

Soil Fauna Communities and Soil Attributes in the Agroforests of Paraty

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ABSTRACT

We conducted the present study in Paraty, southeastern Brazil, in areas under different management regimes and plant cover. The study comprised two different agroforestry systems (AFS-1 and AFS-2), a secondary forest, and a cassava monoculture. We aimed at assessing the effects of land use on the soil fauna and its relationship with soil chemical (pH, Al, Ca, Mg, P, K, carbon, and organic matter) and microbiological attributes (soil microbial biomass carbon - SMB-C, soil respiration - SR, metabolic quotient - qCO_2 , microbial biomass carbon - C-mic). During winter, AFS-2 showed higher abundance of microphagous, saprophagous, and total individuals than the other areas. AFS-1 and the forest showed an increased abundance of Formicidae and phytophagous groups from winter to summer. The soil fauna and community structure showed that the studied agroforests are under regeneration, becoming more similar to the native forest, where ecological processes are considered efficient.

Keywords: soil organisms, organic matter, nutrient cycling.

Comunidades da Fauna do Solo e Atributos Edáficos sob Agroflorestas em Paraty, RJ

RESUMO

Este estudo foi conduzido no município de Paraty, RJ, em áreas com diferentes manejos e plantas de cobertura, sendo: dois diferentes sistemas agroflorestais (AFS-1 e AFS-2), uma área de floresta secundária e uma área com monocultivo de mandioca. O objetivo foi avaliar os efeitos de diferentes usos da terra na fauna do solo e sua relação com os atributos químicos (pH, Al, Ca, Mg, P, K, carbono e matéria orgânica) e microbiológicos (carbono da biomassa microbiana - C-BMS; respiração do solo - RS; quociente metabólico - qCO_2 ; quociente microbiano - Cmic) do solo. No inverno, o AFS-2 apresentou maior abundância de indivíduos micrófagos, saprófagos e totais, em relação a outras áreas. Do inverno para o verão, o AFS-1 e a área de floresta mostraram um aumento na abundância de formicidae e grupos fitófagos. A avaliação da fauna edáfica e a estrutura da comunidade mostraram que os AFSs estão sob processo de regeneração, aumentando a semelhança com florestas nativas, nas quais os processos ecológicos são considerados eficientes.

Palavras-chave: organismos do solo, matéria orgânica, ciclagem de nutrientes.

1. INTRODUCTION

Agroforestry systems are known as an intercropping practice involving tree species, which improves soil physical and chemical properties by decreased erosion and increased organic matter content in the topsoil (Mendonça et al., 2001). This practice is meant for the recovery of degraded areas and to preserve the surrounding of protected areas. Particularly in the Atlantic forest, agroforestry systems have already been recognized as an alternative to buffer zones around reserves (Cullen et al., 2001).

Agroforestry systems improve soil fertility and the microclimate. They also create more habitats, and so provide greater diversity of food sources than conventional monocultures (Tscharrntke et al., 2011). This increase in soil quality results from a different input of organic material from tree and crop species. A better soil quality stimulates the reestablishment of soil invertebrate community, which results in decomposition, nutrient cycling, and soil physical structure more similar to forests (Lima et al., 2010; Silva et al., 2012a; Tarrá et al., 2012).

The soil fauna is composed of invertebrates at different proportion and with different functions. These invertebrates interact with each other and with the microbial community, thus regulating decomposition (Lavelle et al., 2006; Resende et al., 2013). The soil fauna acts mainly in the fragmentation of plant debris, which makes them available to microorganisms. Microorganisms have a more suitable enzyme complement to break the litter fall, and, therefore, play an active role in decomposition (Wardle & Lavelle, 1997; Correia & Oliveira, 2000; Resende et al., 2013). According to Correia & Oliveira (2000), the effects of these microorganisms on the microbiota can be either stimulatory or inhibitory (Theenhaus & Scheu, 1996) and are reflected in the soil microbiological attributes (soil microbial biomass carbon - SMB-C, soil respiration - SR, metabolic quotient - qCO_2 , etc.).

Soil invertebrates are sensitive to changes in plant cover (Barros et al., 2003), management regime (Aquino et al., 2008; Silva et al., 2012a), and microclimate (Vasconcelos et al., 2009). In the conversion of natural ecosystems into productive systems, the structure of the soil fauna is less altered when the structure of the plant cover is kept similar, i.e., when fields are grown

in pastures, and silvopastoral and agrosilvicultural systems are grown in areas originally covered with forests (Lima et al., 2010).

