EDAPHIC MITES AND THEIR RESPONSE TO THE INCORPORATION OF ORGANIC MATTER FROM VARIOUS SPECIES OF FABACEAE INTO THE SOIL BENEATH COFFEE TREES

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ABSTRACT: The aim of this study was to evaluate the presence and abundance of mites in the soil beneath cultivated coffee trees (*Coffea arabica* L.). In particular, we compared mite communities in three different soil environments: 1) native forest; 2) soils from underneath the coffee trees "under full sun" (i.e. areas that received no additional shade or organic treatment); 3) soils that incorporated organic matter from four leguminous windbreak plant species — acacia, *Acacia mangium* Wild.; pigeon pea, *Cajanus cajan* (L.) Millsp.; gliricidia, *Gliricidia sepium* (Jacq.) Walp.; and leucaena, *Leucaena leucocephala* (Lam.) de Wit. Sampling was conducted at the ends of the dry and rainy seasons, in São Sebastião do Paraíso, Minas Gerais, Brazil. Undisturbed soil samples were acquired using a cylinder and a Berlese-Tullgren funnel was used to extract the mites in the laboratory. A total of 1,014 mite specimens, assigned into 143 different species, were collected. The forest soil had the greatest richness and total abundance of edaphic mites, while the coffee plantation soils, enriched with acacia, had the lowest richness and abundance. The mite communities of treated soils were less than 18% similar to that of the forest soil. These results suggest that the substitution of native forests with cultivated systems can cause significant changes in the abundance, richness and structuring of edaphic mite communities, particularly of oribatid mites. Maintenance of mite communities should be one of the goals of agricultural practices, since these organisms are important for maintenance of biological cycles, especially for the decomposition of organic matter.

KEY WORDS: Acarology, edaphic mites, mesofauna, Coffea arabica.

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INTRODUCTION

Efforts have been made with several agricultural crops to implement responsible crop cultivation practices. Namely, researchers have been trying to come up with management systems that are minimally invasive to the diverse fauna and flora of such areas (Perfecto *et al.* 1996; Moguel and Toledo 1999). The search for such sustainable cultivation models has been the focus of studies of coffee growing practices (Perfecto *et al.* 1996; Moguel and Toledo 1999).

Based on the concept of sustainable production, the research, focused on soil-plant interaction, has abandoned the idea that soil fertility is solely the result of its intrinsic chemical and physical properties (Oliveira *et al.* 2006). Thus, a fertile soil that is suitable for cultivation, should not only be rich in nutrients and free from physical processes that impede crop implementation—such soil should also be biologically active, so that nutrient cycling processes may be completed (Oliveira *et al.* 2006).

The biological community of soil consists of numerous organisms including plants, lichens, bacteria, fungi and a wide range of arthropods, which are responsible for several activities that directly and indirectly influence the chemical and physical properties of the soil (Pankhurst and Linch 1994, Theenhaus and Scheu 1996). In addition, these organisms may play a role in soil regeneration, thereby benefiting vegetation by making nutrients available and maintaining equilibrium among the populations of organisms present (Uhlig 2005).

Mites are an important component of the edaphic fauna, and they may represent as many as 95% of all microarthropods in the soil (Seastedt 1984). Edaphic mites are characterized by a vast diversity of species and feeding habits. They can be strictly predatory, or they can feed on microorganisms and decomposing matter; their feeding habits may also vary throughout their life cycle, e.g., parasitic immatures transforming to predacious adults. Among predatory mites, there are those that feed on nematodes, larvae and adults of other invertebrates, including mites (Evans and Campbell 2003; Castilho *et al.* 2009; Moreira *et al.* 2015). Given this variation in feeding habits,

some predatory species have been widely used in agriculture. Others, namely the species of the families Macrochelidae, Laelapidae and Rhodacaridae, may be used as biological control agents for the soil (Wallace *et al.* 1979; Evans and Campbell 2003; Castilho *et al.* 2009; Moreira *et al.* 2015).

