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Article

Soil Free Living Nematode Composition in the Vicinity of Vineyard Rhizosphere in a Basaltic Soil

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Abstract: Soil free living nematodes community composition was compared for five agro-management systems in a basaltic soil. The sites differed in their long-term management system and were designated as organic farming (OM), conventional (MI), natural pasture (NP), foliage farming (FF) and open field (OF). The effect of five diverse land use types in vineyard growth in basaltic soils on soil free living nematodes, density, diversity, and functionality in the plant rhizosphere was examined. We found 22 families, 42 genera of which only 14 genera were present in all soil samples, three bacteria feeders, 4 fungi feeders, 3 omnivores-predators and 4 plant parasites, in the soil collected from the five sampling sites. The organic and conventional managements were found to influence soil free living nematode community and its trophic composition. The higher abundance of bacteria feeders included the Cephalobidae and Panagrolaimidae families. The fungi feeders included the Anguinidae and Aphelenchidae families. Ominvores-predators included the Aprocelaimidae and Qudsianematidae, and the plant parasites included the Anguinidae (genus *Anguina*), Nothotylenchidae and Tylenchidae families. The changes in plant parasite nematodes were one of the main pests in the below-ground rhizosphere biome. Promoting the bacterial community could be a beneficial and promising way for a considerable increase in organic management.

Keywords: soil nematodes; community structure; basaltic soil; vineyard

1. Introduction

In the last 20 years, the rate of vineyards planted has increased dramatically all over the world. It was estimated to be 7.3 mha in 2021, with a world market of 34.3 bn EUR (Roca, 2022). The viticultural areas use a broad spectrum of soils, although experts suggest that the soil should be well drained and loamy, containing a mix of sand, silt, and clay. However, grapes also tolerate poor alkaline soils. Regarding climate, grapevines can grow in a wide range of climates, from mild tropical to arid or cold. However, grape quality is substantially affected by soil type and origin. Volcanic origin soils contain a wide range of minerals, have low water retention and organic content, are of great potential for viticultural use (Maltman, 2018; Szabo, 2016; Wilson et al., 2016). In Israel, the Golan Heights volcanic soils are basaltic clay soils found at an altitude of 400 to 1,200 meters above sea level. These soils are in demand by many vineyards that consider the basaltic soils, temperature, and altitude as conditions in which grapes of high-quality grow.

The capacity of soil to function as a vital living ecosystem that fulfills all its nutrient supply functions is a fundamental requirement for the plants grown within it. These functions are dependent on the soil biota that maintains diverse soil organisms with high activity, retains and decomposes organic matter, which can meet the plant's nutrient demands.

Soil quality is predominantly assessed by the physical structure and chemical nutrient levels, followed by its biological components (Jing et al., 2015; Paz-Ferreiro and Fu, 2013). The composition and function of the biotic components of the soil food web are determined by resources supplied by organic matter. In agricultural systems, organic and mineral amendments can be supplied from other

sources that replace their enrichment role. The applied organic substrates are degraded by bacteria and fungi, that can potentially be used as indicators representing the nature of the organic matter. However, according to Ferris and Bongers (2006) available methods do not reliably indicate the soil microbial community composition, while the nematodes that regulate the microbial community and enhance plant nutrition, are a good indicator of substrate quality and nutrient release along the decomposition process (Christensen et al., 1992; Griffiths, 1994; Neher, 2001). The nematodes constitute a numerically important component of the soil biotic community, and their community composition follows the changes in food source availability depending on their trophic functions, which include bacteria, fungi, plant parasites, omnivores, and predators (Ferris and Bongers, 2006; Georgieva et al., 2005; Yeates, 2003; Yeates et al., 1993). Earlier studies had elucidated their importance due to their ability to reflect changes in soil physico-chemical properties and function based on nematode community structure (Neher et al., 1997). In agricultural management the use of diversity indices that combine between taxa richness and evenness may contribute to elucidate the frequency of less abundant trophic groups (Ludwig and Reynolds, 1988; Neher et al., 1997). According to Wardle and Yeates (1993), the most dominant interaction in the soil food web is determined by bacteria and fungi predation and competition, that may be limited by resource quality. In agro-ecosystems, where soil bacteria and fungi are closely linked with input of organic matter, the nematode community, and its trophic components are positively correlated to the amount and turnover of organic matter. Since nematodes occupy an important position in the detritus food web, they can be used as indicators for determining the effects of the different agro-managements on the soil detritus food web (Steinberger et al., 2001; Steinberger and Sarig, 1993). Application of organic materials from different sources to the soil will trigger changes in soil physical and chemical components, and in the bacterial and fungal communities, which will influence the trophic composition of the soil nematode community. Alteration between the different vineyard farming practices affect soil properties, some of them can have negative effects on soil quality (Coll et al., 2012). Recent studies conducted in vineyard soil have elucidate the importance in the use of nematodes in soil quality determination as playing a key role in soil organic matter decomposition. The use of soil free living nematodes that had presented to be a good bioindicators of vineyard soil quality is based on trophic groups diversity and characteristics demographic groups such as colonizers and persisters (Bongers, 1990; Yeates et al., 1993).