Lima et al. (2010) observed that the implementation of agroforestry systems in the Cerrado (Brazilian savanna) increases the richness and density of the edaphic macrofauna compared to native forests (Lima et al., 2010). However, Silva et al. (2012a) did not observe significant differences in those variables between agroforestry systems and the native forest of Zona da Mata in the state of Minas Gerais, southeastern Brazil. According to some authors (Moço et al., 2009; Lima et al., 2010; Silva et al., 2013), plant diversity in agroforestry and forestry systems improves the diversity of microhabitats, which contributes to higher biodiversity.

Thus, agroforestry systems are models for good soil management, which most resemble a natural forest. Therefore, they are considered an alternative for a sustainable ecosystems use. The present study aimed to assess the effect of two agroforestry management regimes on soil fauna abundance and diversity, as well as their relationship with soil chemical and microbiological attributes.

2. MATERIAL AND METHODS

We carried out the present study in Campinho da Independência, a maroon community (locally known as *quilombo*) located in Paraty-Mirim, Paraty, state of Rio de Janeiro, southeastern Brazil (44°42'W, 23°17'S, 60 m a.s.l.). The community is located at the center of Cairucu, a protected area, by the Paraty-Mirim River and the BR101 Highway. The area still has native vegetation: submontane tropical rainforest (Brasil, 1983).

The studied area has a clay soil covered with secondary forests (SF), cassava monocultures, and two different agroforestry systems (AFS-1 and AFS-2). The distance between AFS-1 and AFS-2 and the cassava monoculture is approximately 3 m, and the distance thereof to the SF is approximately 15 m. The SF has approximately 1 ha, and most of its species occur at low density. According to Tavares (2014), it could still be at an early stage of succession, despite being located in a landscape characterized by large forest fragments.

The cassava monoculture is under traditional management for food production (maize, beans, yam,

plantain, and cassava), based on intercropping, crop rotation, and fallow. The monoculture had approximately 600 m² of cultivated area near a forest without tillage or fertilization, similar to the AFSs.

Agroforestry systems were established in 2003 based on the Agroforestry Systems Regenerative and Analogues (ASRA; Piña-Rodrigues, 2006) model. Each experimental unit consisted of a 20 x 20 m plot, corresponding to treatments of mixed legume (AFS-1) and non-legume species (AFS-2). The main culture of economic interest in both AFSs was peach palm (*Bactris gasipaes* Kunth.). They also had short-cycle fruit species such as banana (*Musa* sp.) and papaya (*Carica papaya* L.), subsistence food species such as black beans (*Phaseolus vulgaris* L.), corn (*Zea mays* L.), and cassava (*Manihot esculenta* L.), and green manure species such as jack bean (*Canavalia ensiformis* (L.) DC.), pigeon pea (*Cajanus cajan* (L.) Millsp.), and sunnhemp (*Crotalaria juncea* L.).

AFS-1 and AFS-2 differed in the composition of long-cycle fruit species. Abiu (*Pouteria caimito* Radlk.) and jackfruit (*Artocarpus heterophyllus* Lam) were cultivated in AFS-1 and graviola (*Anona muricata* L.) and starfruit (*Averrhoa carambola* L.) in AFS-2. They also differed in the composition of fertility renewing species, as *Inga* sp and *Lonchocarpus guilleminianus* (Tul.) Malme were present in AFS-1, and *Bixa orellana* L. and *Cytharexylum myrianthum* Cham in AFS-2. The composition of climax species and species with

potential use of seeds, oils, and hardwood also differed between AFSs. *Schizolobium parahyba* (Vell.) Blake, *Centrolobium tomentosum* Guill. ex Benth., *Hymenaea courbaril* L., and *Copaifera langsdorffii* Desf. were present in AFS-1, whereas, *Cedrela fissilis* Vell. (*Nectandra lanceolata* Nees et Mart. ex Ness., *Cariniana legalis* (Mart.) Kuntze and *Virola bicuhyba* (Schott) Warb were present in AFS-2.

The spacing between rows was 2 m, alternating planting pupunha (*Bactris gasipaes* Kunth) with another arborous species, or arborous species intercropped with fruit species. Spacing within the row was 1 m for the pupunha and 1.5 m for arborous and fruiting species. A total of 1561 seedlings were planted, including 462 seedlings of pupunha per treatment. Agricultural and green manure species were planted in the interrows of arborous species.

