsen and Sommers 1982).

AMF root length colonization and spore density sampling

Root samples of all plants were collected by excavating soil using a handheld hoe starting from the plant's trunk base in four directions of the plant and working out towards to get fine roots within 3–5 m radius. Forty-gram root samples from all four directions were collected from 0 to 30 cm depth, mixed and then 10 g fine roots (diameter < 2 mm) were taken from the composite root samples. The collected fine roots were put into beaker and carefully washed with tap water until the soil was removed and became free of any soil particles and put into plastic jar and filled with 97% ethanol to preserve the roots until they were ready for processing. Soil samples were taken at 50 cm depth using auger (Chanie and Assefa 2013) in four directions of each plant and gave 300 g soil sample, in total 1200 g soil sample were mixed to make one composite soil sample. For each plant, 1 kg composite soil samples was taken from the 1200 g of soil and put in a sealed plastic bag. The sampled roots and soils were transported to Mekelle University laboratory for estimating root colonization and spore extraction.

AMF spores were separated from the soil by the wet sieving and 50% sucrose centrifugation method (Brundrett et al. 1996). For spore extraction, the soils first air-dried and pass through 0.75 mm sieve and weighed using sensitive balance to get a subsample of 25 g. The weighed dry soil was wet with enough amount of water for at least 30 min. The wet soil was stirred for 15 to

30 min and then decanted through a series of sieves by putting in ascending order with the smallest sieve size (50 μm) at the bottom, followed by 100, 300, and 750 μm sieve size at the upper most. The 750 μm sieve size used to remove unnecessary materials. The different sieve sizes were given codes as 50 μm = A, 100 μm = B, and 300 μm = C, to designate the spore sizes captured by each sieve sizes. Spores were counted from 25 g soil using dissecting microscope with $\times 100$ magnification.

AMF spores were grouped into genera using their morphological characteristics such as spore size, shape, color, wall structure, and hyphal attachment (simple, swollen, or bulbous). Permanent slides were prepared for each different spore morphotype with polyvinyl alcohol plus Melzer's solution. The PVLG mountant was prepared using polyvinyl alcohol 8.33 g (polyvinyl alcohol should be used and can be dissolved in water by heating at 90 $^{\circ}\text{C}$ overnight), distilled water 50 ml, lactic acid 50 ml, and glycerin 5 ml) and prepared Melzer's reagent (1.5 g iodine, 5 g potassium iodide, and 100 ml distilled water). The Melzer's reagent then mixed 1:1 (v/v) with PVLG (Brundrett et al. 1996). Small droplets of the mountant was added using pipet (1:1 PVLG and Melzer's reagent) on to the microscope slide then the spores put on it using paint brush and pressed with coverslip to easily see and identify the internal structures of the spores. The genera of each spore were identified using a compound microscope with $\times 400$ magnification based on descriptions in Brundrett et al. (1996) and information from the INVAM website (<http://www.invam.caf.wvu.edu>).

The procedure in Brundrett et al. (1996) was used to determine root colonization. To identify root colonization, the roots were washed to remove ethanol, cut into 1 cm length, and insert into the heat resistant bottle containing 10% KOH solution then autoclaved for 20 min at 120 $^{\circ}\text{C}$. The roots were washed to remove KOH and put into 10% H_2O_2 for further bleaching and clearing for about 15 min. Cleared roots were captured on a fine sieve and rinsed with water before transferring them into the HCl solution. Roots were acidified with 3% HCl for about 30 min and stained in trypan blue (0.05% in 5:1:1 lactic acid: glycerol: distilled water) over night. The stained roots were washed and added to 50% glycerol for destaining and preserved until further processing. Then roots were prepared on slide lengthwise by selecting 20 subsample roots. The proportional root colonization by AMF was estimated using the magnified intersection method with hair line graticule inserted into an eyepiece acted as the line of intersection

with each root at 400 magnification power under the compound microscope. Percentage of root length colonization (RLC %) was calculated from 100 or more intersections for each root sample. At each intersection, there were six possible mutually exclusive outcomes (Brundrett et al. 1994).

Statistical analyses

The differences in spore density, root colonization, and soil parameters between an enclosure and its adjacent communal grazing land at different age groups and landscape position were assessed using two- and three-way ANOVA with Tukey HSD test after checking normality test. Pearson correlation test was used to conduct relationships between spore density and root colonization and soil parameters. Statistical package for social sciences (SPSS) version 20 was used to analyze spore density and root colonization of AMF and the soil.