The present study aimed to determine the interaction between the soil free living nematodes community structure, from both a taxonomic and a functional aspect, and how it is affected by five long-term managements on basaltic soil in Northern Israel. The objective of the study was to compare the soil free-living nematode community in terms of density, diversity, and functionality between the five agro-managements in the vineyard rhizosphere of a basalt soil ecosystem. The rhizosphere is recognized as a hotspot for biota activity, functioning as a precursor to soil organic matter affecting community composition and diversity.

We hypothesized the following: (1) Organic and foliar managements, that increase organic matter, will stimulate bacteria and fungi feeder nematodes more than conventional, open field and natural pasture; (2) In undisturbed sites, there will be a decrease in predation pressure; (3) Guilds of nematodes (bacteria and fungi feeders - e.g., Cephalobidae, Aphelenchidae, Aphelenchoididae, respectively) will be responsive to changes in the abundance of their food.

2. Materials and Methods

2.1. Study Site and Sampling

The soil samples were collected from five different sites, northern Golan Heights ($33^{\circ}04'10''N$ $35^{\circ}46'11''E$) next to Mount Shifon at 820 m above sea level. This area has a humid Mediterranean climate, with cold winters ($6-8^{\circ}C$ range in January) and hot summers ($24-26^{\circ}C$ in August). The average multiannual rainfall is 760 mm y^{-1} (Israel Meteorological Service gov.il), and the topography of the area is of slopes with moderate gradient. On the surface, which is covered with a thick layer of basalt, tuff, and scoria, stands out a tall volcanic mound: Shifon Mountain, which creates a mountainous morphology (Dan et al., 1972). The soil is a Mediterranean reddish-brown basalt soil, on the slopes of the volcanic cone (Yaalon et al., 1974).

Five replicates of soil samples were collected by using a randomized sampling technique at a depth of the upper soil layer (0-10 cm) adjacent to the trunk of the vine plants rhizosphere, under five different plants in December 2021. Each soil sample were place in a individual plastic bag and placed in an insulation container transported to the laboratory and stored in $4^{\circ}C$.

After sampling, the soils were at the existing moisture to minimize changes in nematode populations. Subsamples were taken from each sample for estimation of nematode populations and different soil parameters. Soil moisture was determined gravimetrically by drying samples at $105^{\circ}C$ for 48 h and expressed as a percentage of dry weight.

Soil was sampled from five sites: 1. Merom Golan *Organic Vineyard* (MO) ($33^{\circ}03'50.9''N$ $35^{\circ}45'10.6''E$). A certified organic vineyard planted in 2014. Fertilization is based on compost, applied once every 2 years; 2. Merom-Golan Intensive – *Conventional Vineyard* (MI) ($33^{\circ}03'29.5''N$ $35^{\circ}44'57.4''E$). An intensive vineyard that was planted in 2017. Fertilization is applied according to leaf analysis and compost is applied, on demand. Helium (strong a long-lasting herbicide) was applied a year ago. 3. *Natural pasture* (NP) ($33^{\circ}03'44.4''N$ $35^{\circ}45'06.9''E$); 4. *Foliar fertilization* (FF) was planted in 2014 ($33^{\circ}04'45.8''N$ $35^{\circ}46'21.8''E$). Fertilization is based on foliar application of mainly N, K and micro-Mg, Zn and compost application, whiteout herbicide application. 5. *Open field* (OF), ($33^{\circ}04'45.6''N$ $35^{\circ}46'22.5''E$). Each sample included 3 replications.

