SHORT NOTE



Stable C and N isotope ratios reveal soil food web structure and identify the nematode *Eudorylaimus antarcticus* as an omnivore–predator in Taylor Valley, Antarctica

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Abstract

Soil food webs of the McMurdo Dry Valleys, Antarctica are simple. These include primary trophic levels of mosses, algae, cyanobacteria, bacteria, archaea, and fungi, and their protozoan and metazoan consumers (including relatively few species of nematodes, tardigrades, rotifers, and microarthropods). These biota are patchily distributed across the landscape, with greatest faunal biodiversity associated with wet soil. Understanding trophic structure is critical to studies of biotic interactions and distribution; yet, McMurdo Dry Valley soil food web structure has been inferred from limited laboratory culturing and microscopic observations. To address this, we measured stable isotope natural abundance ratios of C (¹³C/¹²C) and N (¹⁵N/¹⁴N) for different metazoan taxa (using whole body biomass) to determine soil food web structure in Taylor Valley, Antarctica. Nitrogen isotopes were most useful in differentiating trophic levels because they fractionated predictably at higher trophic levels. Using ¹⁵N/¹⁴N, we found that three trophic levels were present in wet soil habitats. While cyanobacterial mats were the primary trophic level, the nematode *Plectus murrayi*, tardigrade *Acutuncus antarcticus*, and rotifers composed a secondary trophic level of grazers. *Eudorylaimus antarcticus* had a ¹⁵N/¹⁴N ratio that was 2–4‰ higher than that of grazers, indicating that this species is the sole member of a tertiary trophic level. Understanding the trophic positions of soil fauna is critical to predictions of current and future species interactions and their distributions for the McMurdo Dry Valleys, Antarctica.

Keywords Dry Valleys · Predator · Trophic levels · Isotopic fractionation · Feeding ecology · Connectivity

Introduction

The McMurdo Dry Valleys, Antarctica are an extreme ecosystem: they are very cold, dry, and windy, their soils have high pH, low moisture, and low organic carbon, and are

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often saline (Campbell et al. 1998; Fountain et al. 1999; Burkins et al. 2001). These are simple ecosystems compared to temperate systems: they lack vascular plants, soil fauna biodiversity is 1.1–2.6% of temperate soils, and approximately 30-40% of studied soils lack soil fauna (Freckman and Virginia 1997). While the presence of soil cyanobacteria, microbes, nematodes, rotifers, tardigrades, collembolans, and mites has been recorded in locations throughout the dry valleys, their distributions are patchy, with greatest invertebrate diversity in wetted lake and stream margins (Freckman and Virginia 1997; Adams et al. 2014). For example, soil invertebrates found in Taylor Valley include four species of nematode: Scottnema lindsayae, Plectus murrayi, Geomonhystera antarcticola, and Eudorylaimus antarcticus, a tardigrade, Acutuncus antarcticus, and rotifers. Of these, S. lindsayae is the only taxon whose most suitable habitat is the dry soil (e.g., Freckman and Virginia 1997; Treonis et al. 1999; Adams et al. 2006; Ayres et al. 2007) that makes up > 95\% of the landscape. The low biodiversity in both wet and dry soils makes each trophic link significant, due to the



lower redundancy in function compared to more temperate soils (Freckman and Virginia 1997). However, in situ observation of feeding is difficult due to the opaque nature of soil and the small size of these fauna. The current understanding of dry valley soil trophic structure is limited to laboratory studies and microscopic observations.

