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## Impact of crop management systems on soil nematode communities in South Brazil

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Received April 06, 2020 Accepted September 28, 2020 ABSTRACT: The analysis of nematode communities allows inferring consequences of management practices on the soil food web. We studied the taxonomic structure of nematode communities in preserved areas of the Atlantic forest (native forest = NF) in Ponta Grossa, Paraná State, Brazil, and in three different agroecosystems in neighboring areas to assess the effect of agricultural land use on nematode assemblages. Agroecosystems were located in a red latosol cropped during 30 years under conventional tillage (CT), no-tillage (NT), and minimal tillage (MT). We collected ten composite soil samples in each area in the summer. Nematodes were extracted by Baermann funnel and fixed with formalin. Subsequently, individuals were classified into taxonomic groups and counted on a Peters slide to determine densities of each taxon. Plant-parasitic and free-living nematodes were classified at the genus level. Data were analyzed with the parameters abundance, Bray & Curtis, Shannon-Weaver, and Simpson indexes. We recorded 35 genera and abundance of nematodes in MT and NT areas was more similar. Higher richness was observed in NF in relation to cropped areas, especially under NT and CT. The PCA and clustering analyses from both nematode communities and soil chemical characteristics showed that MT and CT were more similar and NT was clustered near NF. The replacement of native vegetation by cropping systems caused a reduction of nematode diversity, demonstrating the influence of agricultural practices on nematode communities.

Keywords: abundance, diversity, taxonomic structure, soil indicators

## Introduction

Nematodes are some of the most diverse soil animals (van den Hoogen et al., 2020) and are capable to respond quickly to environmental changes; therefore, they have been used as indicators of soil health in different crop systems and native areas (Neher, 2001; van den Hoogen et al., 2020). Bacterial-feeding nematodes can ingest a great number of bacterial cells per day and are attracted by high  ${\rm CO_2}$  concentrations in their soil. Their preying capacity releases  ${\rm NH_4}^+$  readily available to plants and other animals in the soil solution (Lavelle, 1997).

Abundance and diversity of nematodes in fields under different agricultural systems are used to infer disturbances to the soil fauna (van den Hoogen et al., 2019), as well as enrichment and disturbance of soil communities (Bongers, 1990; Ferris et al., 2001; Berkelmans et al., 2003). The intensification of agricultural practices generally leads to biodiversity loss (Matson et al., 1997; Tscharntke et al., 2005), reducing the taxonomic richness of soil nematode communities (Yeates and Bongers, 1999; Yeates and Stirling, 2008).

Reduced tillage practices are used to minimize soil disturbance, as they benefit many soil characteristics, such as soil structure and fauna (Sánches-Moreno et al., 2006). However, most studies focus on the effects of tillage and cropping patterns on plant-parasitic nematodes and, to a lesser extent, on the free-living nematode communities (Sánches-Moreno et al., 2006).

This study investigated the effects of different management systems on nematode communities in preserved areas of the Atlantic Forest in Ponta Grossa, Paraná State, Brazil, and in neighboring areas, based on plant-parasitic and free-living nematodes. The results are based on studies conducted under field conditions for over 30 years to (a) contribute to the knowledge of nematode diversity (plant-parasitic and free-living) and (b) study the influence of agricultural activities on nematode communities.

### **Materials and Methods**

#### Study site

The samples were collected from a long-term experiment at the municipality of Ponta Grossa, Paraná State, Brazil (25°05'42" S, 50°09'43" W, altitude of 875 m), which has a Cfb climate, as described by Zagatto et al. (2018). In this study, three types of land use systems were considered, which were kept for 30 years in a Geric Ferralsol (rhodic) according to FAO (FAO, 2006): notillage (NT), conventional tillage (CT), minimum tillage (MT) and a native forest (NF), used as a reference. Soil management was conducted according to Zagatto et al. (2018).

The dimensions of the experimental plots under CT and MT managements were  $100 \times 50$  m, and under no-tillage,  $100 \times 100$  m, with slopes of 6.9, 7.1 and 8.1 %, respectively. In these agricultural systems, we collected data on crop succession and weed management since 1995, as described by Zagatto et al. (2018): maize/ black oat/ soybean/ wheat/ soybean/ black oat + vetch. The NF was a remnant of mixed Ombrophilous forest and comprised an adjacent area with  $100 \times 100$  m.