Since mites live in soil, they may serve as indicators of soil quality (Rieff 2010). Therefore, the collection of certain edaphic organisms and the analysis of their population parameters (e.g., richness, abundance, exclusive species and quality indicator species) permits the evaluation of soil quality in cultivated areas and natural environments (Ruf 1998; Ruf and Beck 2005; Bedano and Ruf 2010). Such information can contribute to the maintenance, recovery and restoration of the environmental health, and thus, to the sustainability of ecosystems (Ruf 1998; Ruf and Beck 2005; Bedano and Ruf 2010).

Although the greatest richness of edaphic mites is found in native forests, several species of this group can be found in other systems, including anthropic environs. However, even in situations characterized by lower abundances and fewer species, mites may perform important biological functions, such as pest control and nutrient cycling (Crossley et al. 1992; Hülsmann and Wolters 1998; Bedano et al. 2006; Morais et al. 2010). The same occurs in coffee cultivation areas, where the substitution of native vegetation tends to negatively affect soil acarofauna, with the reduction in the richness and abundance of soil species (Rojas et al. 2009; Marafeli et al. 2018). However, coffee cultivation with a shading system may reduce the loss of biodiversity. This type of management, which maintains part of the native vegetation integrated into the cultivation system, allows the maintenance of a greater species richness than conventional types of agricultural practices (Perfecto et al. 1996; Moguel and Toledo 1999). Thus, one of the issues that must be considered is how to reconcile agricultural production and expansion with practices that are less detrimental to biodiversity. The objective of the present study was to evaluate the occurrence and variation in the edaphic mite fauna that occurs in the soil beneath coffee trees with and without the incorporation of organic matter into the soil. The present study compared three environments: coffee crops grown in the shade produced by rows of arboreal Fabaceae; coffee grown in full sun; and native forest.

MATERIAL AND METHODS

Mite community sampling

The present study was conducted at the EP-AMIG Sul Experimental Farm in São Sebastião do Paraíso, in the state of Minas Gerais, Brazil (20°54'37.32" S; 47°06'43.8" W). The site is 837 m a.s.l. The region has a mean annual temperature of 20.8 °C and a mean annual rainfall of approximately 1,400 mm. The rains mainly occur between October and April. The soil of the region is the Dystroferric Red Latosol.

The experiment was conducted in an open area of a conventional coffee plantation without any control of the environmental factors. As a reference farm for research activities, numerous experiments regarding coffee management are conducted in the area. These research activities are conducted with the aim of better understanding the dynamics of coffee growing areas. Nonetheless, during the present experiment, no other activities were conducted that could have interfered with the results.

At the site, there was only Topázio (Coffea arabica L.), which was planted in December 1999. The coffee plant rows were spaced 3.5 m apart, with 0.5 m between plants in a given row. The total plant density was approximately 5,714 plants/ ha. Interlaced with every five rows of coffee plants, trees of Fabaceae family were planted perpendicular to the prevailing winds to serve as windbreaks. The plantation is managed in a conventional manner, as is recommended for cultivating arabic type coffee, including annual fertilization. One aspect, in which the farm deviates from the convention, has to do with using legume plant matter as windbreaks. This incorporation of vegetal organic matter from leguminous plants is the central focus of the present study. Weeding of grasses and growing between the rows of coffee is performed manually and only when this undergrowth exceeds approximately 20 cm.

The forest area located next to the coffee plantation (less than 1 km away) was used as a control area. This forest is in the state of secondary succession; it has been left untouched and unmanaged since 1972, when the farm was acquired by EPA-MIG Sul.

To evaluate whether the organic matter from Fabaceae plants influences the edaphic mite community, the soils from the areas, shaded by these plants, were evaluated. The incorporation of organic matter occurs when the Fabaceae plants are annually thinned, and their branches, leaves and fruits fall beneath the coffee trees, thus coming into contact with the soil. This practice is common in organic coffee plantations and has the objective of enriching the soil by nutrients (e.g., nitrogen) and organic matter, derived from the decomposition of Fabaceae vegetal pieces (Ricci 2005).