Some Taylor Valley soil fauna have been grown in the laboratory, helping to identify their feeding ecology. For example, P. murrayi feeds on bacteria in laboratory cultures (Adhikari et al. 2010; de Tomasel et al. 2013), while the tardigrade A. antarcticus feeds on algae (Cecilia Milano de Tomasel, personal communication). Additionally, S. lindsayae is a known microbivore, feeding on yeast and bacteria in a laboratory study (Overhoff et al. 1993). Furthermore, mouthparts are often used to identify nematode trophic groups under microscopic observation (Yeates et al. 1993). While P. murrayi, S. lindsayae, and G. antarcticola all possess the tube-like esophagus of typical bacterivore nematodes, E. antarcticus bears an odontostylet—a piercing structure inside its mouth which can puncture food, such as plant or animal tissue. Previous studies predicted that E. antarcticus was a likely omnivore-predator in the dry valleys (Wall and Virginia 1999) due to its mouthparts and the trophic classification of the genus Eudorylaimus as an omnivore-predator in temperate habitats (Yeates et al. 1993). Other Antarctic studies hypothesized that soil algae was the only food source for *E. antarcticus* (Powers et al. 1998). More recently, E. antarcticus was revealed to be an algal feeder through observation of chlorophyll in the intestine using a fluorescent microscope and acridine orange filter (Wall 2007). While other locations in Antarctica have predaceous taxa (Sohlenius and Boström 2005), no studies to date have confirmed a metazoan predator in Taylor Valley.

The nematode genus *Eudorylaimus* is widespread in soils (McSorley 2012). This odontostylet-bearing group is considered omnivorous or predaceous in temperate ecosystems (Yeates et al. 1993; McSorley 2012; Stirling 2014). As early as 1929, Cobb observed Eudorylaimus feeding on mite eggs. Additionally, *Eudorylaimus* in laboratory culture preys on nematodes, such as the bacterivore genera Acrobeloides, Plectus, and Panagrellus (Tjepkema et al. 1971; Ferris and Ferris 1989), but its food sources are diverse: it also feeds on algae, enchytraeids, fungi, protozoa, and mites, but with reproduction only observed on nematodes, algae, moss, and protozoa (Hollis 1957; Wood 1973; Ferris and Ferris 1989). Its diverse feeding strategy and occupation of a top trophic position may make it an important driver of food web structure in temperate climates. However, the high taxonomic and functional biodiversity in soil makes these relationships difficult to discern.

The natural abundance of stable isotopes fractionate predictably up trophic levels at approximately +3-4% for $\delta^{15}N$ and +0.5-1% for $\delta^{13}C$ (Post 2002; McCutchan et al. 2003).

Previous stable isotope studies showed that nematode orders and families differ in their trophic structure (Kudrin et al. 2015). Eudorylaimus antarcticus belongs to the order Dorylaimida, and Kudrin et al. (2015) found this order to have a similar isotopic composition to predaceous nematodes in boreal forests. However, in hot desert soils, Dorylaimida have isotopic composition similar to bacterivore nematodes, probably due to feeding on cyanobacteria (Darby and Neher 2012). Following this logic, we used natural abundance of stable isotopes to characterize the trophic levels for the three most abundant nematode species in Taylor Valley (E. antarcticus, P. murrayi, and S. lindsayae), the tardigrade A. antarcticus, and rotifers (grouped at phylum level). We did not include G. antarcticola in this study because it is an extremely rare species, and we did not find it at our study sites. We hypothesized that the grazers P. murrayi, S. lindsayae, A. antarcticus, and rotifers would be 3–4% for $\delta^{15}N$ and around + 0.5–1% for δ^{13} C higher than the microbial mat values. We also expected that E. antarcticus would be 3-4% for $\delta^{15}N$ and around +0.5-1% for $\delta^{13}C$ higher than P. murrayi, S. lindsayae, A. antarcticus, and rotifers.