2.2. Soil Free-Living Nematode Community Analysis:

Soil free-living nematodes were extracted using a modified Baermann funnel procedure (Cairns, 1960) with 200 grams of soil. The extracted nematodes were counted under a binocular to determine total nematode amount per 100 grams dry soil. The nematodes were transferred to a 1.5 ml tube and centrifuged for 10 min at 10,000 rpm to reduce the amount of remaining water. The obtained pellets were placed at $-20^{\circ}C$ until DNA extraction. Nematode DNA was extracted from the pellets using PureLink Genomic DNA Mini Kit (Invertogen, Thermo Fisher Scientific Inc., Waltham). The eluted DNA was stored at $-20^{\circ}C$ until PCR amplification. DNA was amplified by PCR using SimpliAmp™ thermal cycler (Thermofischer Scientific), by mixing 12.5 μl PCR BIO HS Taq Mix Red, 9.5 μl ultrapure water, 1 μl extracted DNA, 1 μl CS1-NF1 (ACACTGACGACATGGTTCTACAGGTGGTGCATGGCCGTTCTAGTT) and 1 μl CS2-18sr2b (TACGGTAGCAGAGACTTGGTCTTACAAAGGGCAGGGACGTAAT). The thermal cycling program was set to: $98^{\circ}C$ for 30 sec, 20 cycles of $98^{\circ}C$ for 10 sec, $58^{\circ}C$ for 30 sec, $72^{\circ}C$ for 1.5 min, and after the cycles $72^{\circ}C$ for 10 min. Sequencing (Miseq) was performed at the Hylabs Laboratory Ltd. (Rehovot, Israel), (www.hylabs.co.il) sequencing facility using an Illumina sequencing platform (Illumina Inc., San Diego, CA, USA).

2.2.1. Ecological Indices and Statistical Analysis

The characteristics of the nematode communities were described by means of indices. The classification of trophic groups was assigned to: (1) bacterivores; (2) fungivores; (3) plant-parasites; and (4) omnivores-predators (Liang et al., 2000; Steinberger and Sarig, 1993; Yeates et al., 1993). The nematode community was analyzed by the following approaches: (1) absolute abundance of

individuals expressed per 100 g⁻¹ dry soil; (2) trophic structure; (3) FB: (Fungi feeders/Bacteria feeders, or - FF/BF) reflects the structure of the microflora community. Bacteria-based food webs (lower values) exhibit higher decomposition rates than fungi-based webs (Twinn et al., 1974) ; (4) Wasilewska index (WI) = ratio of (fungivores+bacterivores) to plant parasites [(FF+BF)/PP], which represents substantial changes in the trophic structure of the community, indicates the dominant pathway of mineralization (Wasilewska, 1994, 1991); (5) Shannon index (H'), a species diversity measure which gives more weight to rare species, H'+Pi(ln Pi), where Pi is the proportion of each trophic group in the total population (Pielou, 1975); (6) trophic diversity, TD= 1/Σ P_{2i}, where Pi is the proportion of the individuals in the i-th trophic group (Higgins and Thiel, 1988; Neher, 2001) and describes the trophic group distribution; (7) genus dominance, $\lambda = \frac{P_{2i}}{\sum P_{2i}}$ (Neher, 2001; Simpson, 1949); (8) modified maturity index (ΣMI), is the c-p value assigned by (Bongers, 1990) of the i-th taxon and q_i is the proportion of the i-th taxon in the nematode community. The c-p values describe the nematode life strategies and range from 1 (for colonizers and individuals tolerant to disturbance) to 5 (for persisters and individuals sensitive to disturbance), including plant-feeding nematodes (Yeates et al., 1994). ΣMI incorporates ecological characteristics of families based on a colonizer-to-persister scale of 1–5, where lower ΣMI values indicate more disturbed environments; (9) evenness J' = H'/ln (S), where S is the number of taxa (Simpson, 1949) is highest when all species in a sample have the same abundance.