Four species of Fabaceae were used to enrich the soil in the experimental area: acacia, *Acacia mangium* Wild.; pigeon pea, *Cajanus cajan* (L.) Millsp.; gliricidia, *Gliricidia sepium* (Jacq.) Walp.; and leucaena, *Leucaena leucocephala* (Lam.) de Wit. Each of them was incorporated into their respective coffee tree row. As previously mentioned, some rows received no shade and no organic matter from the Fabaceae plants (i.e. full sun).

Soil sampling was conducted during two collection events: one during the end of the dry season (5 November 2012) and another at the end of the rainy season (15 April 2013). In each season, soil samples were taken as follows. Four samples were taken from the areas where four different types of Fabaceae plants were incorporated into the soil. In addition, a sample was taken from within a coffee area without the incorporation of organic matter. Finally, a sample was taken from within an area of forest with native vegetation. Thus, the samples encompassed a total of six areas, each of them was sub-sampled four times in the dry season and four times in the rainy season. The soil samples taken from the different systems were spaced four meters from each other. Soil samples taken during the rainy season were obtained near the same spots that were used during the dry season; samples taken during both seasons were subsequently combined into one. Soil sampling was accomplished with a stainless-steel cylinder probe (50 mm internal diameter, height of 53 mm; 100 cm³) for the collection of undisturbed soil samples (BRAVIFER-Indústria de Equipamentos e Assessoria Agronômica Ltda, ME).

Edaphic organisms were extracted in the laboratory using a Berlese-Tullgren extracting funnel (Walter and Krantz 2009; Camargo *et al.* 2015). The samples were extracted into collecting flasks with 70% alcohol, using light and heat for seven days.

After extraction, mites were mounted on glass slides in a Hoyer's medium, counted with the assistance of a binocular stereomicroscope and identified with the use of a phase-contrast binocular microscope.

Mite sampling extraction and identification were conducted in the Laboratory of Agricultural Acarology EPAMIG Sul/Research Center, in Ecological Management of Pests and Plant Diseases— EcoCenter, in the city of Lavras, Brazil.

The mites were identified to the order and family levels following Krantz and Walter (2009), with the elevation of suborder Endeostigmata to order, as suggested by Pepato and Klimov (2015). In addition, to better visualize the results, the order Sarcoptiformes was split into the Cohort Astigmatina and the remaining Oribatida.

After the initial separation into orders and families, specimens were identified to the most specific taxonomic level possible. When species identification was not possible, the mites were separated into morphospecies to facilitate statistical analysis. Whenever possible, species identification and separation into morphological types (recognizable taxonomic units) was accomplished with the aid of specialists on various groups of mites.

Voucher specimens for the species and morphospecies of mites were deposited into the following collections: the acarological collection of the Laboratory of Agricultural Acarology EPA-MIG Sul (Lavras, Brazil); the Museo Nacional de Ciencias Naturales Bernardino Rivadavia (Buenos Aires, Argentina)—voucher specimens of Oribatida; the mite reference collection of Escola Superior de Agricultura "Luiz de Queiroz", (Piracicaba, Brazil)—voucher specimens of Acaridae and Mesostigamata; and the Museu de Ciências Naturais (ZAUMCN), UNIVATES—Centro Universitário (Lajeado, Brazil)—voucher specimens of Cunaxidae.

Data analysis

Mean species richness (number of species found) and mean total abundance were calculated for each sampling area. Generalized linear models (GLM) with contrast analysis were used to determine if there were significant differences in edaphic mite richness and abundance among the six sampled areas (native forest; coffee trees in full sun without the incorporation of nutrients; and coffee trees with the incorporation of organic matter from one of the species of Fabaceae—*A. mangium*, *C. cajan*, *G. sepium* and *L. leucocephala*),

Similarities among the communities of six sampling areas were quantified using Jaccard's index and represented graphically using non-metric multidimensional scaling (n-MDS). An analysis of similarities (ANOSIM) was conducted to test for significant differences in the composition of mite communities. The analyses were conducted using R (R Core Development Team 2014).

RESULTS AND DISCUSSION

Across all samples, 1,014 mite specimens of 143 different species were collected. Sarcoptiformes (71 morphospecies and 581 specimens) was the order with the greatest richness and abundance, followed by Mesostigmata (37 morphospecies and 293 specimens), Trombidiformes (28 morphospecies and 128 specimens) and Endeostigmata (7 morphospecies and 12 specimens), as presented in Table 1 and Appendix.