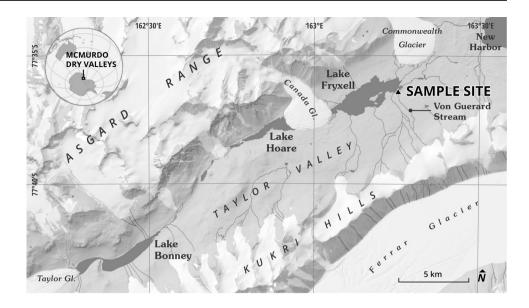
Methods

To test our hypothesis, we chose a well-studied site in Taylor Valley (e.g., Spaulding and McKnight 1998; Treonis et al. 1999) near Von Guerard stream in the Lake Fryxell basin of Taylor Valley (77.608 S, 163.254 E). In January 2014, six soil samples were collected near Von Guerard stream (Fig. 1). Three random soil samples (sites 1, 2, 3) were collected in dry soil more than 5 m from the stream (but not more than 10 m) and three random soil samples (sites 4, 5, 6) were collected within 5 m of the stream (within wetted margin, but not within stream sediments). Sites 1-3 were considered 'dry' soil replicates and sites 4-6 were considered 'wet' soil replicates. These samples were collected into Whirl-Pak® bags to 10 cm using a clean plastic scoop. Approximately 500 g of soil was collected per sample. The soils were stored in an ice chest and transported to the Crary Laboratory at McMurdo Station (United States Antarctic Program) where they were slowly cooled from +4to -20 °C over 4 days (i.e., 24 h at +4 °C, 24 h at -4 °C, 24 h at -10 °C, and 24 h at -20 °C). Samples were shipped frozen (- 20 °C) to Colorado State University, Fort Collins, Colorado, USA, and slowly defrosted (as described by De Tomasel et al. 2013) before extracting soil fauna.

For each of the six sites, fauna were extracted from 100 g of soil via cold sugar centrifugation technique (Freckman and Virginia 1993). Within 48 h of extraction, taxonomic groups were identified (nematodes and tardigrades identified to species and rotifers to phylum, Olympus CKX41, 200X magnification) and counted. An additional 50-g subsample



Fig. 1 Location of our sample site near Von Guerard stream in the Lake Fryxell basin in Taylor Valley, Antarctica. Taylor Valley is located in the McMurdo Dry Valleys of Southern Victoria Land, Antarctica (*inset*). Map by Brad Herried, Polar Geospatial Center



of soil was used to determine gravimetric soil moisture (water mass per unit soil mass) by mass loss from soils dried at 105 °C for 48 h. The soil fauna counts were then corrected for soil moisture content and expressed as kg⁻¹ dry soil. After identification and enumeration, each group was separated and collected into tin capsules $(8 \times 5 \text{ mm})$ Elemental Microanalysis BN/170056) with an eyelash tool (Superfine eyelash with handle, Ted Pella, Inc., Prod no. 113) under a dissecting microscope (Olympus SZX10, 30X magnification) as per the method described by Shaw et al. (2016). A minimum of 0.02 mg biomass dry weight (approximately 50-100 live individuals) was collected for each taxon. For each taxon identified at each site, we aimed to triplicate fauna collections for isotope analysis. Extractions from 100 g of soil were repeated by site until sufficient biomass was achieved for each group found at that site. For this study, it was possible to collect eight total replicates for E. antarcticus (three replicates from sites 4, 5, and two replicates from site 6), nine total replicates for P. murrayi (three replicates from each of the sites 4, 5, and 6), five total replicates for S. lindsayae (two replicates from site 2, and three replicates from site 3), nine total replicates for A. antarcticus (three replicates from each of the sites 4, 5, and 6), three replicates for rotifers (one replicate from each of the sites 4, 5, and 6), and nine total replicates for the cyanobacterial mat (three replicates from each of the sites 4, 5, and 6). Samples were dried for 48 h in a desiccator prior to shipment to the Stable Isotope Mass Spectrometry Laboratory (SIMSL) at Kansas State University for isotope analysis (CE-1110 EA coupled via Conflo II interface to an IRMS, Thermo Finnigan DeltaPlus).

We also extracted *P. murrayi* from cultures grown on Bold's Modified Basal Freshwater Nutrient Media with Ottawa Sand (Adhikari et al. 2010) by both modified

Baermann funnel technique, which uses water only (Hooper 1970), and the sugar centrifugation technique to check for any effect of sugar (which the fauna are submerged in for ~ 2 min) on the carbon isotope composition of the animals. We used 3 replicates for each extraction technique. Our tests revealed that there was no significant difference (p = 0.25, df = 4, unpaired t test) between cultured *P. murrayi* extracted via Baermann funnel or sugar centrifugation, which had δ^{13} C isotopic compositions of -17.81 ± 0.76 and -19.26 ± 0.77 (Mean \pm SE), respectively.