Results and discussion

AMF spore density and root colonization between enclosures and adjacent grazing lands

Six AMF genera were recovered and identified in enclosures (Table 2). *Glomus* was the dominant genus found in all soils followed by *Acaulospora*, *Gigaspora*, *Scutellospora*, *Entrophospora*, and *Sclerocystis*. *Glomus*, *Acaulospora*, and *Gigaspora* were found in all the 12 sites followed by *Scutellospora* (11), *Entrophospora* (9), and *Sclerocystis* (1). This finding was similar to Vyas and Gupta (2014) where they found similar dominance order of genera. Enclosures had significantly higher spore density (4385 spores 100 g^{-1} dry soil and 6 genera) than grazing land (176 spores 100 g^{-1} dry soil and 5 genera). The genus *Sclerocystis* was retrieved from enclosure but not in the samples from grazing land.

Enclosures had higher spore density mainly dominated by the genus *Glomus* in Northern Ethiopia (Birhane et al. 2010). The low spore density in grazing land could be due to high soil disturbance, changes in land use and management, and changes in plant community composition (Birhane et al. 2017; Santos-González et al. 2007).

The spore density recovered in this study was higher than the total number of spores recovered from forest and grazing land (Chris 2010). *Glomus* was the most dominant genus in enclosure and in grazing land. *Glomus* is a predominantly genus found to be distributed in soils all over the world (Minal and Anil 2012; Chris 2010) and is the dominant genus in disturbed soils (Tanvir et al. 2011;

Table 2 Spore density and spore genera in enclosures and grazing land (100 g^{-1} dry soil)

LU	<i>Gigaspora</i>	<i>Scutellospora</i>	<i>Glomus</i>	<i>Sclerocystis</i>	<i>Acaulospora</i>	<i>Entrophospora</i>	Sum
EX	206.3 \pm 5.5a	105.2 \pm 3.3a	2468.2 \pm 42.2a	0.013 \pm .009a	1582.2 \pm 30.8a	23.2 \pm 1.3a	4385.3 \pm 80.4a
GL	2.3 \pm 8.5b	0.271 \pm 5.1b	110.2 \pm 64.3b	0.000 \pm .013a	63.3 \pm 47.0b	0.068 \pm 1.9b	176.3 \pm 122.5b

Means followed by the same letter in the same column are not significantly different at $P < 0.05$ level, mean \pm SEM
GL grazing land, EX enclosure, LU land use types

Table 3 Root length colonization (%) of AMF between exclosures and open grazing lands

Land use types	HC	MHC	AC	VC
Exclosure	78.4 ± 0.4a	70.1 ± 0.5a	50.3 ± 0.5a	38.1 ± 0.5a
Grazing land	55.7 ± 0.6b	47.5 ± 0.8b	35.5 ± 0.8b	23.4 ± 0.8b

Means in the same column followed by same letter do not differ significantly at $P < 0.05$, mean ± SEM

HC hyphal colonization, MHC mycorrhizal colonization, AC arbuscular colonization, VC vesicular colonization

Burni et al. 2009; Sharma et al. 2009; Wang and Zao 2008; Mridha and Dhar 2007; Panwar and Tarafdar 2006; Burni and Illahi 2004). The predominance of *Glomus* species under varying soil conditions might be due to its wide adaptability to the varied soil conditions and survive in acidic as well as in alkaline soils (Tanvir et al. 2011; Pande and Tarafdar 2004).

Spore density in undisturbed natural environment is higher than disturbed sites (Chanie and Assefa 2013; Muleta et al. 2008). AMF plays a role in the formation of stable soil aggregates, building up a macroporous structure of soil that allows penetration of water and air and prevents erosion (Rillig 2004; Borie et al. 2008; Aggarwal et al. 2011); therefore, the application of AMF is of interest for the reclamation and revegetation of degraded lands (Miller and Jastrow 1992).

The influence of grazing on soil nutrient availability and host plant productivity (Frank and McNaughton 1993) may cause variable effects on AMF community composition and structure (Bai et al. 2013; Eom et al. 2001). Grazing intensity might change the level of mycorrhizal infection in a community by altering the plant composition and is an important factor in regulating the cycling of nutrients in undisturbed ecosystems (Abbott and Robson 1991). Grazing can influence the dynamics of nutrient exchange between host plants and AMF (Gehring and Whitham 2002). Herbivore grazing can alter leaf photosynthetic rates (McNaughton 1979), the above ground production (Frank and McNaughton 1993) and the C allocation below ground (Frank et al. 2002). The allocation of AMF morphological structures can either increase or decrease depending on the timing and severity of herbivores (Gange 2007).

Land use significantly affected AMF root colonization (Table 3). AMF root colonization was significantly higher

in exclosures compared to the adjacent grazing land ($P < 0.05$). Hyphal colonization was the highest followed by MHC and AC while the least was VC. The grazing land showed low root colonization, and the higher colonization was for hyphae followed by MHC, AC, and VC, respectively.