### Soil sampling and nematode identification

Soil samples were collected in the summer (25 Feb 2011). Ten samples from each treatment (n = 10) were collected at 0-10 cm deep, totaling 1 kg of soil. Nematodes were extracted from a subsample of 50 cm³ using the Baermann funnel method, as described in Machado et al. (2010), and later fixed in TAF (formalin 40 %, triethanolamine, distilled water). All nematodes in each sample were counted at low magnification  $(40\times)$  and then identified at the genus level at higher magnification  $(100\times)$ , according to De Ley and Blaxter (2002).

At each sampling point, soil samples were also analyzed for chemical attributes and characterization is shown in Table 1.

#### Statistical analysis

Relative abundance of each taxon was calculated from the proportion of that taxon among the nematodes identified. Abundance was expressed as the number of nematodes in each taxon. Posteriorly, usual indicators for ecological studies were calculated: Richness (number of genera), the Simpson diversity index, the Shannon-Weaver diversity index, and the Bray and Curtis similarity index, using the vegan package (Oksanen et al., 2019) in R 2.15 program.

The dataset was submitted to principal component analysis (PCA) and later to the hierarchical clustering analysis. Based on the cophenetic matrix, the grouping method was the UPGMA. This multivariate analysis allowed determining the most robust discriminant variable and obtaining an overview of similarities between the management systems, based on nematode communities and the soil chemical analysis. We used the canonical correspondence analysis (CCA), a multivariate constrained ordination technique, to observe the combinations between chemical characteristics and nematode-feeding groups. All multivariate analyses were performed with the vegan package (Oksanen et al., 2019) in R 2.15 program.

### Results

# Influence of management practices on soil nematode community

Thirty-five nematode genera were identified in the samples collected (Table 2), which were affiliated to six orders of Adenophorea (Araeolaimida, Diplogasterida, Dorylaimida, Enoplida, Mononchida, and Triplonchida) and two orders of Secernentea (Rhabditida and Tylenchida). We verified 20 genera in CT, 32 in MT, 14 in NT and 32 in NF (Table 3).

We identified 15,571 nematodes/kg of soil in CT, with more than 70 % belonging to five genera/family: Aphelenchus (17.9 %), Cephalobus (5.4 %), Helicotylenchus (16.5 %), Prismatolaimus (5.2 %) and Rhabditidae (25.9 %). In MT, we verified 27,212 nematodes/kg with more than 77 % belonging to six genera/family: Aphelenchus (23.4 %), Dorylaimoides (4.4 %), Helicotylenchus (18.1 %), Labronema (5.1 %), Tylenchus (4.3 %) and Rhabditidae (22.5 %). On the other hand, in NT, we identified 6,891 nematodes/kg of soil, with more than 63 % belonging to two genera: Aphelenchus (19.6 %) and Helicotylenchus (44.1 %), while NF showed 12,338 nematodes/kg of soil, with more than 60 % belonging to three genera: Meloidogyne (21.8 %), Mesocriconema (24.8 %) and Xiphinema (13.7 %).

The Shannon-Weaver (H') diversity index confirmed the indications based on number of genera (or richness) relatively to the high diversity found in NF and MT samples (Table 3). The Bray and Curtis index (Table 4), based on quantitative values of abundance, evidenced that MT and CT areas were more similar, while MT was more dissimilar to NF.

# Soil chemical attributes effects on nematode community

The PCA and clustering analyses showed variability between the management systems for nematode communities and soil chemical attributes (Figures 1A and 1B). Based on nematode community composition

**Table 1** – Chemical analysis of soil from different management systems (NT = no tillage, CT = conventional tillage, MT = minimal tillage, NF = native forest) collected at 0-5 and 5-10 cm, in the municipality of Ponta Grossa, Paraná State, Brazil.