The forest soil had the greatest edaphic mite total abundance and richness (475 specimens and 69 species), while the soil enriched with acacia was

the least abundant and rich (27 specimens and 15 species), with the remaining areas having intermediate values (Tables 2 and 3).

As shown in the species discovery curve, which shows the number of species as a function of the number of specimens caught (Fig. 1), the sampling was not sufficient to completely sample the mite community present in the soil of the forestry system, because the asymptote was not achieved. In turn, in the environments with coffee plants, the species discovery curves approach more closely the asymptote, demonstrating that the number of species in these locations is lower than that in

Table 1

List of mite specimens collected in the order of abundance per family and percentage of occurrence in relation to the total.

Order	Abundance (total specimens)	%	Rate (Abundance/Richness)
Sarcoptiformes	581	59.4	Astigmatina: Acaridae (135 ind., 08 ssp.), Histiostomatidae (23 ind., 06 ssp.); Oribatida: Astegistidae (18 ind., 01 ssp.), Brachychthoniidae (06 ind., 02 ssp.), Camisiidae (01 ind., 01 ssp.), Ctenacaridae (01 ind., 01 ssp.), Cultroribula (02 ind., 01 ssp.); Damaeidae (16 ind., 01 ssp.), Damaeolidae (02 ind., 01 ssp.); Epilohmannidae (21 ind., 01 ssp.), Eremulidae (08 ind., 03 ssp.), Euphthiracaridae (05 ind., 02 ssp.), Galumnidae (11 ind., 01 ssp.), Haplozetidae (01 ind., 01 ssp.); Hypochthoniidae (85 ind., 02 ssp.), Licnodamaeidae (41 ind., 01 ssp.); Lohmaniidae (02 ind., 01 ssp.), Nothridae (01 ind., 03 ssp.), Microzetidae (01 ind., 01 ssp.), Oribatellidae (11 ind., 01 ssp.); Scheloribatidae (25 ind., 01 ssp.), Suctobelbidae (52 ind., 6 ssp.), Tectocepheidae (08 ind., 1 ssp.); youth Oribatida not identified (73)
Mesostigmata	293	27.2	Ameroseiidae (01 ind., 01 ssp.), Ascidae (86 ind., 10 ssp.), Blasttisociidae (18 ind., 01 ssp.), Eviphididae (01 ind, 01 ssp.), Rhodacaridae (74 ind., 04 ssp.), Laelapidae (29 ind., 04 ssp.), Melicharidae (23 ind., 02 ssp.), Ologamasidae (28 ind., 03 ssp.), Podocinidae (01 ind., 01 ssp.), Veigaiidae (01 ind., 01 ssp.) youth Uropodina. (31 in., 08 ssp.), youth Gamasina (01 ind., 01 ssp.)
Trombidiformes	128	12.4	Cheyletidae (02 ind., 01 ssp.), Cunaxidae (08 ind., 06 ssp.), Ereynetidae (02 ind., 01 ssp.), Erythraeidae (01 ind., 01 ssp.), Eupodidae (62 ind., 02 ssp.), Scutacaridae (36 ind., 06 ssp.), Neopygmephoridae (06 ind., 03 ssp.), Rhagidiidae (06 ind., 02 ssp.), Ereynetidae (02 ind., 01 ssp.), Stigmaeidae (03 ind., 02 ssp.), Tydeidae (01 ind., 01 ssp.), Smaridae (01 ind., 01 ssp.).
Endeostigmata	12	1.0	Alycidae (09 ind., 4 ssp.), Alicorhagiidae (02 ind., 02 ssp.), Nanorchestidae (01 ind., 1 ssp.)
Total	1,014	100.0	

ind.-abundance of specimens; ssp.-number of morphospecies.

natural forest environments, and that the sampling was close to ideal, having sampled almost the entire diversity of the mite communities present in the local soils (Fig. 1).

