RESEARCH ARTICLE

Extremely high abundances of *Prasiola crispa*-associated micrometazoans in East Antarctica

Dzmitry A. Lukashanets,[1,2] Yury H. Hihiniak[3] & Vladislav Y. Miamin[3,4]

[1]Marine Research Institute, Klaipėda University, Klaipėda, Lithuania
[2]Institute of Oceanology, Polish Academy of Sciences, Sopot, Poland
[3]State Scientific and Practical Center for Bioresources, National Academy of Sciences of Belarus, Minsk, Belarus
[4]Biology Faculty, Belarusian State University, Minsk, Belarus

Abstract

To elucidate poorly known aspects of the microscopic metazoan distribution in ice-free parts of the Antarctic, we examined samples of the multicellular terrestrial alga *Prasiola crispa*, collected over the last decade in different parts of continental East Antarctica and Haswell Island. We found that the micrometazoans inhabiting the algae consist of remarkably abundant bdelloid rotifers (subclass Bdelloidea), followed by tardigrades. We did not find nematodes. The rotifer assemblages were characterized by low diversity (only six species). Nevertheless, rotifer densities were extremely high: mean densities ranged from 75 to 3030 individuals per 100 mg of the dry sample weight and the maximum value numbered in excess of 8000 per 100 mg of the dry sample weight. These data show that terrestrial algae, along with mosses, are a very attractive habitat for rotifers and tardigrades in the Antarctic. The statistical analysis showed a lack of correlations between rotifer and tardigrade densities and nutrients (N, C, P, K and Na). Our findings are consistent with the patchy distribution of terrestrial micrometazoans in the Antarctic that has previously been found.

To access the supplementary material, please visit the article landing page

Introduction

Terrestrial life in the Antarctic is concentrated in the few ice-free areas; is well adapted to extremely low temperatures, strong winds, freeze–thaw cycling and nutrient deficiency (Convey 1996; Rogers et al. 2012); is highly endemic (Pugh & Convey 2008; Convey et al. 2020) and has a long evolutionary history (Convey & Stevens 2007; Convey et al. 2008; Fraser et al. 2012; Convey et al. 2020). Microscopic metazoans, which include rotifers, nematodes and tardigrades, are important components of terrestrial species assemblages and inhabit both soils, the so-called 'chalikosystems' proposed by Janetschek (1963, 1967), and microbiotopes, forming in moss cushions, lichen and algal thalli (the so-called 'bryosystems'). Simple intraspecific interactions and low taxonomic diversity allow for the consideration of terrestrial micrometazoan assemblages as models for ecology studies (Adams et al. 2006; Wall et al.

2006; Heatwole & Miller 2019). This is especially important in the context of climate change and anthropogenic impact, both of which are dramatically affecting the components of Antarctic biota (Convey 2010; Chown et al. 2012; Convey & Peck 2019).

The Antarctic terrestrial biota is commonly considered to be relatively poorly known (Convey 2010; Convey et al. 2014). Nevertheless, a number of genetic studies of micrometazoans have enhanced our knowledge of the diversity of particular groups (Velasco-Castrillón, Page et al. 2014; Velasco-Castrillón & Stevens 2014; Velasco-Castrillón et al. 2015) and have revised earlier published data (Iakovenko et al. 2015; Velasco-Castrillón et al. 2018). Other publications pertaining to micrometazoan diversity have a more global basis and describe general patterns (e.g., Fontaneto et al. 2015), with some offering a review of the literature (e.g., Adams et al. 2006; Velasco-Castrillón, Gibson et al. 2014) or a data synopsis (Garlasché et al. 2020).