AMF spore density and root colonization with increasing age of exclosures and slope position

Spore density increased significantly as the age of exclosure increased ($P < 0.05$). An exclosure with greater than 30 age class counted higher spore density followed by age class between 20 and 30, age class between 10 and 20, and age class less than 10, respectively (Table 4). AMF spore density ranged from 25 to 2550 in 100 g^{-1} of dry soil with increasing stability of ecosystem (Zhao et al. 2003). High AMF diversity was found in undisturbed and old woodlands and forests (Don-Rodrigue et al. 2013; Mathimaran et al. 2007; Muthukumar and Udaiyan 2000; Wilson et al. 1992). *Glomus*, *Acaulospora*, and *Gigaspora* were highly significantly different in all age classes while *Scutellospora* and *Entrophospora* significantly differed only among age classes greater than 30, age class from 20 to 30, and 10–20 (Table 4). Except *Sclerocystis* and *Entrophospora*, all the genera were found in all age classes. The dominant genera at all stages of wheat crop development were *Acaulospora*, *Glomus*, and *Scutellospora* (Arpita et al. 2012; Schalamuk et al. 2006) and the dominant genera in tropical rainforest were *Acaulospora* and *Glomus* (Zhao et al. 2003). The increased trend in spore density with increased age of exclosures is related to the low disturbance and high organic matter in old age exclosure (Burni and Illahi 2004; Anwar and Jalaluddin 1991). High organic matter increases the water holding capacity of soil which may enhance the sporulation of AMF.

All AMF genera showed significant difference between slope positions except for *Sclerocystis* (Table 5). The foot and middle slope had six genera whereas the upper slope had five genera (Table 5). The highest spore density was recorded at the foot slopes followed by the middle and upper slopes. The interactions between age × land use, age × elevation, land use × elevation, and age × land use × elevation also showed significant difference in spore density ($P = 0.000$). This high spore density and genera

Table 4 AMF spore density and spore genera in four age classes of exclosures (100 g^{-1} of dry soil)

Age (year)	<i>Gigaspora</i>	<i>Scutellospora</i>	<i>Glomus</i>	<i>Sclerocystis</i>	<i>Acaulospora</i>	<i>Entrophospora</i>	Sum
< 10	6.1 ± 10.6d	2.9 ± 6.3c	326.2 ± 91.7d	0.000 ± 0.01a	226.6 ± 64.2d	0.000 ± 2.5c	562.0 ± 169.3d
10–20	39.7 ± 8.4c	13.3 ± 5.0c	1135.9 ± 72.6c	0.000 ± 0.01a	603.4 ± 50.8c	2.5 ± 2.0c	1794.9 ± 134.1c
20–30	162.7 ± 8.5b	75.6 ± 5.1b	2002.2 ± 74.1b	0.000 ± 0.01a	1259.3 ± 51.8b	13.6 ± 2.0b	3513.6 ± 136.7b
> 30	300.8 ± 7.9a	165.2 ± 4.7a	2903.5 ± 68.6a	0.030 ± 0.01a	1976.0 ± 48.0a	39.9 ± 1.9a	5385.6 ± 126.6a

Means followed by the same letter in the same column are not significantly different at $P < 0.05$ levels, mean ± SEM

Table 5 AMF spore density (100 g⁻¹ of dry soil) across foot, middle, and upper slope position

Slope	<i>Gigaspora</i>	<i>Scutellospora</i>	<i>Glomus</i>	<i>Sclerocystis</i>	<i>Acaulospora</i>	<i>Entrophospora</i>	Sum
Foot	192.3 ± 8.2a	109.2 ± 4.8a	2578.0 ± 66.0a	0.006 ± 0.01a	1747.6 ± 45.6a	22.8 ± 1.8a	4650.1 ± 122.9a
Middle	133.8 ± 9.0b	60.3 ± 5.2b	1491.4 ± 72.1b	0.021 ± 0.01a	947.4 ± 49.9b	14.2 ± 1.9b	2647.3 ± 134.3b
Upper	96.8 ± 9.2c	42.8 ± 5.4c	1006.4 ± 4.1c	0.000 ± 0.01a	527.1 ± 51.2c	10.1 ± 2.0b	1683.4 ± 137.9b

Means followed by the same letter in the same column are not significantly different at $P < 0.05$ level, mean ± SEM

richness increment from upper to down slope might be due to natural forces such as flood and wind. AM fungal spores rely on some forces that can move soil, such as strong winds, water, gophers, and worms, to migrate to a new location (David and Jeff 2010). Removal of surface soil layers mainly by water erosion decreased markedly both the number of propagules of AMF and the extent of mycorrhiza formation (Habte 1989; Powell 1981) thereby increase spore density and genera diversity down slope gradient.