The abundance of each edaphic mite group varied, and the number of specimens depended on the type of soil and the quantity of organic matter, as well as on other environmental factors, such as moisture content and pH (Badejo 1990). However, mites are one of the edaphic groups with the largest number of species, both in natural and in cultivated areas (Hijii 1987; Minor and Cianciolo 2007; Mussury *et al.* 2008; Souto *et al.* 2008; Lorenzon and Machado Filho 2012). In the present study, as in other similar studies conducted in Brazil, oribatid mite species (Acari: Oribatida), the abundance and richness categories dominated (Badejo 1990; Badejo *et al.* 2004; Duarte 2004; Morais *et al.* 2010). In addition, it has been noted that the

edaphic arthropod fauna, including mites, can vary with changes to the soil management system, especially when natural systems are replaced by monocultures (Crossley *et al.* 1992; Minor and Cianciolo 2007). In the present study, significant differences between samplings conducted in the forest and in the coffee cultivation systems were clear (Fig. 2, Tables 2 and 3), with a decrease in the species richness and abundance occurring in the latter environment.

As previously mentioned, the native forest had the greatest abundance of mites in the soil. However, among the coffee plantation treatments, there were no significant differences in edaphic mite abundance between those areas where organic matter was not incorporated, and those where organic matter of *C. cajan*, *G. sepium* or *L. leucocephala* were incorporated. The areas where organic matter of *A. mangium* was incorporated, had

Table 2

Total abundance of edaphic mite specimens, sampled at EPAMIG Sul Experimental Farm, São Sebastião do Paraíso, MG, 2012–2013. Forest—native forest; CPFS—coffee tree plantation in full sun without organic matter; *C. cajan* (CC), *L. leucocephala* (LL), *G. sepium* (GS) and *A. mangium* (AM)—coffee plantation areas with the incorporation of organic matter from species of Fabaceae.

Taxon	Forest	CPFS	CC	LL	GS	AM
Oribatida	301	52	18	25	21	6
Astigmatina	4	22	36	59	32	5
Mesostigmata	102	47	52	64	13	15
Trombidiformes	58	9	14	33	13	1
Endeostigmata	10	0	1	0	1	0
Total	475	130	121	181	80	27

Table 3

 Total species richness of edaphic mites sampled at EPAMIG Sul Experimental Farm, São Sebastião do Paraíso, MG, 2012–2013. Forest—native forest; CPFS—coffee tree plantation in full sun without organic matter;
C. cajan (CC), *L. leucocephala* (LL), *G. sepium* (GS) and *A. mangium* (AM)—coffee plantation areas with the incorporation of organic matter from species of Fabaceae.

Taxon	Forest	CPFS	CC	LL	GS	AM
Oribatida	43	21	11	12	6	3
Astigmatina	4	5	5	8	3	2
Mesostigmata	15	13	16	8	8	9
Trombidiformes	7	6	2	3	1	1
Endeostigmata	0	1	0	1	0	0
Total	69	46	34	32	18	15

significantly lower abundance than the others (Tables 2 and 3, Fig. 2).

Significant differences were also observed in edaphic mite species richness, with the highest being associated with the native forest environment. Among the treatments, the leucena, *L. leucocephala*, had the greatest richness, followed by areas where no organic matter was incorporated. The areas where pigeon pea, *C. cajans*, and gliricidia, *G. sepium*, were incorporated, did not differ significantly. The areas where acacia, *A. mangium*, was incorporated, had the lowest species richness and abundance (Tables 2 and 3, Fig. 2).

The edaphic mite group that was most affected by the changes to the crop management system was Oribatida: its richness and abundance decreased even in the areas where organic matter of Fabaceae species was incorporated into the soil (Tables 2 and 3). This group of mites is sensitive to environmental changes. Oribatid mites are an important component of the edaphic fauna, since they play a role in nutrient cycling. Therefore, these mites are essential for maintaining natural cycles of renovation and incorporation of organic matter into the soil (Crossley *et al.* 1992; Behan-Pelletier 1999; Minor and Cianciolo 2007; Norton and Behan-Pelletier 2009). The loss of these species can cause an imbalance in natural nutrient cycles. Thus, practices that assure presence of this group of species in cultivated areas are important for maintaining soil quality. Although the practice of incorporating organic matter into the soil is important for maintaining nutritional and physical properties of the soil (Ricci 2005), it is not sufficient for preserving the mite community. It was initially hypothesized that places where organic matter was incorporated into the soil, would have greater abundances of soil mites, and that their abundance and richness would be similar to those found in the natural forest. This, however, turned out not to be the case. Further research is needed to better understand how this practice may be implemented more efficiently in order to better maintain population sizes of mites in the soils of cultivated systems.