The ecological patterns of Antarctic micrometazoans are characterized by great variability and inconsistency. Serial surveys conducted in both the chaliko- and bryosystems of Dronning Maud Land found no (Sohlenius et al. 1995; Sohlenius et al. 2004) or very little (Sohlenius et al. 1996; Sohlenius & Boström 2005) correlation between the presence of large micrometazoan taxa and biotic and abiotic factors. This has led to the conclusion that the micrometazoan distribution in Antarctica is largely a result of colonization processes and historical factors, apart from the environmental conditions (Sohlenius & Boström 2008). However, in a more recent study, micrometazoans in the Larsemann Hills (Heatwole & Miller 2019) were found to have habitat preferences, and some groups of micro-invertebrates were strongly associated with one another in terms of abundance and diversity. Studies of the microbial mats of maritime Antarctica (Velázquez et al. 2017; Almela et al. 2019) and the soils of the McMurdo Dry Valleys (Shaw et al. 2018) have shed light on the general outlines of the trophic interactions and nutrient fluxes among the main micrometazoan taxa. Moreover, another recent study has found, in contrast to earlier studies, that the nematode occurrence in soils in this region is determined by biotic interactions (Caruso et al. 2019) with geochemistry and altitude (Zawierucha et al. 2019) and by moisture (Andriuzzi et al. 2018). Soil chemistry and moisture were correlated with the diversity and distribution of micrometazoans at Edmonson Point, where a predominance of rotifers and their species-specific habitat preferences were reported (Smykla et al. 2018). In general, the distribution of soil microfauna in the Antarctic is patchy and appears to be determined by soil chemistry rather than geography (Velasco-Castrillón, Schultz et al. 2014). The sometimes contradictory findings across different habitats and regions underscore the necessity of further investigations.

In our study, reported herein, we focused on terrestrial algae, which are among the substrates that have not been well studied in connection with micrometazoans. We examined samples of the commonly found green macroalga, *Prasiola crispa* (Lightfoot) Kützing 1843, collected during the 2011–19 summer seasons at different sites in continental East Antarctica and on Haswell Island. We observed extremely abundant micrometazoans (rotifers first and then tardigrades). We consider several issues here: (1) the structure of the micrometazoan community in the algae; (2) how habitat characteristics appear to be linked with the high abundance of inhabiting animals, especially the rotifers; (3) a comparison of the observed densities and diversity with those of other Antarctic regions and types of habitats.

Materials and methods

Characteristics of study material

The study material included lumps of the terrestrial algae, *Prasiola crispa*, along with the underlying ground substrate, from different parts of East Antarctica (Fig. 1). We collected samples at three sites in the Thala Hills, in Enderby Land, near the Belarusian station Vechernyaya Mount. We also collected samples at two sites in the Larsemann Hills, near the Russian station Progress. Algae were collected manually during the 2012/13, 2016/17 and 2018/19 summers (Table 1). Algal growth in a variety of settings was sampled, including ground depressions, crevices among rocks and, in the case of sites in the Larsemann Hills, pond-like hollows (Fig. 2). Between seven and 24 samples, each about 2.0 cm³ in size, were collected at sites in different parts of the algal growth. We also examined older material, which had been collected at one site on Haswell Island and one site in the vicinity of the Druzhnaya Russian field station by Sandefjord Bay during the 2010/11 season. Old photographs and oral descriptions revealed that these sites strongly resembled those from which samples had been collected between 2012 and 2019. Seven samples of the 2010/11 specimens from near Druzhaya and 24 samples from the 2010/11 specimens collected on Haswell Island were examined. These older samples were similar in size to the samples collected between 2012 and 2019. Algae were stored frozen (collected in 2016/17 and 2018/19) or dried (collected in 2012/13 and the older specimens, collected in 2010/11), without being chemically preserved. All materials have been deposited in the collections of the Scientific and Practical Centre for Bioresources (Belarus).

Sample processing

Before analysis, all stones, large sand particles, penguin feathers and other inclusions were removed from the samples. The concentrations of the main elements—the different nutrients—were measured in the samples for each site (for details, see the Supplementary material).