	pH CaCl <sub>2</sub>	Р	K	Al <sup>3+</sup>	Ca <sup>2+</sup>	Mg <sup>2+</sup>	H+AI	Т	٧	С
		mg dm <sup>-3</sup>			cmolo	c dm <sup>-3</sup>			%	g dm <sup>-3</sup>
NT 0-5 cm	4.33	21.58	0.24	0.53	3.32	1.35	10.18	15.09	32.4	39.8
NT 5-10 cm	4.31	11.55	0.16	0.84	2.20	0.94	8.56	11.86	27.2	28.4
MT 0-5 cm	4.40	8.68	0.20	0.42	2.96	1.23	8.24	12.63	34.5	31.6
MT 5-10 cm	4.49	7.62	0.11	0.38	2.87	1.17	7.54	11.68	35.3	26.9
CT 0-5 cm	4.34	7.31	0.22	0.48	2.57	1.08	15.62	19.49	32.1	27.6
CT 5-10 cm	4.37	7.09	0.12	0.50	2.56	1.04	7.62	11.34	32.8	26.7
NF 0-5 cm	4.24	4.16	0.32	1.13	3.68	1.68	11.97	17.65	32.1	46.2
NF 5-10 cm	4.11	1.89	0.21	1.57	1.60	0.82	11.49	14.11	17.7	32.7

P, K (Mehlich-1); Al, Ca, Mg = (KCl); T = cation exchange capacity at pH 7.0; V % = saturation of the cationic complex by bases.

**Table 2** – Effect of tillage system on total abundance of nematode taxa (number of nematodes / 50 cm<sup>3</sup> soil). Data from four sampling experimental plots (n = 40).

	Trophic group		Conventional tillage	Minimal tillage	No tillage	Native Forest
Aphelenchida	PP/FF	Aphelenchoides	-	$8.10 \pm 25.61$	-	16.55 ± 41.44
	PP/FF	Aphelenchus	139.25 ± 75.32	318.45 ± 200.37	67.60 ± 35.49	15.15 ± 21.00
Araeolaimida	BF	Plectus	1.25 ± 3.95	$3.60 \pm 7.59$	-	$0.65 \pm 2.05$
Diplogasterida		Mononchoides	18.40 ± 15.07	16.05 ± 29.63	-	$0.90 \pm 2.84$
Dorylaimida	PR	Carcharolaimus	-	$0.90 \pm 2.84$	-	-
	PR	Discolaimus	-	$9.75 \pm 18.80$	-	$8.80 \pm 23.61$
	PR	Discolaimoides	$4.35 \pm 13.75$	-	-	$6.95 \pm 16.24$
	OM	Dorylaimoides	24.75 ± 19.69	$59.85 \pm 65.59$	$25.90 \pm 36.02$	$15.35 \pm 23.60$
	OM	Dorylaimus	-	$39.75 \pm 54.72$	-	$4.80 \pm 10.97$
	OM	Labronema	$3.85 \pm 9.10$	69.85 ± 80.12	-	$0.55 \pm 1.74$
	OM	Laimydorus	$32.65 \pm 39.95$	$19.00 \pm 56.65$	-	$0.90 \pm 2.84$
	OM	Mesodorylaimus	4.30 ± 9.57	$9.20 \pm 12.79$	15.75 ± 12.63	$4.25 \pm 5.97$
	OM	Prodorylaimus	18.30 ± 15.57	$1.25 \pm 3.95$	-	-
	PP	Trichodorus	14.80 ± 20.28	$15.30 \pm 25.73$	$3.70 \pm 7.92$	$3.90 \pm 7.28$
	FF	Tylencholaimus	-	-	-	$0.55 \pm 1.74$
	PP	Xiphidorus	-	$7.50 \pm 20.75$	-	9.60 ± 14.47
	PP	Xiphinema	-	$2.80 \pm 4.51$	-	85.15 ± 61.74
Enoplida	BF	Prismatolaimus	40.40 ± 43.37	22.90 ± 31.42	13.80 ± 23.54	21.05 ± 33.57
Mononchida	PR	Mononchus	15.00 ± 16.67	$7.90 \pm 8.56$	$2.40 \pm 5.25$	5.65 ± 12.85
	PR	Mylonchulus	1.40 ± 4.42	21.70 ± 28.14	-	$3.00 \pm 9.48$
	PR	Prionchulus	-	$1.65 \pm 5.22$	-	2.75 ± 8.69
Rhabditida	BF	Acrobeles	3.25 ± 5.36	$7.80 \pm 7.99$	8.22 ± 8.21	2.35 ± 5.76
	BF	Acrobeloides	-	5.40 ± 17.07	-	8.25 ± 17.13
	BF	Cephalobus	42.35 ± 38.61	16.60 ± 27.04	8.00 ± 13.75	0.65 ± 2.05
	BF	Eucephalobus	-	$2.55 \pm 5.66$	-	0.55 ± 1.74
	BF	Panagrolaimus	-	13.05 ± 41.27	-	7.85 ± 16.55
	BF	Rhabditidae	202.00 ± 122.43	305.90 ± 179.78	17.15 ± 22.22	24.80 ± 29.45
Triplonchida	PP	Paratrichodorus	-	15.00 ± 21.21	-	1.50 ± 4.74
Tylenchida		Filenchus	1.90 ± 4.03	16.95 ± 20.05	4.45 ± 8.64	8.55 ± 15.83
	PP	Helicotylenchus	128.60 ± 65.17	246.45 ± 250.07	151.95 ± 76.53	19.30 ± 27.34
	PP	Meloidogyne	78.40 ± 64.22	-	-	135.05 ± 208.52
	PP	Mesocriconema	-	$10.35 \pm 26.19$	-	154.10 ± 415.89
	PP	Paratylenchus	-	3.70 ± 8.77	-	10.60 ± 17.59
	PP	Pratylenchus	3.35 ± 5.68	20.85 ± 62.83	$0.95 \pm 2.00$	-
	PP	Tylenchus	-	58.85 ± 41.72	18.25 ± 16.47	36.90 ± 65.89
		•				