Results and discussion

Across the six sites near Von Guerard stream, the nematodes *S. lindsayae*, *P. murrayi*, *E. antarcticus*, tardigrade *A. antarcticus*, and rotifers were found. However, their distributions were not equal across all sites. Wet sites had soil moistures of $17.91 \pm 0.29\%$; all groups were found in these samples but *S. lindsayae* was in very low abundance compared to dry sites (e.g., sites 1-3; Table 1). Dry sites had gravimetric soil moistures of $0.70 \pm 0.12\%$; only *S. lindsayae* was found in these samples (Table 1). This is unsurprising; many studies have found *S. lindsayae* dominating dry habitat, while other taxa prefer wetter habitats in the dry valleys (e.g., Freckman and Virginia 1997; Treonis et al. 1999; Adams et al. 2006; Ayres et al. 2007).

The stable isotope composition of soil fauna biomass revealed a wet soil food web with three trophic levels: a basal level of cyanobacterial mat, a secondary level of rotifers, tardigrade *A. antarcticus*, nematode *P. murrayi*, and a tertiary level occupied by nematode *E. antarcticus* (Fig. 2). The primary trophic level had an overall isotopic composition of δ^{15} N -5.36 \pm 0.96‰, δ^{13} C -24.99 \pm 0.63‰. These



Table 1 Abundance of taxonomic groups from Taylor Valley soil samples in total live individuals kg⁻¹ dry soil

Taxonomic group	Overall abundance	Dry soil abundance	Wet soil abundance
Nematoda			
Scottnema lindsayae	1822.98 ± 213.22	533.07 ± 402.76	54.59 ± 054.59
Plectus murrayi	300.23 ± 172.73	0.00 ± 0.00	600.45 ± 243.00
Eudorylaimus antarcticus	1325.61 ± 640.76	0.00 ± 0.00	2651.22 ± 543.74
Tardigrada			
Acutuncus antarcticus	545.67 ± 287.81	0.00 ± 0.00	1091.33 ± 341.23
Rotifera			
	1299.48 ± 667.53	0.00 ± 0.00	2598.95 ± 734.38

Data are shown as mean \pm standard error. Dry soil (n=3) corresponds to samples 1, 2, and 3, which were > 5 m from the stream and had soil moisture < 1% (gravimetric). Wet soil (n=3) corresponds to samples 4, 5, and 6, which were < 5 m from the stream and had soil moisture > 17% (gravimetric)

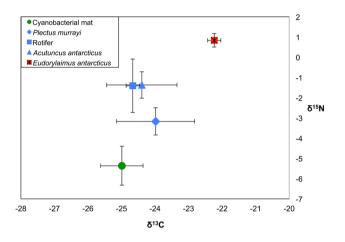
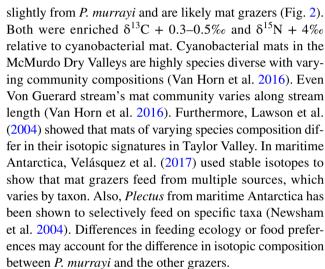


Fig. 2 δ ¹³C and δ ¹⁵N signature of fauna groups from wet soil within 5 m of Von Guerard stream in Taylor Valley, Antarctica. Data presented are % and are means with standard error bars

cyanobacterial mats from the Von Guerard stream margin have a similar isotopic ratio to the organic matter measured by Lawson et al. (2004) for streams in Taylor Valley, Antarctica. While these mats are composed of multiple groups including cyanobacteria and diatoms, which have differing isotopic signatures (Lawson et al. 2004; Velázquez et al. 2017), our analysis is limited to composite mat samples.