After hydrating the samples, we observed numerous individuals of the phyla Rotifera and Tardigrada. To count all the individuals, we used the method suggested by Peters et al. (1993) for the extraction of rotifers from mosses, which constitutes multiple washings (up to 10) and counting the number of animals in each washing. For each washing, we shook the samples vigorously in a 50-ml vial with 20 ml of distilled water. This procedure allowed for the almost complete disintegration of the sample, making all metazoans visible (Fig. 3).
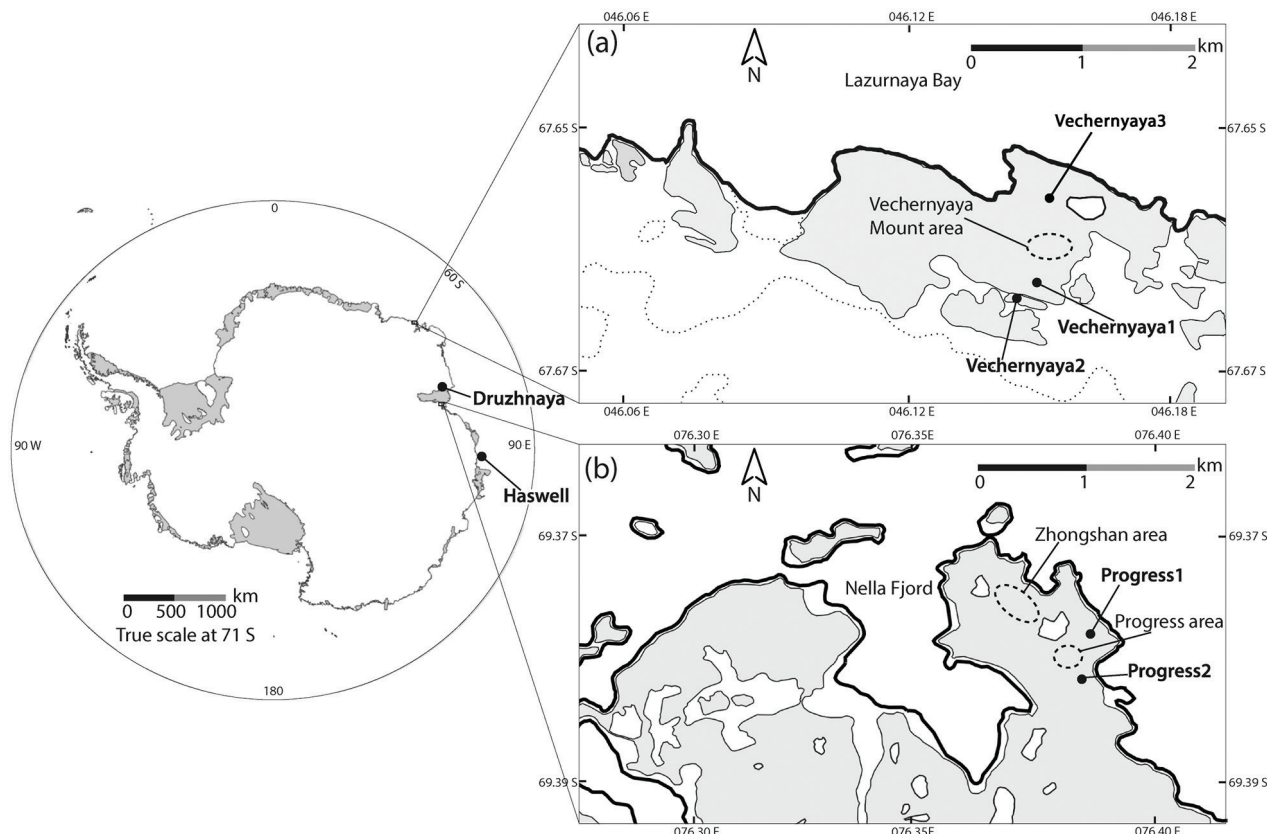
Fig. 1 Location of sampling sites: (a) Thala Hills (near the Belarusian Antarctic Station, Vechernyaya Mount) and (b) Larsemann Hills (near the Russian Antarctic Station, Progress).