We collected soil samples 18 (winter) and 24 (summer) months after the AFS implementation. In AFSs and cassava areas, we used the total area for soil collection, whereas, in the SF area, we used a plot of 400 m². We collected five soil samples at 0-10 cm depth to measure chemical (EMBRAPA, 1997) and microbiological attributes (Soil microbial biomass carbon – SMB-C (Vance et al., 1987; Bartlett & Ross 1988); soil basal respiration – SR (Jenkinson & Powelson, 1976); and microbial biomass carbon – C-mic and metabolic quotient- qCO_2 (Anderson & Domsch, 1985, 1990) (Table 1). We also collected five random samples

Table 1. Chemical and microbiological characterization* of the soil layer at 0-10 cm under agroforestry systems (AFS-1 and AFS-2), cassava monoculture (CAS), and secondary forest (SF) during the winter (W) and summer (S). Se= season; C: total organic carbon; OM: organic matter.

		AFS 1		AFS 2		CAS		SF	
		W	S	W	S	W	S	W	S
C	(g kg ⁻¹)	26.5	26.9	20.5	24.1	25.5	30.6	29.1	44.0
OM	(g kg ⁻¹)	45.84	46.54	35.46	41.69	44.11	52.94	50.34	76.12
N	(g kg ⁻¹)	3.3	3.3	2.4	2.7	3.2	3.7	3.8	5.9
pH		5.6	5.0	5.4	5.1	5.2	4.8	4.9	4.5
Al	(cmol _c dm ⁻³)	0.07	0.48	0.33	0.44	1.03	0.96	1.60	1.02
Ca	(cmol _c dm ⁻³)	3.70	2.70	1.80	2.24	1.70	1.76	1.70	3.24
Mg	(cmol _c dm ⁻³)	1.53	1.04	1.30	1.14	1.43	1.04	1.17	1.72
P	(mg dm ⁻³)	11.0	7.0	6.3	7.0	7.6	11.0	8.3	15.0
K	(mg dm ⁻³)	267	51	73	43	88	152	110	91
SMB-C	(μg g ⁻¹)	282	254	307	241	233	160	357	472
SR	(μg C-CO ₂ g ⁻¹ h ⁻¹)	0.96	1.32	1.01	1.23	0.47	0.86	1.15	3.95
qCO_2	(mg C-BMS h ⁻¹)	3.4	5.2	3.3	5.1	2.0	5.4	3.2	8.4
C-mic	(%)	1.06	0.94	1.50	1.01	0.91	0.52	1.23	1.10

*Adapted from Silva et al. (2012b).

per area per season to assess soil fauna using a 25-cm wide square-shaped metal template. We divided each sample into litter fall and soil (0-5 cm of depth). For the extraction of soil fauna, we used a Berlese-Tullgren extraction funnel for 15 days. After that period, we collected organisms for counting and identification of taxonomic groups using binocular lens.

We estimated the abundance of edaphic fauna by transforming the number of individuals found in each sample into the number of individuals per square meter (individuals.m⁻²). We made a qualitative analysis of the soil fauna community by classifying taxonomic groups into nine functional groups according to habitat use and food resources (Seastedt, 1984): microphagous (Acari and Collembola), Formicidae (ants), saprophagous (Blattodea, Diplopoda, Diplura, Enchytraeidae, Gastropoda, Isopoda, Orthoptera, Pauropoda, Psocoptera, Symphyla, and Thysanoptera), phytophagous (Heteroptera and Homoptera), holometabolous (Coleoptera, Diptera, Lepidoptera, and Trichoptera), larvae (Coleoptera, Diptera, Lepidoptera, and Trichoptera), predators (Araneae and Chilopoda), parasitoids (Hymenoptera), and termites (Isoptera).

To describe the structure of the soil fauna community in each area we calculated the Shannon diversity index (H') (Shannon & Weaver, 1964) and the Pielou evenness index (J) (Pielou, 1975). H' considers the species (S) richness and their relative abundance (pi), defined by $H' = -\sum pi \ln pi$. J considers differences in abundance distribution among species, defined as $J = H' / \ln S$.

The agroforestry systems sampled in the present study were implemented as demonstration units at Quilombo do Campinho and in other communities of Paraty. Therefore, there was no experimental design with randomly distributed blocks, which impairs the analysis of variance and the comparison of means using the Bonferroni t test mandatory (Bailey, 1977). We used the software CANOCO to make a principal component analysis (PCA) of the edaphic fauna and soil chemical (pH, Al, Ca, Mg, K, P, N, C, organic matter) and microbiological variables (soil microbial biomass carbon - SMB-C, soil respiration - SR, metabolic quotient - qCO_2 and microbial biomass carbon - C-mic) (Table 1). We calculated the statistical significance of correlations between soil microbiological and chemical variables and fauna variables using a Monte Carlo permutation test (Ter Braak, 1995).