2.3. Statistical Analysis

The data were subjected to statistical analysis of variance using the SAS model (ANOVA and Duncan's multiple range test, T-test, genera indicator analysis and Pearson correlation coefficients) to evaluate differences between the treatments, by using the statistical package Statistica 4.3. Differences obtained at levels of p<0.05 were considered significant.

The Indicator Species Analysis was run in R using the "multipatt" function from the "indicspecies" package. This function calculates the indicator value for each genus in relation to each type of land management. For those genera that were found to be significantly different (p <= 0.05), we provide bar plots created in R using gplot of their relative abundance across the different land management systems.

3. Results

3.1. Soil Nematode Community

3.1.1. Total Number of Nematodes

The mean total number of soil free living nematodes ranged between 108 individuals 100⁻¹ g dry soil in samples collected from the conventional management to 776 individuals 100⁻¹g dry soil (Figure 1) in the foliar fertilization treatment. No significant differences were obtained in the total number of nematodes between the natural pasture (NP) and open field (OF) that represent control sites. The total number of nematodes in the organic (OM) management site (339.8 individuals 100⁻¹g dry soil) was almost two-fold lower compared to the FF management site (775.5 individuals 100⁻¹g dry soil).

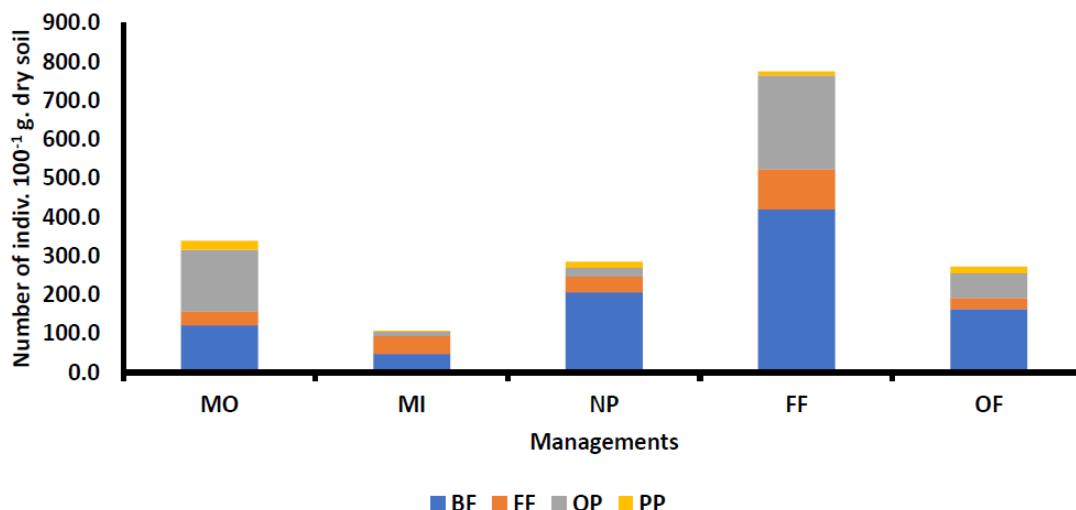


Figure 1. Total nematode abundance of the different samples, with different trophic groups in different colors. MO – organic vineyard; MI – conventional vineyard; NP – natural pasture; FF – foliar fertilization; OF – Open field.

The mean percentage of bacterial feeders (BF) from the five sampling sites were NP > OF > FF > MI > MO, where the two control NP and OF sites showed 72 and 60% bacterial feeders out of the total population, followed by about 50% in the soil obtained from the foliage management. Lower values for bacteria feeders and fungi feeders were obtained in the MO (organic farming). However, the highest percentage of omnivore-predator and plant parasite nematodes were present in the MO compared to the other managements.