PP = plant-parasitic; PR = predator; BF = bacterial-feeder; OM = omnivores; FF = fungal-feeders; PP/FF = variable between plant-parasitic and fungal-feeder.

**Table 3** – Nematode diversity in the soil sampled in different cropping systems and native forest in South Brazil.

Index	Conventional tillage	Minimal tillage	No tillage	Native Forest
Richness	20	32	14	32
H'	0.97	1.04	0.80	1.06
J'	0.74	0.69	0.70	0.70

Richness = number of genera; H' = Shannon-Weaver diversity index; J' = Simpson index.

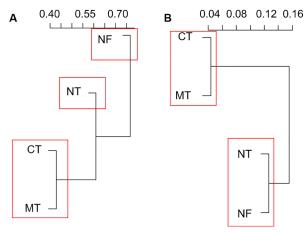
(Figure 1A), CT was grouped by the clustering analysis with MT, whereas NT formed a separate group near NF, in accordance with to the Bray and Curtis index classification. The soil chemical analysis (Figure 1B) showed that NF was grouped with NT, while MT was grouped with CT.

**Table 4** – Relationships between native forest (NF), minimal tillage (MT), conventional tillage (CT) and no-tillage (NT) areas, based on nematode community similarities accessed by Bray and Curtis index calculated from total abundance.

	Bray and Curtis
$NF \times MT$	0.77
$NF \times NT$	0.75
NF × CT	0.71
$MT \times NT$	0.61
$MT \times CT$	0.43
NT × CT	0.50

The canonical correspondence analysis (Figure 2) performed with data of nematode community and the soil chemical analysis showed that NF was positively correlated to higher concentrations of Al and C, whereas

MT and NT were correlated to P concentrations. Other soil characteristics were not significant in the analysis and were excluded from the graphic representation. Regarding nematode communities (Figure 2), MT and

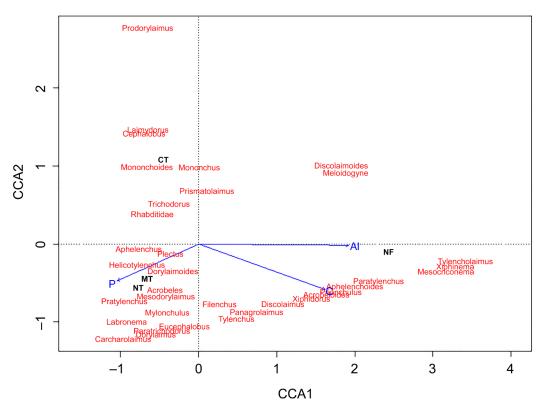


**Figure 1** – Dendrogram illustrating the dissimilarity between treatments (NF = native forest; NT = no-tillage; MT = minimal tillage; CT = conventional tillage) obtained by the UPGMA algorithm, based on the Euclidean distance, calculated from the average of nematode-feeding groups (A) and the soil chemical analysis (B).