3. RESULTS AND DISCUSSION

We collected a total of 8,206 individuals, 41% during the winter and 59% during the summer. During the winter, AFS-2 had the highest abundance of individuals (4320 ind.m⁻²), followed by the secondary forest (3094 ind.m⁻²), AFS-1 (1878 ind.m⁻²), and the cassava monoculture (1450 ind.m⁻²). AFS-2 also had the highest abundance of individuals in the litter fall (2864 ind.m⁻²) (Figure 1a), which represented 33% of the total abundance, followed by the forest, with 30%, and the AFS-1, with 10%. During the winter, the abundance of individuals in AFS-2 and SF was statistically higher

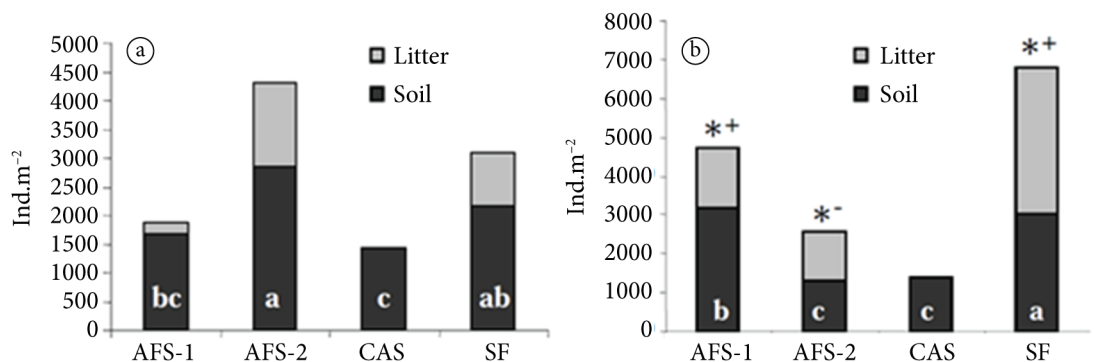


Figure 1. Abundance of fauna (individuals per m²) in the litter fall and soil under agroforestry systems (AFS-1 and AFS-2), cassava monoculture (CAS), and secondary forest (SF) during the winter (a) and summer (b). Same letters represent statistically similar areas in the same season (Total = soil + litter fall), whereas, an asterisk represents a significant difference in the mean of each treatment (Total = soil + litterfall) in relation to the previous season; (*+) means an increase and (*-) means a decrease.

than in the cassava monoculture, whereas during the summer AFS-1 and SF showed higher abundance of individuals than the cassava. (Figure 1a).

This result shows that the edaphic fauna responds positively to agroforestry management. The management practices used in a productive system can affect directly and indirectly the soil fauna, especially in terms of density and diversity (Nunes et al., 2008). Silva et al. (2012b) found higher amount and better nutritional quality of litter fall in AFSs and SF areas than in cassava monocultures. This fact might explain the higher abundance of individuals in AFSs and SF found in the present study. Some studies show that management systems that promote the maintenance of plant residues on the soil provide a more favorable environment for colonization by most groups of edaphic fauna (Moço et al., 2005).

We observed an increase in the abundance of individuals per square meter in the AFS-1 and forest (Figures 1a and 1b) from winter to summer. Silva et al. (2009) also observed an increase in the abundance of soil fauna from winter to summer in a secondary forest in Ubatuba, state of São Paulo, southeastern Brazil. These authors attributed that pattern to high moisture content in the soil, commonly observed during the summer (rainy season).

We found 24 taxonomic groups, which we divided into nine functional groups (Table 2). Microphagous were the most abundant group in the samples, with 4,277 individuals, which represented 52.12% of the total individuals collected in the study. Ants were the second

most abundant functional group, with 30.55% of the total individuals, followed by saprophagous (5.53%).

During the winter, ants showed higher average abundance in the forest than in all other areas. From winter to summer, there was an increase in the abundance of this group in AFS-1, and also in the forest, which showed a higher abundance of ants compared to AFS-2 and cassava monoculture, but similar to AFS-1 during the summer. Phytophagous also showed an increased abundance in AFS-1 and forest areas during the summer. In the AFS-1 the abundance was higher than in the AFS-2 and cassava monoculture, but not significantly different from the forest (Table 2). This result showed that the increase in abundance of individuals found in those areas during the study corresponded to the same functional groups. This fact corroborates the hypothesis of an intense biological flow between the forest and the AFS-1, which is a symptom of the edge effect (Didhan & Lawton, 1999; Copatti & Gasparetto, 2012). In a study conducted in southern Brazil, there was no difference in functional groups of soil fauna between agroforestry and native forest systems (Albuquerque et al., 2009).