All the age groups in both land uses exhibited all structures of AMF. The root colonization showed significant difference among the four age classes (Table 6). There was higher colonization in % HC, % MHC, % AC, and % VC with increasing age from < 10 years of age to 10–20 years of age followed by 20–30 years of age, and the highest colonization was recorded in the age group > 30 years old. The root colonization of *Juniperus procera* ranged from 59 to 79% and increased with the age of the tree (Amal and Hasnah 2012). Natural environment increased spore density and root colonization, indicating stability of ecosystems (Megan and Kirkegaard 2012).

Plant roots at the foot slope showed higher colonization compared to the plants at the middle and upper slope positions. Mycorrhizal hyphal colonization was significantly different between the foot slope and the other two slope position ($P < 0.05$). The interaction between age × land use and land use × elevation were sources of variation ($P = 0.000$) in root length colonization. Soil disturbance, degradation, and erosion can reduce levels of AMF propagules (Birhane et al. 2010; Carpenter et al. 2001; Brundrett et al. 1996). Vesicular, arbuscular, and hyphal colonization in soils of soybean plant root found colonized with AMF in 133 samples out of 167 samples examined at different

slope positions (Bansode et al. 2014). Current distribution and density of AMF are the result of contemporary ecological processes that are under control of several factors such as altitude, soil chemical properties, soil disturbance, and above-ground vegetation (Sturmer and Siqueira 2011; Yang et al. 2010; Shukla et al. 2009).

AMF in different woody plant species and their families

All rhizosphere soil samples under each plant species generally displayed higher spore density as compared to other studies and were shown significant variation in spore density between plant species and their family (Table 7), though the grazing intensity varies with species. The highest spore density was found under *Ficus vasta*, *Cordia africana*, *Ekebergia capensis*, and *Justicia schimperiana* while *Aloe vera*, *Opuntia ficus-indica*, and *Rumex nervosus* exhibited the lowest spore density. Families of Acanthaceae, Boraginaceae, Meliaceae, and Moraceae had significantly higher spore density 100 g⁻¹ dry soil while lowest spore counts were found in Aloeaceae, Cactaceae, and Polygonaceae rhizosphere soils. Low AMF spore count was reported by Muleta et al. (2008) for coffee and in a dry savannah wood land ecosystem under *Acacia polyacantha* rhizosphere soil (Yonas 2005).

Glomus was found in all the 32 plant families and 47 plant species followed by *Acaulospora*, *Gigaspora*, *Scutellospora*, *Entrophospora*, and the lowest found in *Sclerocystis*, respectively (Additional file 1: Table S1 and S2). *Glomus* is the most common AMF genus distributed globally and dominant in the tropical areas (Chaurasia 2000). Wide occurrence of *Glomus* in the present study suggested that the genus *Glomus* has wide ecological amplitude that is responsible for its adaptability and survival in different habitats and vegetation composition. The dominance of *Glomus* species has been reported from tree species in Ethiopia (Birhane et al. 2010; Wubet et al. 2003, 2009; Muleta et al. 2008). The variation in spore density between plants species may be due to differences in host trees which can be attributed to the type of AMF species associated with their respective hosts (Cardoso et al. 2003; Pringle and Bever 2002).

Typical structures like hyphae, arbuscules, and vesicles were observed in all tree species (Additional file 1: Table S3 and S4). This study was in agreement with Allen et al. (1998) and found all plant species to be colonized by AMF. There were a significant difference

Table 6 Root colonization (%) of AMF structures at four age groups of enclosure

Age (year)	HC	MHC	AC	VC
< 10	57.0 ± 0.9d	39.9 ± 1.0d	23.7 ± 0.9d	10.3 ± 0.9d
10–20	65.8 ± 0.7c	55.5 ± 0.8c	37.3 ± 0.7c	23.3 ± 0.7c
20–30	74.0 ± 0.7b	68.9 ± 0.8b	51.7 ± 0.7b	39.7 ± 0.7b
> 30	82.7 ± 0.6a	78.4 ± 0.7a	60.7 ± 0.6a	50.7 ± 0.6a

Means in the same column followed by same letter do not differ significantly at $P < 0.05$, mean ± SEM

HC hyphal colonization, MHC mycorrhizal hyphal colonization, AC arbuscular colonization, VC vesicular colonization

Table 7 AMF spore density (100 g⁻¹ dry soil) and root colonization (%) of plant species and their families