NT were correlated to P content and omnivores, plant-parasitic, and to predators, to a lesser extent. NF was correlated to fungal-feeders, plant-parasitic nematodes, and high Al content, while CT was not correlated to the chemical analysis; nevertheless, it was correlated to bacterial-feeders and predators.

### **Discussion**

Studies conducted in Uruguay and Brazil showed that discrimination of different cropping systems using only abundance values was not applicable, when native areas were compared to cropped areas (Korenko and Schmidt, 2007; Tomazini et al., 2008). However, in systems under more intensive cropping, such as CT and MT, values of total density were higher. Several authors reported the predominance of higher density values in areas submitted to more intensive cultural traits when compared to native areas (Andrade et al., 2004; Goulart and Ferraz, 2003; Mattos et al., 2006), in Brazilian ecosystems. This can be explained as some crops host plant-parasitic species in the field, increasing the number of nematodes in the samples, as observed for Helicotylenchus and Aphelenchus in our study. Yeates and King (1997) and Valocká et al. (2001) compared nematode communities in native pastures with



**Figure 2** – Canonical correspondence analysis bi-plot of association of soil properties (P = phosphorus; C = carbon; Al = aluminum) and nematode community in different crop systems (NF = native forest; MT = minimal tillage; CT = conventional tillage; NT = no-tillage). Data represent all samples combined (*n* = 40). Eigenvalues (lambda) are 0.5113, 0.1273, and 0.1001 for the first (horizontal), second (vertical) and third axes, respectively. The first two axes explained 64.8 % of the variation.

cropped areas and observed that disturbances caused by agricultural practices increased the total abundance and richness of nematodes. This is possibly due to the increase in plant-parasitic nematode population, which could be so small in the soil and not be detected without the presence of the host plant.

Dorylaimida, such as *Labronema* and *Discolaimoides*, is cited as abundant in Cerrado soils and little detected in cropped soils (Goulart and Ferraz, 2003); however, in this study, both species occurred in cropped areas, demonstrating low sensibility to agricultural land use. In addition, agricultural crops could increase abundance of bacterial-feeding nematodes, especially due to the presence of organic matter (Shaw et al., 2019) and fungal-feeders, which sporadically parasitize on aerial parts of plants and could be favored or introduced by seeds of the agricultural crops (Yeates, 1991).

Mesocriconema, of the order Tylenchida, was observed in higher abundance in NF, demonstrating that this nematode is more sensitive to disturbances caused by the agricultural practices, such as tillage (Cares and Huang, 1991; Yeates, 1991; Gomes et al., 2003). In the Cerrado region of Brazil, nematodes of Criconematidae family have played an important role in distinguishing native from cropped areas (Mattos et al., 2008), since they are very sensitive to disturbances resulting from the replacement of native areas by annual crops (Goulart and Ferraz, 2003). Other genera from Tylenchida were present in all sampled areas, but with great variation in the density values, such as Helicotylenchus, the dominant genus in Tylenchida in this study, possibly favored by soybean cropping in CT and MT (Machado et al., 2019). The same can be concluded for Mononchida, found in all sampled areas and in great variation. These nematodes are predators and supposedly sensitive to environmental changes (Niles and Freckman, 1998; Wasilewska, 1997); nevertheless, they were not observed in our study.

The Shannon-Weaver and the Bray and Curtis indexes showed high diversity in NF and MT samples and that these areas were more similar. Conversely, CT was more dissimilar to NT. These observations corroborated previous reports in which the number and richness of genera or species sometimes constitutes better tools to evaluate nematode diversity in different ecosystems (Mattos et al., 2006); nevertheless, the efficiency of these parameters is controversy in literature. Freckman and Ettema (1993) considered richness and the Shannon-Weaver index little useful in distinguishing annual from perennial crops, while Yeates and Bird (1994) and McSorley and Frederick (1996) considered these approaches efficient.