The SF showed the highest abundance of holometabolous species during the winter compared to the AFS-1 and cassava monoculture. From winter to summer, there was a significant increase in microphagous and parasitoids in the SF, in addition to the ants and phytophagous previously mentioned. This pattern may result from a higher moisture content in the soil during summer (rainy season) (Nunes et al., 2012;

Table 2. Abundance of soil fauna (individuals per m²) classified by functional group under agroforestry systems (AFS-1 and AFS-2), cassava monoculture (CAS), and secondary forest (FS) in two seasons (summer – S; winter – W).

Areas	Se	Micro	Ants	Sapro	Phyto	Holo	Larvae	Preda	Para	Termites
AFS-1	W	1398bA	294bB	99bA	22abB	51bA	13aA	0aB	0aA	0aA
	S	1661bA	1651abA	198aA	570aA	499aA	109aA	16aA	10abA	3aA
AFS-2	W	3222aA	621bA	253aA	61aA	112abA	38aA	6aA	3aA	3aA
	S	1283bB	797bA	221aA	144bA	77aB	42aA	10aA	3abA	3aA
CAS	W	1066bA	202bA	86bA	6bA	38bA	48aA	0aB	0aA	3aA
	S	746bA	435bA	90aA	51bA	58aA	16aA	10aA	0bA	0aA
SF	W	1485bB	1174aB	77bA	26abB	195aA	125aA	10aA	3aB	0aA
	S	2826aA	2848aA	426aA	198abA	368aA	99aA	32aA	19aA	0aA
Total		13686	8022	1450	1078	1398	490	83	38	12

Same lowercase letters in the same column represent significantly similar areas in the same season, whereas same uppercase letters in the same column represent significantly similar seasons in the same treatment, according to the t test with Bonferroni correction ($p < 0.05$). Se: seasons, W: winter, S: summer, Micro: microphagous, Sapro: saprophagous, Phyto: Phytophagous, Holo: holometabolous, Preda: predators, and Para: parasitoids.

Fernandes et al., 2011), and higher organic matter content (Silva et al., 2013; Tripathi & Sharma, 2006) as shown in Table 1. During the summer, saprophagous showed no significant differences between seasons or areas (Table 2), whereas during the winter, a higher abundance of this group was found in the AFS-2. That result may indicate an increased litter fall decomposition rate during the winter in AFS-2 compared to other areas. Saprophagous feed directly on plant litter (Moço et al., 2005), fragmenting them and contributing to litter fall decomposition, organic matter redistribution, and nutrient cycling in the environment (Correia & Andrade, 1999).

AFSs showed the lowest Shannon diversity (H') and evenness (J) during the winter, and only the J of AFS-2 was significantly lower than that of cassava monoculture (Table 3). From winter to summer, AFS 1 and 2 showed an increase of approximately 60% in H' , associated with an increase of approximately 25% in richness in AFS-1, and an increase of 56% in J in AFS-2. The cassava monoculture showed high H' and J in both seasons, but its taxonomic group richness was lower than that of AFS-2 during the winter and all other areas during the summer. The forest and the cassava monoculture showed no significant increase in H' , due to different reasons. In the forest, there was a high increase in richness but not in evenness, which remained unchanged. The cassava monoculture showed a small increase in richness, which did not surpass 15 groups during the summer (Table 3), suggesting a potential for diversity increase in AFSs, both in richness and evenness components.

Motta (2005) found similar results in Atlantic Forest fragments in Rio de Janeiro, with richness ranging from 15 to 22. However, these authors observed that J was below 0.31, resulting in an H' below 1, which shows a strong dominance in the community.

Figures 2a and 2b show the principal component analysis (PCA), where eigenvectors (arrows) represent the taxonomic groups and points represent the samples of each study area. During the winter, the areas were able to explain 30.6% of the fauna data variance (Figure 2a). Axis 1 explained 41.9% of data and axis 2 explained only 3%. Most eigenvectors pointed to the right on axis 1. The AFS-2 showed more samples on the right. This pattern indicates abundant and diverse fauna, which matches the high richness and abundance of individuals per square meter of this area in the winter. The forest showed samples in the central part, with average status. Finally, the cassava monoculture and AFS-1 showed samples on the left; the latter showed also samples in the right quadrant.