The hypothesis (which may explain the results of the present study) that the quantity and quality of the vegetable matter deposited under the coffee trees, is not adequate. A greater amount and diversity of vegetable substrates may be necessary. Determining the ideal amount and type of vegetable matter may be the topic of future research. However, there is evidence that changes in soil management practices, and an adequate incorporation of organic matter may significantly increase the abundance of oribatids and other edaphic mites (Khalil *et al.* 2016). In addition, it is known that



Fig. 1. Species accumulation curves for the soil mite communities, sampled at EPAMIG Sul Experimental Farm, São Sebastião do Paraíso, MG, 2012–2013. Forest—native forest; CPFS—coffee tree plantation in full sun without organic matter; *C. cajan, L. leucocephala, G. sepium* and *A. mangium*—coffee plantation areas with the incorporation of organic matter from species of Fabaceae.

the leaves of acacia, *A. mangium*, contain the chemical compounds, such as tannin (Correa 1984), which is potentially harmful to mite communities. Such compounds may reduce the populations of mite species (Fernández-Salas *et al.* 2011). Therefore, choosing the most appropriate plant species to be used in soil management practices must be an initial step towards preserving the mite community.

The practice of converting natural ecosystems into agricultural ones usually decreases microarthropod population in the soil. In addition, removing native vegetation may negatively affect the species richness and abundance of edaphic mite communities (Edwards and Lofty 1975; Wallwork 1976; Blevins *et al.* 1984; Minor and Cianciolo 2007). Such negative effects are also a result of drastic changes in microclimatic conditions, such as humidity and temperature, chemical changes, and the destruction of microhabitats (Crossley et al. 1981; Blevins et al. 1984; Perdue 1992). In addition to changes in species richness and abundance, alterations to the crop management system can lead to a complete change in the soil mite community with the substitution of species. All of the studied treatment environments possessed a low degree of similarity with the native forest. More specifically, the treated and forest environments frequently shared less than 18% of their species, which shows that different soil management practices strongly affect edaphic mite communities (Table 4).



Fig. 2. Comparison of means (GLM) for total species richness (A) and total abundance (B) of mites found in the studied systems at EPAMIG Sul Experimental Farm, São Sebastião do Paraíso, MG, 2012. Different letters indicate significant differences between means (p<0.05). Forest—native forest; CPFS—coffee tree plantation in full sun without organic matter; *Cajanus cajan* (CC), *Leucena leucocephala* (LL), *Gliricidia sepium* (GS) and *Acacia mangium* (AM)—coffee plantation areas with the incorporation of organic matter.

As shown in Fig. 3 and Table 5, the composition of the forest system mite community is significantly different from that of the cultivated systems; whether they are managed with the Fabaceae organic matter or not.

The substitution of native forests with cultivated systems may cause significant changes in the edaphic mite species richness, abundance and community structure. Nevertheless, the improvements in soil management techniques and agricultural production procedures may revert these impacts, as it can be observed in organic production systems (Khalil *et al.* 2016).

Biological indicators of soil quality are often used to understand complex environmental changes because they are easily measured, responsive to a wide range of stresses, and they can be employed rather quickly (Pulleman *et al.* 2012). Mites have a high potential of serving as biological indicators of soil quality (Gulvik 2007). Among the mite groups observed in the present study, oribatid mites exhibited the most evident variation. They responded with a strong reduction in species richness and abundance, especially in comparison of the natural forest with the cultivated systems. According to Gulvik (2007), using oribatids as indicators of soil quality is efficient because a large number of these species possess long lifespans, low fertility rates, lengthy development and low mobility. Therefore, changes in the environment, especially those caused by human activities, are likely to reduce the presence of Oribatida in the environment. It should be mentioned that species identification in tropical environments is often impossible because there are few qualified professionals with the necessary expertise. Nonetheless, oribatids can still be useful in comparing the parameters of richness and abundance among ecosystems with different levels of degradation.