3.1.2. Ecological Parameters

A total of nine ecological indices (Table 1) were used to assess management differences between the sites. The relative high variability in data obtained we had observed significant difference in the bacterivores between MI and NP ($p < 0.0053$) were no significant differences in FF, OP, and PP were obtained between the treatments. Moreover, as result of it no significant difference was found in H' (Shanon indices). From all the indices only the dominance (λ) that measure the information energy elucidating that only one of a few representatives' individuals in the genera that are found in high numbers will have more dominance. In this case the λ had been found to show significant difference between MO and MI ($p < 0.007$). Only two genera showed a specific signal (Figure 2) the *Aphelenchoïdes* in representing the MI treatment with a $p = 0.034$, and in the MO treatment the *Acrobeloides* were the $p = 0.079$ showing difference in comparison to the other four treatments.

Running a multilevel pattern analysis at a significant level (alpha) at level $p = 0.05$ – on the total number of 42 genera, the number of selected genera were found to be tow (2), were number of associate genera were two in the first group, in comparison to the other groups were 0 (zero). The group number one (MI treatment) in the present case were represented by *Aphelenchoïdes* $p = 0.0278$, and for MO treatment $p = 0.0329$.

In both above cases the two genera were found to determine the differences between the treatments.

Table 1. Mean for selected ecological indices of nematode community structure, in soil samples collected from the five managements system (MO – organic vineyard; MI – conventional vineyard; NP – natural pasture; FF – foliar fertilization; OF – Open Field).

	MO	MI	NP	FF	OF
BF	35.7	44.6	72.5	54.2	59.8
FF	10.6	42.4	14.2	13.3	10.3

OP	46.7	10.4	7.9	31.2	23.2
PP	7	2.7	5.4	1.3	6.6
FF/BF	0.2	1	0.2	0.4	0.4
WI	10.2	33.9	33	82	23.8
H'	1.7	1.5	1.5	1.5	1.3
TD	0.5	0.4	0.6	0.6	0.6
λ	0.2	0.3	0.4	0.4	0.5
ΣMI	3	1.9	1.6	2.2	2
J'	0.7	0.5	0.5	0.5	0.4

BF – bacteria feeder nematode relative abundance (as percentage); FF – fungi feeder nematode relative abundance (as percentage); OP – omnivore and predator nematode relative abundance (as percentage); PP – plant parasite nematode relative abundance (as percentage); FF/BF – fungi feeder and bacteria feeder nematode ratio; WI – Wasilewska index; H' – Shannon index; TD – trophic diversity index; λ – Simpson index; ΣMI – sigma maturity index and J' – Evenness of nematode assemblages.