AMF between plant species			AMF between plant families		
Plant species	Spore abundance	Root length colonization	Plant families	Spore abundance	Root colonization
<i>Abutilon longicuspe</i>	5173.6 ± 194.9bcde	81.7 ± 5.5abcdefg	Acanthaceae	12,463.0 ± 396.2a	80.0 ± 15.9abcdef
<i>Acacia decurrens</i>	2254.2 ± 619.4cdef	68.6 ± 3.1fghij	Aloaceae	312.4 ± 608.9e	57.5 ± 3.1ghi
<i>Acacia etbaica</i>	2768.3 ± 287.5bcdef	73.0 ± 1.4cdefghij	Anacardiaceae	5505.4 ± 586.7bcd	92.1 ± 3.0abc
<i>Acacia lahai</i>	3224.2 ± 194.9bcdef	83.3 ± 5.5abcdefg	Apocynaceae	6018.8 ± 422.5bc	83.9 ± 2.1abcde
<i>Acacia saligna</i>	3065.0 ± 406.6bcdef	72.2 ± 2.0cdefghij	Asclepiadaceae	2365.5 ± 189.3cde	85.5 ± 11.2abcd
<i>Acokanthera schimpri</i>	6868.2 ± 632.1ab	90.5 ± 3.1abcde	Berberidaceae	2023.0 ± 189.3de	80.0 ± 11.2abcdef
<i>Acacia seyal</i>	3893.2 ± 272.6bcdef	78.2 ± 1.3abcdefgh	Boraginaceae	12,982.0 ± 189.3a	100.0 ± 11.2a
<i>Aloe vera</i>	312.4 ± 607.3f	57.5 ± 3.01hij	Buddleiaceae	6031.0 ± 1097.7bc	86.3 ± 5.6abcd
<i>Berberis holstii</i>	2023.0 ± 2181.6def	80.0 ± 10.9abcdefg	Cactaceae	173.7 ± 677.5e	56.9 ± 3.4hi
<i>Becium grandiflorum</i>	2344.2 ± 252.0cdef	65.8 ± 1.2fghij	Capparidaceae	2098.6 ± 896.2de	62.4 ± 4.5fghi
<i>Buddleja polystachya</i>	5144.0 ± 979.3bcde	81.2 ± 4.9abcdefg	Celastraceae	2849.4 ± 240.2cde	68.4 ± 1.2defghi
<i>Calpurinia aurea</i>	3318.8 ± 360.0bcdef	72.7 ± 1.8cdefghij	Combretaceae	5038.9 ± 548.8bcd	72.8 ± 2.8cdefghi
<i>Cadaba farinose</i>	2098.6 ± 894.0def	62.4 ± 4.5ghij	Cupressaceae	2331.1 ± 238.8cde	76.4 ± 1.2bcdefgh
<i>Carissa spinarum</i>	5339.3 ± 565.4bcde	78.6 ± 2.8abcdefgh	Ebenaceae	1769.8 ± 211.7de	62.2 ± 1.0fghi
<i>Clutia lanceolata</i>	2024.0 ± 3085.3def	52.0 ± 15.5j	Ericaceae	5330.5 ± 189.3bcd	73.50 ± 11.2bcdefghi
<i>Combretum aculeatum</i>	5038.9 ± 547.4bcde	72.8 ± 2.7cdefghij	Euphorbiaceae	4839.5 ± 457.7bcd	77.0 ± 2.3abcdefg
<i>Cordia africana</i>	12,982.0 ± 2181.6a	100.0 ± 10.9a	Fabaceae	3185.5 ± 148.6bcde	73.3 ± .7bcdefghi
<i>Croton macrostachyus</i>	5729.4 ± 710.5bcde	83.2 ± 3.5abcdefg	Flacourtiaceae	6738.6 ± 1792.5ab	96.3 ± 9.1a
<i>Cupressus lusitanica</i>	2433.6 ± 710.5cdef	86.7 ± 3.5abcdef	Lamiaceae	2380.7 ± 251.0cde	65.7 ± 1.2efghi
<i>Diospyros abyssinica</i>	4893.0 ± 2181.6bcde	98.0 ± 10.9a	Loganiaceae	5144.0 ± 981.8bcd	81.2 ± 5.0abcdef
<i>Diplostigma canescens</i>	2365.5 ± 2181.6cdef	85.5 ± 10.9abcdef	Loranthaceae	3533.0 ± 189.3bcde	85.5 ± 11.2abcd
<i>Dovyalis abyssinica</i>	6738.6 ± 1788.1ab	96.3 ± 9.0a	Malvaceae	5173.6 ± 1097.7bcd	81.7 ± 5.6abcdef
<i>Dodonaea angustifolia</i>	3857.7 ± 285.1bcdef	76.2 ± 1.4abcdefghi	Meliaceae	10,498.0 ± 1173.5a	80.8 ± 6.0abcdef
<i>Ekebergia capensis</i>	10,498.0 ± 170.5a	80.8 ± 5.9abcdefg	Moraceae	11,797.0 ± 189.3a	88.5 ± 11.2abcd
<i>Erica arborea</i>	5330.5 ± 2181.6bcde	73.5 ± 10.9cdefghij	Myrtaceae	2599.9 ± 275.5cde	69.2 ± 1.4defghi
<i>Euphorbia abyssinica</i>	4213.3 ± 596.0bcdef	72.6 ± 3.0cdefghij	Oleaceae	6890.6 ± 411.2ab	92.9 ± 2.1ab
<i>Eucalyptus camaldulensis</i>	2809.6 ± 319.4bcdef	69.6 ± 1.6efghij	Oliniaceae	3494.5 ± 861.1bcde	70.9 ± 4.4defghi
<i>Eucalyptus globules</i>	2002.6 ± 539.16def	68.1 ± 2.7fghij	Polygonaceae	264.0 ± 694.2e	55.1 ± 3.5i
<i>Euclea racemosa</i>	1769.8 ± 211.2ef	62.2 ± 1.0ghij	Proteaceae	2789.0 ± 2189.3cde	87.0 ± 11.2abcd
<i>Ficus vasta</i>	11,797.0 ± 2181.6a	88.5 ± 10.9abcdef	Rosaceae	7163.5 ± 2189.3ab	100.0 ± 11.2a
<i>Gravillea robusta</i>	2789.0 ± 2181.6bcdef	87.0 ± 10.9abcdef	Rubiaceae	3013.0 ± 2189.3bcde	76.5 ± 11.2bcdefgh
<i>Juniperus procera</i>	2318.1 ± 252.8cdef	75.0 ± 1.2bcdefghij	Sapindaceae	3857.7 ± 285.8bcde	76.2 ± 1.4bcdefgh
<i>Justicia schimperiana</i>	12,463.0 ± 3085.3a	80.0 ± 15.5abcdefg			
<i>Maytenus arbutifolia</i>	2918.5 ± 250.3bcdef	68.6 ± 1.2fghij			
<i>Maytenus senegalensis</i>	2094.7 ± 827.7def	65.8 ± 4.1fghij			
<i>Nuxia congesta</i>	6031.0 ± 194.9bcd	86.3 ± 5.5abcdef			
<i>Olea europaea</i>	6890.6 ± 410.2ab	92.9 ± 2.0abc			
<i>Olinia rochetiana</i>	3494.5 ± 858.9bcdef	70.9 ± 4.3defghij			
<i>Opuntia ficus indica</i>	173.7 ± 675.8f	56.9 ± 3.4ij			
<i>Osyris quadripartite</i>	3533.0 ± 2181.6bcdef	85.5 ± 10.9abcdef			
<i>Otostegia integrifolia</i>	2213.2 ± 185.0def	58.0 ± 7.0hij			
<i>Psudrax schimperiana</i>	3013.0 ± 2181.6bcdef	76.50 ± 10.9abcdefghi			