According to the PCA, the areas explained 33.5% of the variance during the summer, and of these data, 46% are explained on axis 1, and 9.8% on axis 2. In that season, most eigenvectors pointed to the left, indicating the abundance of most soil fauna groups in this direction. The forest was the only area with all samples in the upper left quadrant, forming the largest polygon. The pattern repeated during the winter, which indicates high variability between samples. The AFS-1 showed an intermediate pattern, with three samples near the center and two in the lower left quadrant, indicating abundance of individuals, but in a composition different from that of the forest.

Table 3. Shannon diversity (H'), species richness, and Pielou evenness (J) calculated for the soil fauna under two agroforestry systems (AFS-1 and AFS-2), cassava monoculture (CAS), and secondary forest (SF) in two seasons (winter and summer).

Areas	Season	H'	Richness	J
AFS-1	Winter	1.42 aB	16 abB	0.51 abA
	Summer	2.30 aA	20 aA	0.77 aA
AFS-2	Winter	1.47 aB	21 aA	0.48 bB
	Summer	2.33 aA	22 aA	0.75 aA
CAS	Winter	1.58 aA	13 bA	0.62 aA
	Summer	1.97 aA	15 bA	0.73 aA
SF	Winter	1.82 aA	19 abB	0.62 abA
	Summer	1.93 aA	22 aA	0.62 aA

Same lowercase letters in the same column represent significantly similar areas in the same season, whereas, same uppercase letters in the same column represent significantly similar seasons in the same treatment, according to the t test with Bonferroni correction ($p < 0.05$).

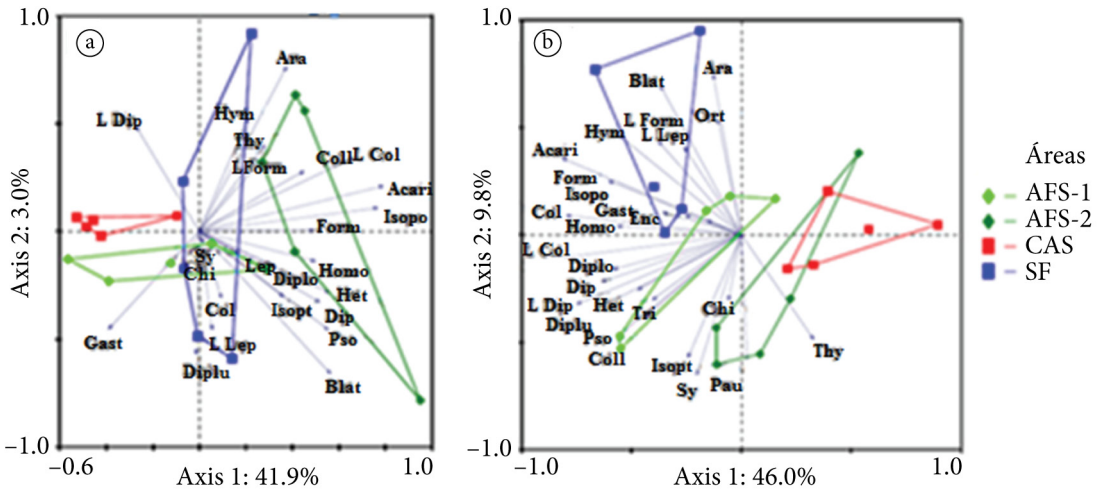


Figure 2. Principal components analysis of the abundance of taxonomic groups of soil fauna and study areas (agroforestry systems - AFS-1 and AFS-2, cassava monoculture - CAS and secondary forest -SF) during the winter (a) and summer (b). Formicidae - Form (ants), Blattodea - Blat, Diplopoda - Diplo, Diplura - Diplu, Enchytraeidae - Enc, Gastropoda - Gast, Isopoda - Isopo, Orthoptera - Ort, Pauropoda - Pau, Psocoptera - Pso, Symphyla - Sy, Thysanoptera - Thy, Heteroptera - Het, Homoptera - Homo, Coleoptera - Col, Diptera - Dip, Lepidoptera - Lep, Trichoptera - Tri, larvae (Coleoptera - L Col, Diptera - L Dip, Lepidoptera - L Lep and Trichoptera - L Tri), Araneae - Ara, Chilopoda - Chi, Hymenoptera - Hym, Isoptera - Isopt, Collembola - Coll.