Table 4

Similarities (Jaccard) among the sampled edaphic mite communities. Forest—native forest; CPFS—coffee tree plantation in full sun and without organic matter; *C. cajan* (CC), *L. leucocephala* (LL), *G. sepium* (GS) and *A. mangium* (AM)—coffee plantation areas with the incorporation of organic matter from the corresponding species of Fabaceae plants.

Environments	Forest	CPFS	CC	LL	GS
Forest					
CPFS	17.83				
CC	15.87	19.82			
LL	12.23	16.07	23.05		
GS	12.74	17.74	18.65	26.25	
AM	7.51	21.26	20.87	19.94	31.2

Table 5

 ANOSIM results, comparing mite communities found in the studied native forest and coffee crop systems. Forest—native forest; CPFS—the system without the incorporation of organic matter; C. cajan (CC),
L. leucocephala (LL), G. sepium (GS) and A. mangium (AM)—systems with the incorporation of organic matter of the corresponding Fabaceae species into the soil.

Environments	Т	P (perm)
CPFS vs forest	1.8014	0.0299
CC vs forest	2.6161	0.0271
LL vs forest	1.8065	0.0296
GS vs forest	3.0390	0.0284
AM vs forest	2.4764	0.0300



Fig. 3. Multivariate ordination (non-metric multidimensional scaling—NMDS) comparing mite species composition among the studied systems at EPAMIG Sul Experimental Farm, São Sebastião do Paraíso, MG, 2012. Forest—native forest; CPFS—coffee tree plantation in full sun without organic matter; *Cajanus cajan* (CC), *Leucena leucocephala* (LL), *Gliricidia sepium* (GS) and *Acacia mangium* (AM)—coffee plantation areas with the incorporation of organic matter.

CONCLUSION

The edaphic mite community in general, and oribatid mites in particular, were found to respond to the replacement of forest by cultivated plantations, as well as to the incorporation of material of different species of Fabaceae plants into the soil in the areas of coffee tree cultivation. The degrees of mite communities' responses were established by measuring the changes in their richness and abundance. Oribatida was one of the most relevant taxa in the present study, since it was widely distributed (i.e. it had the highest richness and abundance) among the studied environments, including the natural forest and areas of soil management practices. The response of orbatids to environmental alterations is expressive, and, according to our results, they can be used as effective biological indicators.

Since the incorporation of organic matter into the soil may minimize the negative effects of cropping, it should be considered as a viable soil management technique.

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Appendix

Taxonomic groups that were identified in each of the systems studied. Forest—native forest;
CPFS—coffee plantation in full sun, with the absence of organic matter; Cajanus cajan (CC),
Leucena leucocephala (LL), Gliricidia sepium (GS) and Acacia mangium (AM)-
coffee plantation areas with the incorporation of organic matter.

Order	Taxa	Genus/Species	Forest	CPFS	CC	LL	GS	AM
Endeostigmata	Alicorhagiidae	*	X					
	Alycidae	Alycus ssp.			X			
	Alycidae	Bimichaelia ssp.	X					
	Nanorchestidae	Nanorchestes ssp.					Х	
Mesostigmata	Ameroseiidae	Ameroseius ssp.	X					
	Ascidae	*	X					
	Ascidae	Asca ssp.	X			Х		
	Ascidae	Protogamasellus sigillophorus	X	Х	X		Х	
	Blattisociidae	Lasioseius ssp.	X		X	Х		
	Eviphididae	*			X			
	Laelapidae	Cosmolaelaps ssp.	X	Х	X			X
	Laelapidae	Gaeolaelaps ssp.			X			
	Laelapidae	Pseudoparasitus ssp.	X					
	Melicharidae	Proctolaelaps paulista		Х		Х		
	Melicharidae	Proctolaelaps ssp.						X
	Ologamasidae	Gamasiphis ssp.	X			X	X	X
	Ologamasidae	Neogamasellevans ssp.	X		X		Х	
	Podocinae	Podocinum ssp.	X					
	Rhodacaridae	Multidentor hodacarus	X	Х	X	Х		X
	Rhodacaridae	Protogamasellopsis ssp.		Х		Х		X
	Rhodacaridae	Rhodacarellus ssp.		Х		Х		X
	Uropodina	*	X			X	X	X
	Veigaiidae	*				Х		
Sarcoptiformes	Acaridae	*		Х	X	X		
	Acaridae	Tyrophagus ssp.	X	Х	X	Х	Х	X
	Histiostomatidae	*	X					
	Astegistidae	Cultroribula ssp.	X	X		X		
	Brachichthoniidae	*	X					
	Brachychthoniidae	Liochthonius aff. fimbriatissimus				X		
	Camisiidae	Camisia ssp.	X					
	Ctenacaridae	*	X					
	Cultroribula	Cultroribula ssp.					X	
	Damaeidae	*	X					