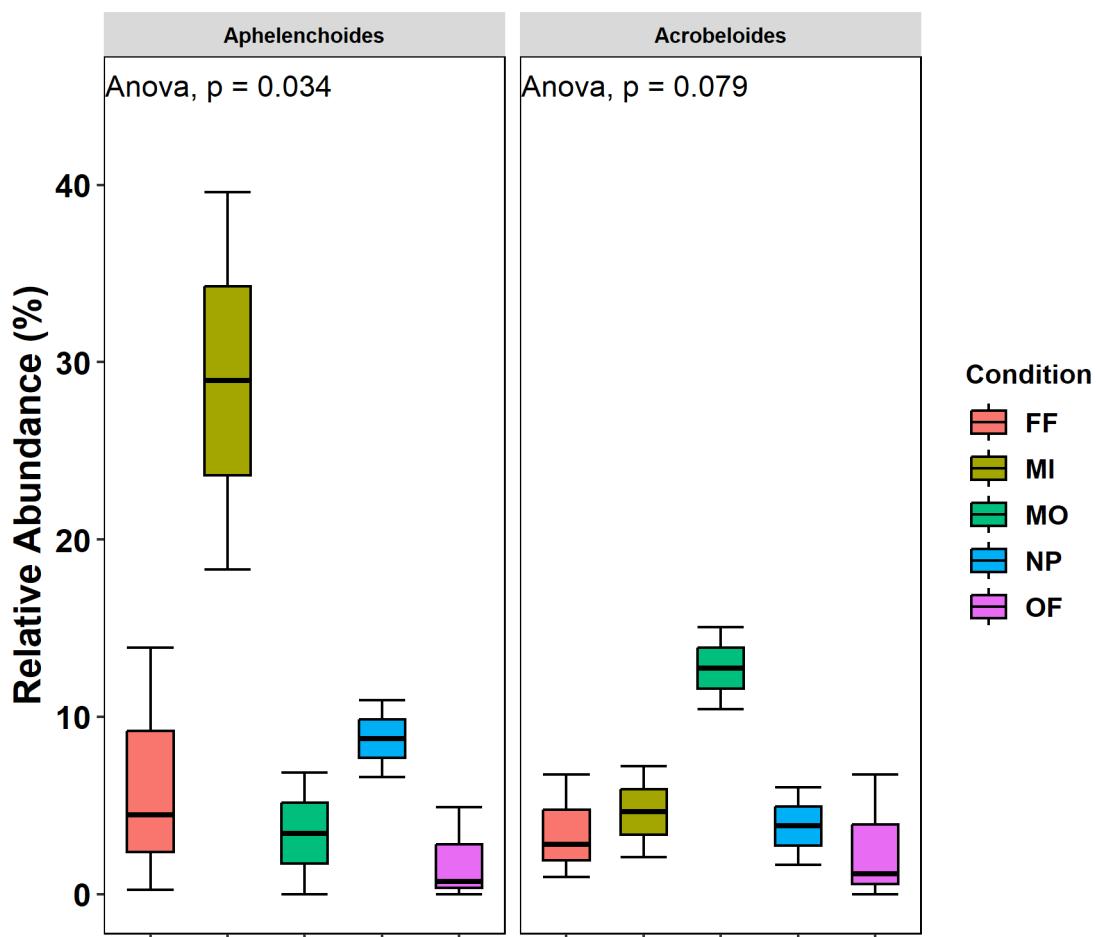


Figure 2. Relative abundance of the two genera *Aphelenchoides* and *Acrobeloides* showed specific signals one in MO and the second in MI treatments.

The Wasilewska index (WI) was significantly ($p < 0.05$) higher in the FF management compared to the other managements, except for the conventional (MI) samples. This shows the significant low number of plant parasite individuals in FF compared to MO managements. Moreover, the H' (Shannon index) showed a significant difference between the MO (organic management) and OF (open field – natural system), without any significant difference relative to the other managements. No significant differences between the managements were observed in trophic diversity (TD). The ΣMI values in the MO management reached a maximum value which was significantly higher compared to the other managements, elucidating a higher disturbance in the MO management.

compared to the other management sites. Evenness values indicated a higher abundance and greater biodiversity in the MO management relative to the other four managements, with the lowest value in the OF site.

3.1.3. Nematode Taxa

A total of six orders, 22 families, 42 genera and 89 species were detected in total from all soil samples collected from the five different managements. Eleven families were found to be dominant: Plectidae, Aporcelaimidae, Panagrolaimidae, Cephalobidae, Tylenchina, Qudsianematidae, Monhysteridae, Aphelenchoididae, Mononchidae, Anguinidae, and Aphelenchidae (Table 2).

A total of 25, 23, 27, 29 and 31 genera were present in the MO, MI, NP, FF and OF soil samples, respectively, from the total of 42 genera (Table 2). Eleven of the genera belonged to seven families of bacteria feeders, six genera belonged to 3 families of fungi feeders, six families represented, by 8 genera belonged to omnivores-predators and six families are represented by 17 genera belonging to plant parasites. The bacteria feeders genera that were present in all samples included *Acroboloides*, *Panagrolaimus* and *Plectus*. Out of the 6 fungi feeders genera found in our study, four were present in all managements: *Ditylenchus*, *Aphelenchus*, *Paraphelenchus*, and *Aphelenchoides*. The omnivores-predators genera present in all soil samples included *Aporceliamellus*, *Allodorylaimus* and *Thonus*. Five out of a total of 17 genera of plant parasites were present in all of the soil samples: *Anguina*, *Mothotylenchidae*, *Aglenchus*, *Basiria*, and *Tylenchus*.

Table 2. Generic composition, feeding group composition and relative abundance (RA) of Nematodes at each management.