Table 1 Information about sampling sites.

No.	Site	Region	Coordinates (decimal)	Altitude (m a.s.l.)	No. of samples	Sampling date	Collected by
1	Haswell	Haswell Island	No exact data	No exact data	24	Jan 2011	Y. Hihiniak
2	Druzhnaya	Sanderfjord Bay	No exact data	No exact data	7	Jan 2011	Y. Hihiniak
3	Vechernyaya1	Thala Hills	S 67.65972 E 046.14619	147	8	Jan 2012	V. Miamin
4	Vechernyaya2	Thala Hills	S 67.66117 E 046.14148	109	7	Dec 2018	D. Lukashanets
5	Vechernyaya3	Thala Hills	S 67.65577 E 046.15339	42	12	Jan 2019	D. Lukashanets
6	Progress1	Larsemann Hills	S 69.37593 E 076.39262	7	24	Feb 2017	Y. Hihiniak
7	Progress2	Larsemann Hills	S 69.38041 E 076.39259	12	20	Feb 2017	Y. Hihiniak

Abundance values, which had been estimated by the simple counting of all animals in the Petri dish for each washing of the sample of known weight, were then recalculated as densities in ind·0.1 gdw^{-1}. Although recalculating to the area each sample could cover is more conventional for quantitative estimates of microscopic animals, we preferred the 'per weight' approach, largely because of the complexity of the *Prasiola crispa* thallus. Even differences of a few millimetres in algal thickness can result in a significant increase in the animal number,

making it very complicated to standardize the sample by its square.

One hundred revived rotifer individuals were chosen randomly and were then transferred to slides and examined using light microscopy (Nikon and Micros). These living, active animals were examined with microphotography (NIS Elements, Microscopy Vision software). We used the most convenient keys (Donner 1965; Kutikova 2005) and articles that contain information regarding Antarctic bdelloid identification (Iakovenko et al. 2015). Additionally,

Fig. 2 Photographs of sampling sites: (a) Vechernyaya1; (b) Vechernyaya2; (c) Progress1; (d) Progress2.
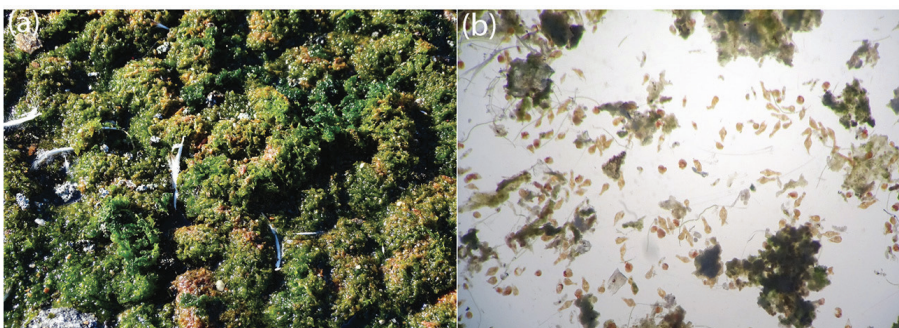

Fig. 3 Sample processing: (a) close-up view of *Prasiola crispa* and (b) bdelloid individuals in the hydrated sample.

we used the morphometry analysis suggested by Iakovenko et al. (2013) and the DNA sequencing of the COXI mitochondrial gene (Supplementary Table S3). Tardigrades, initially not the focus of the research (many of them were in an unidentifiable state), were studied only superficially and were identified to higher taxa using the appropriate keys (Ramazzotti & Maucci 1983; Pilato & Binda 2010; Bingemer & Hohberg 2017).

Data analysis

Spearman correlation tests of rotifer and tardigrade density values against nutrient content and rotifer density values against tardigrade density values were performed using R statistics (R-Studio integrated development environment) and PAST software for scientific data analysis.

Results

Sample characteristics

All samples comprised algal thalli with particles of the hardly visible ground and extraneous organic particles, which remained after primary processing. The samples varied significantly in terms of the Na, P and K contents, whereas N and C concentrations did not differ much (Table 2).