Table 7 AMF spore density (100 g⁻¹ dry soil) and root colonization (%) of plant species and their families (Continued)

<i>Rhus glutinosa</i>	5137.0 ± 692.5bcde	91.1 ± 3.5abcd
<i>Rhus retinorrhoea</i>	6426.5 ± 194.9abc	94.8 ± 5.56ab
<i>Rosa abyssinica</i>	7163.5 ± 2181.6ab	100.0 ± 10.9a
<i>Rumex nervosus</i>	264.4 ± 751.1f	54.9 ± 3.7j
<i>Senna singueana</i>	2307.7 ± 607.3cdef	56.5 ± 3.0ij

Means in the same column followed by same letter do not differ significantly at $P < 0.05$ for spore abundance and root colonization of plants in highlands of Tigray. Mean ± SEM

($P = 0.000$) in root length colonization between plant species and their families (Table 7). *Cordia africana*, *Diospyros abyssinica*, *Dovyalis abyssinica*, and *Rosa abyssinica* showed significant difference while *Clutia lanceolata* and *Rumex netvosus* showed the lowest root length colonization. Boraginaceae, Flacourtiaceae, and Rosaceae were the plant families with higher root length colonization, and Polygonaceae recorded the least colonization and was similar with the results of Chanie and Assefa (2013). Root length colonization by AMF for six Acacia species were found from 56 to 80% (Santhaguru and Sadhana 2000). The variation in the extent of root colonization could be related to the different preferences of the AMF to various species (Khade and Rodrigues 2009; David 2008) and mycorrhizal dependencies among plant species. The variations in plant species with relation to the degree of AMF colonization indicated that environmental factors influence the presence or absence of mycorrhizae and their colonization level (Alexander 1989) and such variation occur at genus and family levels (John 1980).