Family	CP	Genera	MO	MI	NP	FF	OF
<i>Bacteria feeders</i>							
Alloionematidae	1	<i>Rhabditophanes</i>				+	
Cephalobidae	2	<i>Acrobeles</i>		+	+	+	
		<i>Acobeloides</i>	+++	++	++	++	++
		<i>Eucephalobus</i>			+	+	+
Monhysteridae	2	<i>Geomonhystera</i>	+	+	+++		+
Panagrolaimidae	1	<i>Panagrolaimus</i>	+++	+++	+++	+++	+++
		<i>Procephalobus</i>			+	+	
Plectidae	3	<i>Plectus</i>	+	+	+	++	+
Prismatolaimidae	3	<i>Prismatolaimus</i>	+				+
Rhabditidae	1	<i>Pellioiditis</i>				+	+
		<i>Protorhabditis</i>		+	+		
<i>Fungi feeders</i>							
Anguinidae	2	<i>Ditylenchus</i>	++	+++	++	++	++
Aphelenchidae	2	<i>Aphelenchus</i>	++	++	++	++	++
		<i>Paraphelenchus</i>	+	+	++	+	+
		<i>Aphelenchoides</i>	++	+++	++	++	++
Tylenchidae	2	<i>Boleodorus</i>					+
		<i>Dorylillum</i>					+
<i>Omnivores and predators</i>							
Aphelenchoididae	2	<i>Robustodorus</i>	+		+	+	+
Aporcelaimidae	5	<i>Aporceliamellus</i>	+++	+++	++	++	++
Mononchidae	4	<i>Prionchulus</i>			+	+++	+
Qudsianematidae	4	<i>Allodorylaimus</i>	++	+	++	+	++
		<i>Microdorylaimus</i>				+	
		<i>Thonus</i>	+++	+	++	++	++
Sphaerolaimoidea	3	<i>Daptonema</i>			+	+	
Tylenchidae	2	<i>Miculenchus</i>		+			+
<i>Plant parasites</i>							
Anguinidae	2	<i>Anguina</i>	+	+	+	+	+
Meloidogynidae	3	<i>Meloidogyne</i>					+
Merliniidae	2	<i>Geocenamus</i>	+		+	+	+

Nothotylenchidae	2	<i>Nothotylenchidae</i>	++	++	++	+	++
Pratylenchidae	2	<i>Hirschmanniella</i>	+	+			
		<i>Pratylenchus</i>	+			+	+
Tylenchidae	2	<i>Aglenchus</i>	+	+	+	+	+
		<i>Atetylenchus</i>	+		+		+
		<i>Basiria</i>	+	+	+	+	+
		<i>Cephalenchus</i>				+	
		<i>Coslenchus</i>					+
		<i>Discotylenchus</i>		+			
		<i>Filenchus</i>	+		+		+
		<i>Irantylenchus</i>	+	+		+	
		<i>Malenchus</i>					+
		<i>Tylenchus</i>	+	+	+	+	+
		<i>Paratylenchus</i>	+	+	+		

++ - >10%, ++ - >1%, + - >0%. MO – organic vineyard; MI – conventional vineyard; NP – natural pasture; FF – foliar fertilization; OF – Open field.

4. Discussion

In the recent years agricultural practices had overgone enormous changes trying to give answers to increasing demand for higher primary production, soil health maintenance, use of organic amendment more natural and favorable practices, that should go hand in hand with global requests. In order to do so the agroecosystem consequently try to develop novel approaches in developing and improving organic amendments. The effect of organic amendments from different sources had been reported to improve biological, microbiological, and biochemical soil characteristics such as water and nutrient holding capacity.

In the present study, the response of soil free living nematodes community, its density, functional composition, and additional attributes that are known to be good environmental indicators, were used to reveal the differences between long-term forms of managements (organic, conventional, natural pasture, foliage, and open field). Studies conducted by Werner and Dindal (Werner and Dindal, 1990) on conversion of organic agriculture on soil biota had shown that soil nematodes were the most abundant in organic plots. Our results had been found to demonstrate that organic farming contribute in improving the density of microbial feeding that tend to improve nutrient resource availability. As being one of the primary grazers of saprophytic community the nematodes increase nutrient availability to primary producers (Matlack, 2001). The changes in nematodes community features reflect the changes in the below-ground soil milieu, including the below-ground plant rhizosphere biosphere (Bongers and Ferris, 1999; Fitoussi et al., 2016; Neher et al., 2005). Nematode feeding groups indices elucidate the impact of the different managements on accessibility and immediate utilization of food sources (bacteria and fungi) (Yeates et al., 1993). Although the determined nematode indices cannot help in understanding or differentiating between management performances in all cases, we should prolong the use the nematode community food web structure as proposed by Martin et al. (Martin et al., 2022). This will increase our understanding of scale row-crop agriculture.