Taxonomic diversity

The animals in the algal samples included bdelloid rotifers (Rotifera: Bdelloidea) and tardigrades (Tardigrada). Nematodes, the third main group of the micrometazoan community in the Antarctic, were absent.

Table 2 Nutrient concentrations at sampling sites.

No.	Site	Na (g/kg)	P (g/kg)	K (g/kg)	N (%)	C (%)
1	Haswell	59.47	8.56	86.81	3.32	31.35
2	Druzhnaya	155.95	14.736	105.61	3.35	n/d
3	Vechernyaya1	34.32	9.78	194.83	3.68	39.98
4	Vechernyaya2	56.41	17,25	200.53	2.08	25.81
5	Vechernyaya3	60.64	10.13	118.28	2.80	28.23
6	Progress1	100.55	7.11	202.29	3.49	32.44
7	Progress2	31.76	2.338	133.28	3.91	39.36

Species identification of bdelloids was possible only in the case of frozen samples (from three sites: Progress1, Progress2 and Vechernyaya3) that recovered from cryptobiosis almost completely. Most of the individuals in the dried samples retained their unidentifiable tun shape after hydration.

All notes relating to the identification of the rotifer species are presented in the Supplementary material.

Morphological examination of the bdelloids revealed six morphotypes. Four can be identified as Antarctic species. *Adineta grandis* (Murray 1910) and *Philodina gregaria* (Murray 1910) were recorded at Progress1; *Adineta emsliei* (Iakovenko 2015) and *Habrotrocha antarctica* (Iakovenko 2015) were recorded at Progress2. It is noteworthy that the morphometric analysis showed differences between the morphotypes found and the "type varieties" discussed by Iakovenko (2015; Supplementary Table S4). On the basis of morphology, the other two species found in samples from the Vechernyaya3 site can be considered as belonging to the "*Habrotrocha constricta*" (Dujardin 1841) species group and to the "*Adineta vaga*" (Davis 1873) species group, both of which are cosmopolitan taxa; the precise status of which is under debate (Figs. 4–6).

DNA sequencing was carried out on the specimens identified as *A. grandis* and *Ph. gregaria* from Progress1 (Supplementary Table S5). The analysis using BLAST fully supported the morphological identification. DNA samples, which were extracted from individuals with the *Ph. gregaria* morphotype (GenBank identification OK325599 and MT584979), can be attributed to the species with 98–100% certainty, according to the BLAST search. DNA samples from the specimen identified, on the basis of morphology, as *A. grandis* (GenBank ID OK325600) referred to the said species with more than 99% similarity (Supplementary Fig. S1).

As identification without the analysis of nucleotide sequences is arguable, we indicated the other four species as "cf." Such ambiguity is especially suitable in relation to those species which we initially linked with *A. vaga* and *H. constricta*.

All tardigrades were Eutardigrada members. The genera *Macrobiotus*, *Mesobiotus* and *Diphascon* have been identified in the samples collected over the last two seasons.

Abundance

High abundances of micrometazoans were found at all the studied sites (Fig. 7). Assemblages in five of the seven sites consisted mainly of bdelloid rotifers (72.8–100% of individuals), while at Vechernyaya1 and Vechernyaya2 tardigrades predominated (88.7 and 62.0% of individuals, respectively). The sites with the highest densities (with mean and median values over 1000 ind·0.1 gdw^{-1}) were the rotifer-dominated sites—Druzhnaya, Haswell and Progress1—where rotifer mean densities ranged from 1125 to 3030 ind·0.1 gdw^{-1}, and the maximal value was about 8650 ind·0.1 gdw^{-1}. The tardigrade densities at the two tardigrade-dominated sites—Vechernyaya1 and Vechernyaya2—were lower by an order of magnitude and yielded 590 and 281 ind·0.1 gdw^{-1}, which was comparable to that of the rotifers at Vechernyaya3 and Progress2: 400 and 639 ind·0.1 gdw^{-1} (mean values), respectively. At Vechernyaya3, the range of values among samples at a particular site was notable: the maximum exceeded 3000 ind·0.1 gdw^{-1}.