That pattern was also evident when assessing the abundance of functional groups, showing that only some elements of the soil fauna behaved similarly in the AFS-1 and the forest. Considering edge effects (Copatti & Gasparetto, 2012), this pattern may result from group mobility, as in the case of ants or the resource availability in the AFS-1, which would be attracting organisms (e.g., phytophagous) from the forest. AFS-2 showed three samples in the right quadrant, approaching the cassava monoculture, which showed all samples in this quadrant. The decline in soil fauna status in AFS-2 during the summer is associated with a decrease in the abundance of individuals, especially microphagous. However, the AFS-2 maintained high richness of groups and diversity and evenness indexes during the summer.

The patterns found in the PCA can be seen in the principal response curves (PRC), which order the areas according to the fauna, on only one axis and allow comparison of the AFSs and the cassava monoculture with forests throughout the study period. During the winter, AFS-2 showed a higher discriminant coefficient (Cdt) (Figure 3) than the forest, and AFS-1 showed a slightly lower discriminant coefficient than the forest and considerably lower than the cassava monoculture.

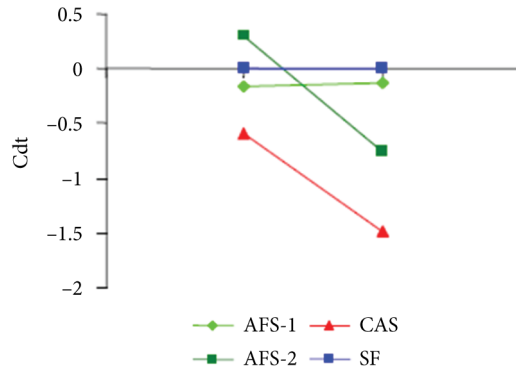


Figure 3. Principal response curves (PRC) of agroforestry systems (AFS-1 and AFS-2) and the cassava monoculture (CAS) according to soil fauna groups, compared to a secondary forest (SF). Cdt = discriminant coefficient.

During the summer, the discriminant coefficient of the AFS-2 and cassava monoculture decreased, and the forest was the area with the best status, followed by the AFS-1, AFS-2, and cassava monoculture.

The decline in the discriminant coefficient of the AFS-2 and cassava monoculture is partly due to the fact that the forest is the benchmark. The cassava monoculture was the most stable area, remaining low

and constant in both seasons in terms of diversity, evenness, and richness; so that its fall on the graph indicates only that the cassava monoculture did not follow the trend of the forest. The AFS-2 showed a decline in the abundance of individuals, but was able to maintain the group richness and increase the evenness index throughout the study period. This is the expected pattern for a biological community throughout ecological succession (Machado et al., 2015). The AFS-1 remained very close to the forest in both seasons on the principal response curves, indicating that variations in the composition of soil fauna groups in these two areas showed the same trend.

Figures 4a and 4b showed the direct redundancy analysis (DRA), which summarizes the situation of the areas according to the explanatory (chemical and microbiological) and response (edaphic fauna) variables. During the winter, the environmental variables (chemical and microbiological) were able to explain 34.4% of the total variance of soil fauna. Regarding the explained variance, axis 1 explained 15.2%, and axis 2 explained 13.4%, showing that the variance was distributed along both axes (Figure 4a).

Eigenvectors related to soil fauna groups pointed mainly to the right, where AFS-2 occurred under the

influence of the Acari group that was abundant in this area, and upwards, where forest occurred. We also noticed the influence of the group Formicidae. The AFS-1 showed an intermediate pattern, near the center. The cassava monoculture was on the extreme lower left, indicating few fauna individuals. The microbiological indicators were in the upper right quadrant, and in addition to fauna, formed a group of biological components in which the best positioned areas were the AFS-2 and forest, followed by the AFS-1 in an intermediate situation, and the cassava monoculture in the opposite quadrant (Figure 4a). The axis of biological components showed the response of the AFS-2 to the litter deposition provided by agroforestry management, in which the microbial activity was stimulated to obtain a higher abundance of microphagous, saprophagous, and total individuals per square meter during the winter.

The axis formed by fertility variables showed a situation of high organic matter and nitrogen in the upper left quadrant, where the forest occurred, and the lower right quadrant, where the AFS-2 occurred with high pH and C/N ratio. The AFS-1 was in an intermediate situation, as well as the cassava monoculture, which stood out in other nutrients. The order observed on the axis formed by fertility variables matched the spatial pattern of those areas in the field, supporting the hypothesis