	Damaeolidae	Fosseremus ssp.		Х				
	Epilohmannidae	Epilohmannia pallida americana	Х	Х	Х	Х		
	Eremulidae	Eremulus ssp.	Х	Х				
	Eremulidae	Eremulus crisspus	Х					
	Euphthiricaridae	*	Х					
	Galumnidae	Galumna ssp.	Х	Х		Х		
	Haplozetidae	Rostrozetes ssp.	Х					
	Hypochthoniidae	Eohypochthonius ssp.	Х					
	Hypochthoniidae	Malacoangelia ssp.	Х					
	Licnodamaeidae	*	Х	Х				
	Lohmanniidae	Lohmannia ssp.	Х					
	Lohmanniidae	Papillacarus ssp.		Х				
	Mesoplophoridae	*	Х					
	Mesoplophoridae	Mesoplophora ssp.	Х				Х	
	Microzetidae	Berlesezetes ssp.		Х				
	Nothridae	Nothrus ssp.	Х					
	Oppiidae	Microppia ssp.	Х					
	Oppiidae	Lanceoppia ssp.	Х	Х				
	Oppiidae	Ramusella ssp.	Х					
	Oppiidae	Striatoppia ssp.	Х	Х	Х			
	Oribatellidae	Lamellobates botari	Х	Х	Х	Х		Х
	Pheroliodidae	Pheroliodes ssp.	Х					
	Quadroppiidae	Quadroppia ssp.	Х					
	Scheloribatidae	Scheloribates praencisus acuticlava	Х	Х	Х		Х	Х
	Suctobelbidae	*	Х			Х		
	Suctobelbidae	Suctobelbella ssp.	Х	Х	Х	Х		
	Suctobelbidae	Suctobelbella aff. ornatissima				Х		
	Suctobelbidae	Suctobelbella variosetosa	Х	Х	Х	Х		
	Suctobelbidae	Suctobelbella elegantula	Х	Х	Х	Х		
	Suctobelbidae	Suctobelbella aff. loksai	Х	Х	Х	Х		
	Suctobelbidae	Cultroribula cf. zicsii	Х	Х	Х	Х		
	Tectocepheidae	Tectocepheus velatus	Х	Х	Х		Х	
Trombidiformes	Cheyletidae	*					Х	
	Cunaxidae	Neocunaxides ssp.	Х	Х				
	Cunaxidae	Cunaxa ssp.	Х		Х			
	Cunaxidae	Coleoscirus ssp.		Х				
	Cunaxidae	Dactyloscirus ssp.		Х				
	Cunaxidae	Parabonzia ssp.			Х			

T.A.F. Carvalho, P.R. Reis, L.F.O. Bernardi et al.

Edaphic mites and their reponse to organic matter

Ereynetidae	*				X	
Eupalopsellidae	*					
Eupodidae	Eupodes ssp.	X	Х	Х	Х	Х
Trombiculidae	*	X				
Pygmephoridae	*	X	Х			
Rhagidiidae	*	X				
Scutacaridae	Imparipes ssp.	X				
Scutacaridae	Scutacarus ssp.	X	Х			
Stigmeidae	*	X				
Tydeidae	*			Х		

*—unidentified species.