Acokanthera schimpri, *Cordia africana*, *Diospyros abyssinica*, *Dovyalis abyssinica*, *Ekebergia capensis*, *Olea europaea*, *Rhus glutinosa*, *Rhus retinorrhoea*, and *Rosa abyssinica* plant species had higher hyphal root colonization (Additional file 1: Table S3). Mycorrhizal hyphal colonization were higher in *Cordia africana*, *Diospyros abyssinica*, *Dovyalis abyssinica*, *Olea europaea*, *Rhus glutinosa*, *Rhus retinorrhoea*, and *Rosa abyssinica* plant species. Mycorrhizal hyphal colonization varied from high to low between different tree species (Chanie and Assefa 2013).

The four AMF structures were identified in all the 32 plants from all the study areas. Anacardiaceae, Boraginaceae, Flacourtiaceae, Meliaceae, Oleaceae, and Rosaceae showed higher hyphal colonization (Additional file 1: Table S4). Similarly, Anacardiaceae, Boraginaceae, Flacourtiaceae,

Oleaceae, and Rosaceae showed high mycorrhizal root colonization while high % AC was recorded in Boraginaceae. High % VC was observed in Rosaceae and Meliaceae. Recently, Burni and Hussain (2011) also reported variations in AMF root colonization in various members of Lamiaceae. Different edapho-climatic factors like soil type, nutritional status of soil, soil pH, organic matter, soil moisture, rain fall, temperature, etc. may be responsible for variations in root colonization (Tanvir et al. 2011).

AMF spore size

High spore density was recovered for the spore groups with 50 µm diameter followed by 100 and 300 µm (Table 8). *Acaulospora* had higher spore density in 50 and 100 µm diameter followed by *Glomus* while it was the most dominant in the 300 µm diameter group. *Gigaspora*, *Scutellospora*, and *Entrophospora* were the third, fourth, and fifth dominants genera in 50 and 100 µm, respectively. *Sclerocystis* was the least dominant found only in 100 and 300 µm. Brundrett et al. (1996) reported AMF produce large asexual spores (20–1000+ µm diameter). The fact that *Acaulospora* and *Glomus* are dominant genera in high lands of Tigray is related to their sporogenous characteristics. It has been found that *Acaulospora* and *Glomus* species usually produce more spores than *Gigaspora* and *Scutellospora* species in the same environment (Bever et al. 1996).

Soil nutrients availability with increasing age of exclosures and slope position

Soil chemical properties of exclosure soil were significantly higher than grazing land (Table 9). This result is supported by Mekuria (2010). Grazing impacts on soil properties depends on grazing intensity, with moderate grazing of 33 years compared with an ungrazed control resulted in higher pH, available P, and Mg (Ayorlo et al. 2011) and

Table 8 Spore density (100 g⁻¹ of dry soil) and types of genera at different spore size (µm) categories

Spore size (µm)	<i>Gigaspora</i>	<i>Scutellospora</i>	<i>Glomus</i>	<i>Sclerocystis</i>	<i>Acaulospora</i>	<i>Entrophospora</i>	Sum
50 µm	87.8 ± 2.1a	45.2 ± 1.2a	263.2 ± 5.3a	0.000 ± 0.003a	626.1 ± 11.9a	10.9 ± 0.4a	1784.2 ± 32.0a
100 µm	48.2 ± 2.1b	24.2 ± 1.2b	270.4 ± 5.3a	0.007 ± 0.003a	389.1 ± 11.9b	3.5 ± 0.4b	1079.1 ± 32.0b
300 µm	8.8 ± 2.1c	4.1 ± 1.2c	129.9 ± 5.3b	0.002 ± 0.003a	109.4 ± 11.9c	1.8 ± 0.4c	254.2 ± 32.0c

Means in the same column followed by same letter do not differ significantly at $P < 0.05$, mean ± SEM

Table 9 Soil properties in exclosures and adjacent grazing lands with increasing age of exclosures and slope position