Haswell and Progress2 were almost barren of tardigrades, whilst no tardigrades were detected at all at Vechernyaya3. In contrast, rotifers were found in every sample at each site. Rotifer and tardigrade densities did not correlate across the samples within each site, except for Progress2, where they demonstrated a high Spearman's rank correlation coefficient between the two groups, with a significant *p* value level (Fig. 8; Table 3).

The correlation analysis of the nutrient content in the samples and the abundance values of the rotifers and the tardigrades did not indicate possible links in any comparison case (Table 4).

Discussion

The first description of the very high rotifer abundances in the terrestrial ecosystems of Antarctica is to be found
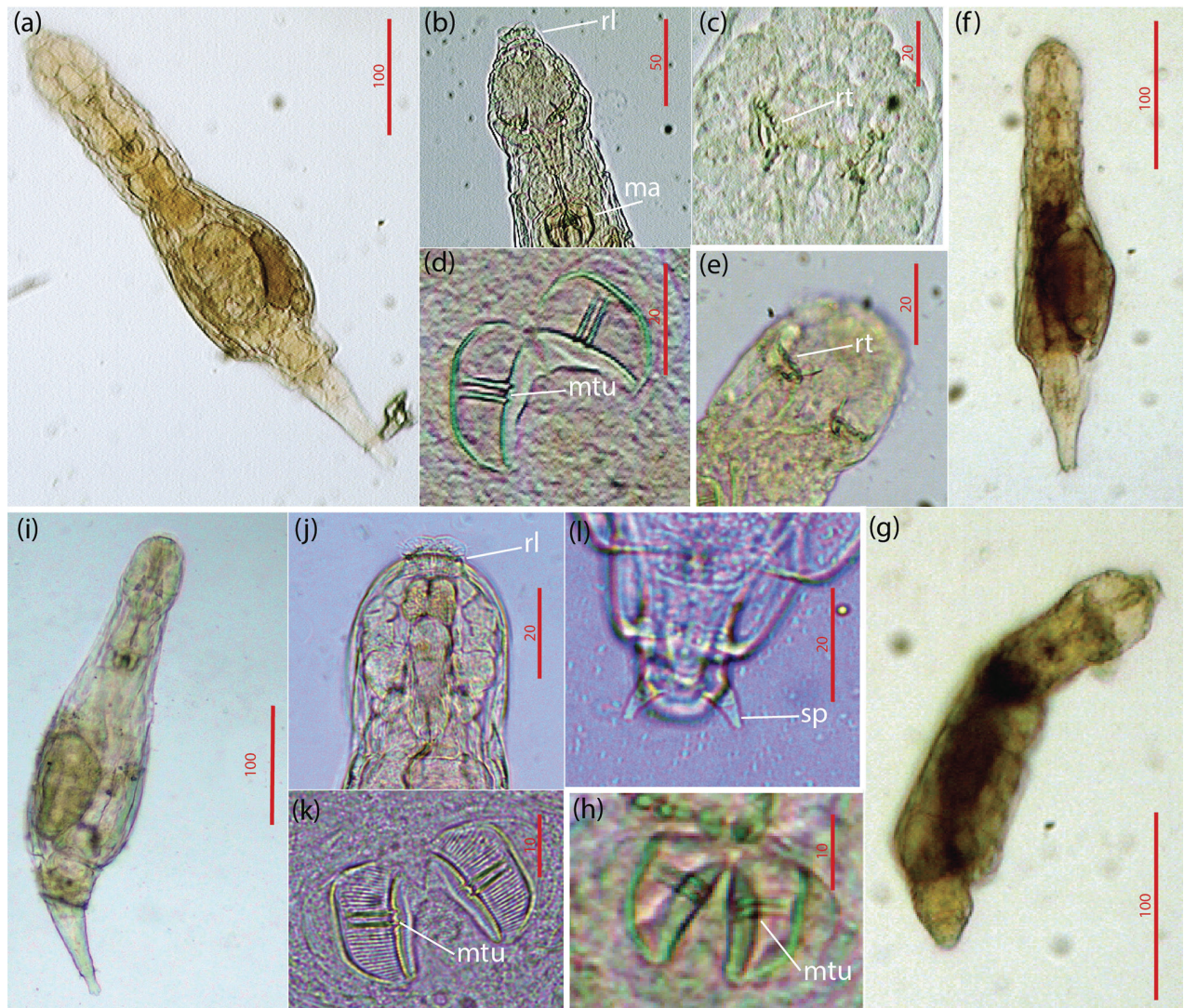
Fig. 4 Morphological details of the rotifers of the genus *Adineta*, isolated from *Prasiola crispa*: (a) dorsal view of crawling specimen of *A. grandis*; (b) and (c) dorsal view of head; (d) masticatory apparatus with trophi; (e) dorsal view of the head of *Adineta* cf. *emsliei*; (f) and (g) dorsal view of a crawling specimen; (h) masticatory apparatus with trophi; (i) dorsal view of crawling specimen of *Adineta* cf. *vaga*; (j) dorsal view of head; (k) masticatory apparatus with trophi; (l) dorsal view of foot. The following terms are abbreviated: rostrum lamella (rl); masticatory apparatus with trophi (ma); rake teeth (rt); major teeth in unci (mtu); spurs (sp). Scale bars are in µm.