The secondary trophic level at the wet sites was composed of rotifers, tardigrades, and the nematode *P. murrayi*. Generally, the trophic position of culturable groups, such as *P. murrayi* and *A. antarcticus* as revealed by stable isotopes, corresponds to the established understanding of their feeding habits. The nematode, *P. murrayi*, was enriched in δ^{13} C relative to cyanobacterial mat (~ + 1‰), but only about + 2‰ for δ^{15} N relative to the cyanobacterial mats (Fig. 2). Other studies have shown that the isotopic composition of Plectidae from boreal forests also supports its trophic position as a bacterivore (Kudrin et al. 2015). The rotifers and tardigrades had similar isotopic compositions to each other, but differed



The tertiary trophic level was occupied solely by *E. antarcticus*. This nematode had an isotopic composition of $\delta^{15}N$ 0.84 \pm 0.44%, $\delta^{13}C$ -22.23 \pm 0.25% (Fig. 2). This is an enrichment of + 2.25% relative to rotifers and tardigrades and + 4% relative to *P. murrayi*. Nematodes of the order Dorylaimida are usually considered omnivore–predator, but have also been shown to eat bacteria, fungi, algae, and plants (Freckman 1988; Yeates et al. 1993; Kudrin et al. 2015). Protozoa may also be a possible food source for nematodes and, although present in dry valley soil (Bamforth et al. 2005), were not included in this study due to difficulty in isolating individuals and collecting sufficient biomass for isotope measurement. Given our isotope results and that *E. antarcticus* also eats algae (Wall 2007), it should be described as an omnivore–predator for the McMurdo Dry Valleys.

The dry soil food web had the lone consumer *S. lind-sayae*, which is a microbivore nematode (Overhoff et al. 1993). *Scottnema lindsayae* had an isotopic composition of δ^{15} N -4.18 \pm 1.05%, δ^{13} C -27.91 \pm 0.39% (Fig. 3), which is considerably lighter than consumers in the wet food web (Fig. 2), making the wet soil's cyanobacterial mats an unlikely primary level for dry soil. Other studies



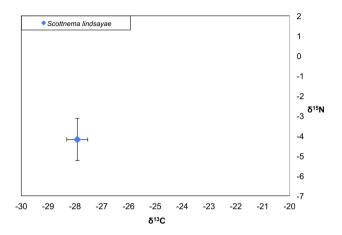


Fig. 3 δ ¹³C and δ ¹⁵N signature of fauna groups from dry soil 5–10 m from Von Guerard stream in Taylor Valley, Antarctica. Data presented are % and are means with standard error bars

have suggested that lithic primary producers (e.g., cryptoendoliths, hypoliths) could be a carbon source in dry soils (Burkins et al. 2000), and these sources have a lower isotopic signature than lake- and stream-associated cyanobacterial mats (Burkins et al. 2000). Additionally, we did not find *E. antarcticus* in the dry soil sites, but it is occasionally found in dry soil in the dry valleys (Freckman and Virginia 1997). The most frequent two-species community in the dry valleys is *S. lindsayae* with *E. antarcticus* (Freckman and Virginia 1997). Further work is needed to confirm trophic positions of soil fauna across the heterogeneous dry valleys landscape, especially in habitats with varied carbon sources (Burkins et al. 2000) and communities of multiple invertebrate species.

Our results have ecosystem implications. Around 95% of the McMurdo Dry Valleys region is considered dry soil habitat ($\leq 2\%$ gravimetric water content), which is most suitable for the dominant nematode, bacterivore S. lindsayae. While E. antarcticus are present in low abundance in dry soil habitat, many are found in moist or wet soils. Around 5% of the McMurdo Dry Valleys are considered moist soil habitat, which is most suitable for E. antarcticus (Virginia and Wall 1999; Burkins et al. 2001). The McMurdo Dry Valleys have recently been termed "a landscape on the threshold of change" because significant increases in the loss of both glacier and buried ice are expected to occur (Fountain et al. 2014). This predicted future increase in glacial melt and permafrost thaw will produce larger amounts of liquid water during the austral summer, generally "wetting up" what is now dry landscape (Gooseff et al. 2017a). With this increased melt from glaciers, massive buried ice, and permafrost, moist habitat area should increase. Such a change could impact the distribution and abundance of soil fauna, which differ in their habitat preferences (Freckman and Virginia 1997), altering community composition (Nielsen et al.

2011), and ultimately having ecosystem-level impacts on soil processes including carbon cycling (Barrett et al. 2008; Gooseff et al. 2017b). Understanding trophic positions and biotic interactions of soil fauna is critical for predicting future changes in species distributions and interactions due to increased connectivity.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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