Manipulating managements in vineyards to reduce disease and improve yields is a popular and accepted practice. Various studies had tried to link cover crops and the soil nematode community – both beneficial and pathogenic – in order to enhance the control over parasitic nematodes. Our results indicated differences in soil food web function between managements and shifting between the three main feeding groups (bacterivores, fungivores, and omnivores-predators) were found to be according to resource availability. Soil organic matter and organic residues are supporting fungi development which in turn is followed by fungivore nematodes abundance that were showed high presence of omnivores and predators community development. These findings are similar to those reported by Ferris (Ferris, 2010; Ferris and Bongers, 2006) and Ugarte et al. (2013). Nematode indices which are developed based on the various functional guilds (Ferris and Bongers, 2006) were

inconsistent comparing the managements, except the Wasilevska index which showed a sharp increase in foliage management, thus elucidating the management effect on plant parasites.

Plant endoparasitic or ectoparasitic nematodes present in grapevine are having a strong effect on plant growth and virus transmission which are widespread all over the globe without any exception of geographical region or soil textures (Aballay et al., 2009). Plant parasitic nematode families followed by their genera were found to vary between the five managements. In the Pratylenchidae family, the genus *Pratylenchus* was present in organic-related management (MO, FF) and in the open field (OF) that can be attributed to development of primary production along time (Coll et al., 2012). It is known as a root lesion and pin nematode, pathogen of grapevine as reported by (Zasada et al., 2012) and (Howland et al., 2014). The Nothotylenchidae family, represented by genus *Nothatylenchidae*, was present in relatively high numbers in all managements. The Nothotylenchidae (Thorne, 1941) is a redescription of *Tylenchorhynchus* (Elmiligy, 1955) and is known to be a pathogen present in the vicinity of clover roots (*Trifolium alexandrinum* L.), guava (*Pisidium guave* L.), squash (*Cucurbita moschata* Duch.), turnip (*Brassica rapa* L.), and near grapevine based on our results from the present study. They can also predispose roots to invasion by other plant pathogens. The main damage of plant pathogen nematodes is in a disturbance of nutrient and water uptake, thus decreasing and yield.

Soil free living plant-parasitic nematodes are very commonly found in vineyards (Schlüter et al., 2022; Zasada et al., 2012). Permanent grass cover, different types of managements, soil aggregate and pore size distribution affect the community structure, diversity, and abundance (Ferris et al., 2001; Neher, 2001; van den Hoogen et al., 2019). Long-term organic, intercrops (grass cover) negatively influence diversity, while positively affecting omnivore-predators that are sensitive to disturbance determined by habitat constraints related to resource availability. Studies of plant parasitic nematodes had showed that competition with arbuscular mycorrhizal and endophytic fungi and predator nematodes can control their community size.

The present study has important implications for understanding the effect of different management types on the soil free living nematode community composition (Siddiqui and Mahmood, 1999). This study emphasized the importance of the bacterial and fungal community that can be used as a front line against pathogens such as nematodes. As the plant rhizosphere determine habitat structure the different management contribute to the physical soil pore structure, moisture availability and resource distribution. The new and the different habitat's structure promoted by the different agro-management stimulate new conglomerates of soil biota composition. This supports our findings that organic managements suppress most parasitic nematode similar to the finding reported by McSorley (2011). These results provide a new perspective on a time axis, where a deeper study to link grapevine plant phenology, soil biota dynamics with emphasis on soil nematode community and its functionality, and physico-chemical components are needed to evaluate and avoid the impact of these pests on vineyard productivity.

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