in Murray's pioneer zoological surveys (1910a, b). He observed (1910b: 18): "I have never anywhere seen Bdelloid rotifers so plentiful as are the two dominant species at Cape Royds (*Philodina gregaria* and *Adineta grandis*). Among the higher Invertebrata, the Rotifera are easily first in numbers, both of individuals and species." This work lacks detailed information regarding the habitat that was so abundantly populated by bdelloids; however, Murray was probably referring to multicellular algae (according to his description, very similar to *Prasiola crispa*) and definitely not mosses.

More recent studies showed that *P. crispa* is inhabited by rotifers (Suren 1990; Everitt 1981) and tardigrades (Dougherty & Harris 1963; Miller et al. 1988; Miller, Heatwole et al. 1994; Miller & Heatwole 1996; Sohlenius et al. 2004; Heatwole & Miller 2019). Other small animals are found in *P. crispa* (Sinclair et al. 2006; Dalto et al. 2010; De Mendonça et al. 2012; Gantz et al. 2018), including *Belgica antarctica*, the Antarctic endemic insect (Sugg et al. 1983). It is likely that terrestrial algae provide favourable microhabitats for invertebrates, and some results indicate that arthropods prefer algae to mosses because of the feeding conditions (Bokhorst et al. 2007).

Fig. 5 Morphological details of the rotifers of the genus Habrotrocha, isolated from *Prasiola crispa*: (a) dorsal view of crawling specimen of *Habrotocha* cf. *constricta*; (b) dorsal view of feeding specimen; (c) dorsal view of head with corona; (d) and (e) dorsal view of foot; (f) and (g) masticatory apparatus with trophi; (h) dorsal view of crawling specimen of *Habrotocha* cf. *antarctica*; (i) dorsal view of feeding specimen; (j) masticatory apparatus with trophi. The following terms are abbreviated: corona (cr); spurs (sp); pedicels (pd); upper lip (up); major teeth in unci (mtu). Scale bars are in μm.

In Svalbard, in the Arctic, the *P. crispa* coverage near depositions of seabird guano is one of the most important environmental factors that positively affect the diversity and the abundance of springtails (Zmudczyńska et al. 2012). However, since Murray's work, no one has confirmed the huge abundances of invertebrate animals in algae, with the exception of one study (Schulte et al. 2008), which documented a high density of Antarctic collembola eggs, presumably as a consequence of global warming.