		pH	EC (msm)	P (ppm)	K (ppm)	N (%)	OC (%)	OM (%)
Land uses	Exclosure	7.7 ± 0.03a	15.8 ± 0.38a	9.1 ± 0.45a	8.8 ± 0.2a	0.5 ± 0.01a	1.7 ± 0.05a	2.9 ± 0.09a
	Grazing land	7.2 ± 0.03b	9.4 ± 0.38b	2.9 ± 0.45b	3.7 ± 0.29b	0.2 ± 0.01b	0.3 ± 0.05b	0.5 ± 0.09b
Age groups	< 10	7.2 ± 0.05c	10.4 ± 0.64b	2.2 ± 0.67c	3.85 ± 0.49c	0.3 ± 0.02b	0.4 ± 0.11c	0.7 ± 0.19c
	10–20	7.4 ± 0.05b	11.9 ± 0.64b	5.7 ± 0.67b	6.1 ± 0.49b	0.4 ± 0.02b	0.95 ± 0.11b	1.6 ± 0.19b
	20–30	7.6 ± 0.05ab	12.4 ± 0.64b	6.16 ± 0.67b	6.7 ± 0.49ab	0.4 ± 0.02ab	1.2 ± 0.11ab	2.1 ± 0.19ab
	> 30	7.7 ± 0.05a	15.8 ± 0.64a	9.9 ± 0.67a	8.3 ± 0.50a	0.5 ± 0.02a	1.4 ± 0.11a	2.5 ± 0.19a
Slope position	Foot	7.7 ± 0.04a	14.2 ± 0.58a	7.2 ± 0.66a	6.6 ± 0.47a	0.4 ± 0.02a	1.2 ± 0.10a	2.1 ± 0.18a
	Middle	7.5 ± 0.04b	12.4 ± 0.58ab	5.7 ± 0.66a	6.3 ± 0.4a	0.4 ± 0.02b	1.0 ± 0.10ab	1.7 ± 0.18ab
	Upper	7.3 ± 0.04b	11.2 ± 0.58c	5.1 ± 0.66a	5.7 ± 0.46a	0.3 ± 0.020b	0.8 ± 0.10b	1.4 ± 0.18b

Means in the same column followed by the same letter do not differ significantly at $P < 0.05$, mean ± SEM

available P, total N, Ca, Mg, and K decreased after 1.5 years of heavy grazing compared with an ungrazed control in a tropical pasture. Soil chemical properties at older age exclosure had better pH, EC, P, N, K, OC, and OM than the three young age groups of exclosures. Availability of soil nutrients decreased with decreasing age of exclosure. Foot slope had better soil chemical properties than the middle and upper slope (Table 9). EC showed significantly higher in foot slope followed by middle and upper slope respectively which is in agreement with Mekuria (2010).

Correlation between AMF and availability of soil nutrients

Spore density had positively correlated with the soil parameters (Table 10) which is in agreement with the studies of Minal and Anil (2012) found a significant positive correlation with pH ($r = 0.85$, $P < 0.01$) and organic carbon ($r = 0.68$, $P < 0.05$). Similarly, Arpita et al. (2012) found root colonization were positively correlated with % OC, pH, available nitrogen, and available phosphorus but negatively correlated with available potassium. Spore densities were positively correlated with all soil parameters except soil pH. Don-Rodrigue et al. (2013) and Amal et al. (2013) also found positive correlation of soil properties with AMF.

Table 10 Pearson correlation between soil parameters and AMF spore density (100 g⁻¹ of dry soil) and colonization (%)

AMF parameters	Soil parameters					
	EC (msm)	P (ppm)	K (ppm)	N (%)	OC (%)	OM (%)
Spore density	0.888*	0.822*	0.897*	0.860*	0.904*	0.904*
HC	0.630*	0.638*	0.720*	0.720*	0.737*	0.737*
MHC	0.678*	0.708*	0.758*	0.731*	0.784*	0.784*
AC	0.599*	0.653*	0.673*	0.647*	0.708*	0.708*
VC	0.628*	0.692*	0.685*	0.645*	0.716*	0.716*

HC hyphal colonization, MHC mycorrhizal hyphal colonization, AC arbuscular colonization, VC vesicular colonization

*Correlation is significant at the 0.05 level

Conclusions

AMF spore density and root colonization were higher in exclosures as compared to grazing land and increased with age of exclosures. Foot slope had greater spore density and root colonization than mid and upper slopes. Soil chemical properties showed significant difference and were highest in the exclosure, oldest age of exclosure and foot slope followed by grazing land, youngest age of exclosure and upper slope. AMF spore density and root colonization were significantly positively correlated to soil chemical properties.

Additional file

Additional file 1: Table S1. AMF spore density (100 g⁻¹ dry soil) and genera by plant species (Mean ± SEM). **Table S2.** AMF spore density (100 g⁻¹ dry soil) and genera by plant families. (Mean ± SEM). **Table S3.** AMF root colonization (%) of different plant species. HC: hyphal colonization, MHC: mycorrhizal hyphal colonization, AC: arbuscular colonization, VC: vesicular colonization. Mean ± SEM. **Table S4** AMF root colonization (%) of plant families in highlands of Tigray. HC: hyphal colonization, MHC: mycorrhizal hyphal colonization, AC: arbuscular colonization, VC: vesicular colonization. Mean ± SEM. (DOCX 38 kb)

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Authors' contributions

EB and KMG generated the idea, designed the study, carried out the experiment works, conducted the sample collection and laboratory analysis, collected and analyzed the data, and wrote the manuscript. TD participated in its design and coordination and helped to draft the manuscript. MH and NS provided the statistical assistance and revised the manuscript draft. All authors read and approved the final manuscript.

Competing interests

The authors declare that they have no competing interests.

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