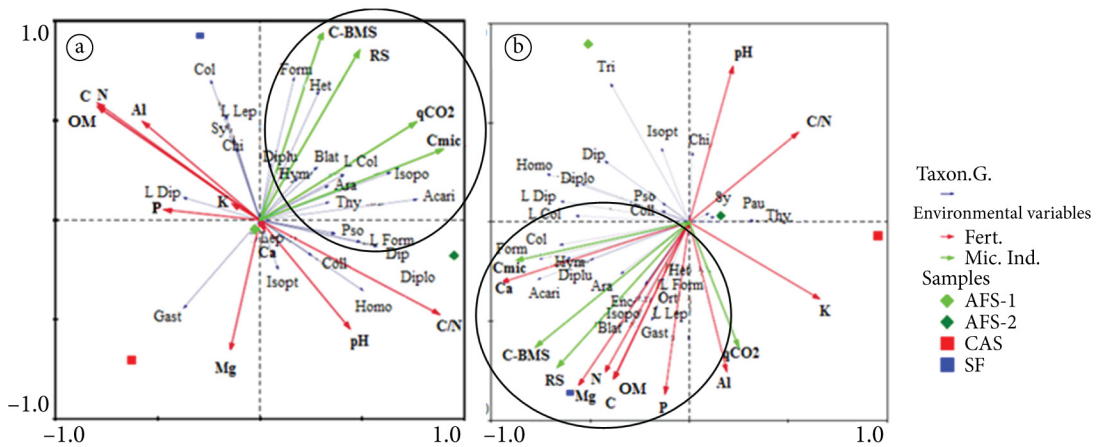


Figure 4. Direct redundancy analysis showing the canonical fauna ordering of soil samples from two agroforestry systems (AFS-1 and AFS-2), secondary forest (SF) and cassava monoculture (CAS) in relation to chemical and microbiological variables of the soil during the winter (a) and summer (b). Taxon. G. = taxonomic group, Fert = fertility, Mic ind = microbiological indicators. Formicidae - Form (ants), Blattodea - Blat, Diplopoda - Diplo, Diplura - Diplo, Enchytraeidae - Enc, Gastropoda - Gast, Isopoda - Isopo, Orthoptera - Ort, Pauropoda - Pau, Psocoptera - Pso, Symphyla - Sy, Thysanoptera - Thy, Heteroptera - Het, Homoptera - Homo, Coleoptera - Col, Diptera - Dip, Lepidoptera - Lep, Trichoptera - Tri, larvae (Coleoptera - L Col, Diptera - L Dip, Lepidoptera - L Lep and Trichoptera - L Tri), Araneae - Ara, Chilopoda - Chi, Hymenoptera - Hym, Isoptera - Isopt.

that the AFS-1 and cassava monoculture were under the influence of the adjacent forest (Figure 4a).

The axis formed by biological components is positioned almost orthogonally to the fertility axis, showing little correlation between them (Figure 4a). This was due to the situation of the AFS-2, which showed positive responses to the agroforestry management on biological components whereas the soil chemical composition (Table 1) was not significantly affected during the winter.

During the summer, the direct redundancy analysis showed that the environmental variables explained 30.3% of the variance of groups in the samples. Most eigenvectors of fauna groups and most fertility and microbiological variables were in the lower left quadrant, where forest samples (Figure 4b), among the fertility variables, calcium (Ca, $p = 0.004$), magnesium (Mg, $p = 0.052$, n.s.), and OM ($p = 0.408$) were selected, and among the microbiological variables, C-mic ($p = 0.002$), SR ($p = 0.042$), and SMB-C ($p = 0.190$, n.s.) were selected. Among all environmental variables, Ca ($p = 0.004$), Mg ($p = 0.052$, n.s.), and OM ($p = 0.408$, n.s.) were selected, and, thus, all the selected variables were positioned in the lower left quadrant.

The forest was the only area with samples in the lower left quadrant, corresponding to the highest fertility, microbial activity, and abundance of edaphic fauna groups. Regarding the fauna, the forest and AFS-1 were positioned closer to the left, where most eigenvectors pointed. This pattern corroborates the principal response curves and agrees with the increase in Homoptera and Formicidae groups. The AFS-2 appeared in the upper left quadrant, near the center, in an intermediate position in relation to the fauna, but similar to the AFS-1 in relation to soil fertility. The cassava monoculture is on the far left, in a position contrary to most fauna arrows, showing low biological activity (Figure 4b).

4. CONCLUSIONS

In general, the studied agroforestry systems have fauna abundance and diversity (richness and Shannon index) similar to those of the forest. The type of agroforestry system is important to determine community composition. Most fauna groups were positively related to microbiological attributes, which indicates that

those invertebrates are probably stimulating the soil microbiota. On the other hand, the correlation of the fauna with chemical attributes was low.

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