In addition, cropped areas, except for MT, have a lower richness than NF. These findings agree with most similar studies in the literature, which report a higher number of taxa in native vegetation than in cropped areas (Freckman and Ettema, 1993; Háněl, 1995; Yeates and King, 1997; Mattos et al., 2006; Goulart and Ferraz, 2003; Gomes et al., 2003; Tomazini et al., 2008).

Regarding the effect of soil chemical attributes on nematode communities, NF was positively correlated to higher Al and C concentrations, whereas MT was correlated to P concentration. Thus, we can infer that high Al concentration is characteristic of this native type of soil where the experiment was conducted, that is, Campos Gerais, characterized by poor and washed soils, deficient in P, Ca, N, and K, with low pH and Al high concentration (Bodziak Júnior and Maack, 2001). The low pH observed in these soils could lead to an increase in the Al toxicity, which might reduce microbial biomass, generally observed after N addition to the soil (Shaw et al., 2019), influencing soil nematode communities, especially those dependent on microbial biomass.

The analysis of feeding groups provides information on changes in decomposition pathways between the sites studied. Abundance of bacterial-feeders could increase due to the microbial activity caused by fertilization or accelerated decomposition (Ettema and Bongers, 1993) and respond to OM input depending on the quantity and quality of OM (DuPont et al., 2009). Nevertheless, several bacteria in agricultural fields can solubilize insoluble forms of P to make them available for plant growth and have been considered as predominant source of P availability (Ilmer and Schinner, 1992). In our samples, bacterial-feeders were also correlated with the P content, possibly due to high concentration of P-solubilizing bacteria.

On the other hand, enrichment opportunist fungal-feeding nematodes react rapidly to the addition of organic materials, because fungal-feeders have high respiration rates and low N needs, mineralizing N to plants (Chen and Ferris, 1999; Ferris et al., 1997). This information if confirmed by our study where fungal-feeders were correlated with high C concentration. Besides, the higher concentration of C was also correlated with NF samples, reflecting the decomposition pathway of natural forests, predominantly by fungal-feeding nematodes. Besides, the black oat and wheat crops could explain high abundance of Aphelenchus in CT and MT in the winter, since these plant species can host this genus and favor the increase of its populations in the soil (Singh et al., 2013).

Therefore, these results may serve as link between microorganism and energy flow in the soils analyzed, providing a valuable tool to access the nature of decomposition pathways. Population of fungal- and bacterial-feeders are useful for tracking disturbances within a crop system, although the responses of nematode communities to agricultural intensification are not easy to interpret and correlate to soil properties (Sánchez-Moreno et al., 2006). Nematode communities can differentiate diverse habitats or cropping systems, especially in large temporal or land scales experiments (Levin, 1992). Therefore, we observed that nematode communities were capable of grouping MT and CT

in a separate cluster from NT and NF, indicating that aggressive cropping systems differ from conservative ones. Popovici and Ciobanu (2000) also found correlation between the composition of the nematode community and habitat characteristics, as different vegetation and crop systems.

In summary, the analyses of nematode communities successfully reported shifts in land use after 30 years in a field experiment in southern Brazil. Abundance of nematodes demonstrated that MT and NT areas were more similar and CT was more dissimilar to NT. Besides, cropped areas, except under MT, have a lower richness than NF. The PCA and clustering analysis also discriminated the impact of land use in the soil nematode communities, since NF was grouped with NT and MT with CT, based on the soil chemical analysis and nematode communities. These results indicated that nematode diversity is important for long-term stability of soil ecosystem and its use as soil health indicators merits further studies covering larger areas under soil conservation systems, such as NT and intercropping in Brazil.

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## **Authors' Contributions**

Conceptualization: Machado, A.C.Z.; Zagatto, M.R.G.; Skora Neto, F.; Silva, S.A.; Zanão Júnior, L.A. Data acquisition: Zagatto, M.R.G.; Skora Neto, F.; Zanão Júnior, L.A. Data analysis: Silva, S.A. Design of methodology: Machado, A.C.Z.; Zagatto, M.R.G.; Skora Neto, F.; Zanão Júnior, L.A. Writing and editing: Machado, A.C.Z.; Zagatto, M.R.G.; Silva, S.A.; Zanão Júnior, L.A.

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