Our finding of thousands of individual bdelloid rotifers in a very limited algal volume illustrates the poorly described phenomenon. The density values per dry weight unit that we found are higher by orders of magnitude than those reported in the literature for algae, lichens, soil and—the most interesting—mosses (Table 5). Bryophytes are sometimes claimed to be the preferred habitat for Antarctic micrometazoans (e.g., Porazinska et al. 2004; Sohlenius et al. 2004; Smykla et al. 2010). In the Larsemann Hills, it has been shown that all
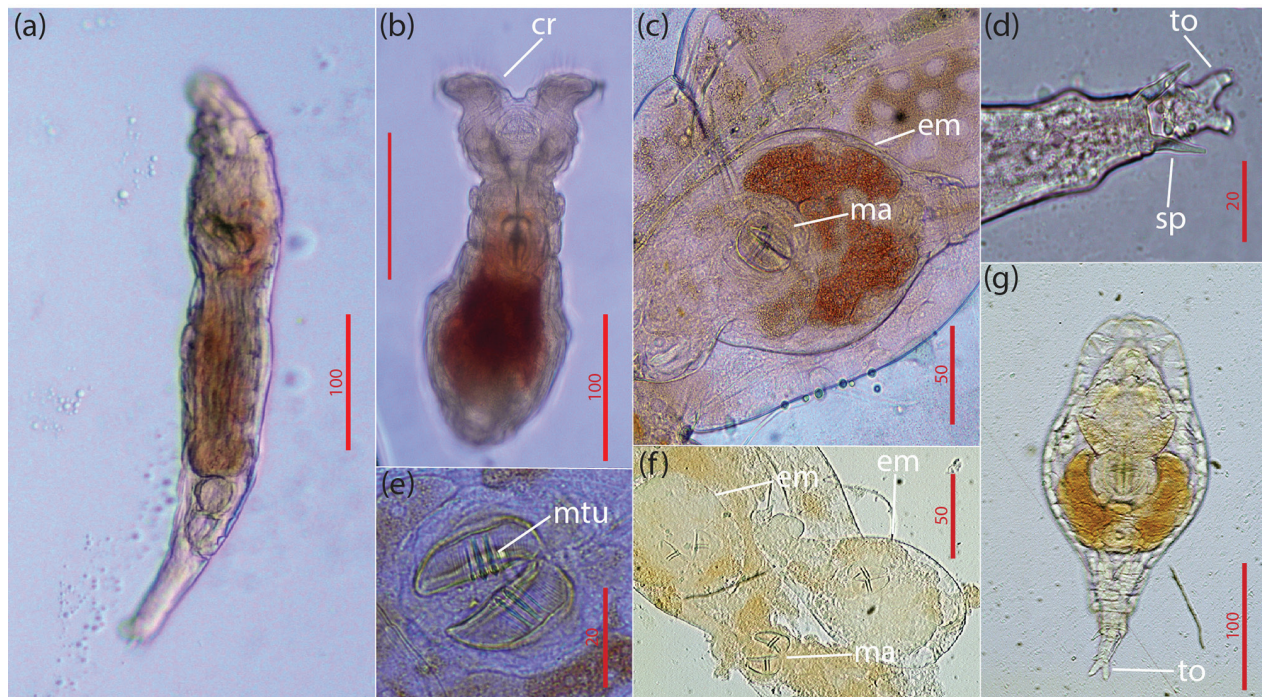
Fig. 6 Morphological details of *Philodina gregaria* isolated from *Prasiola* crispa: (a) lateral view of crawling specimen of *Philodina gregaria*; (b) dorsal view of feeding specimen; (c) and (f) embryos in the maternal specimen; (d) dorsal view of foot; (e) masticatory apparatus with trophi; (g) dorsal view of semi-squeezed specimen. The following terms are abbreviated: corona (cr); masticatory apparatus with trophi (ma); embryos (em); major teeth in unci (mtu); spurs (sp); toes (to). Scale bars are in µm.
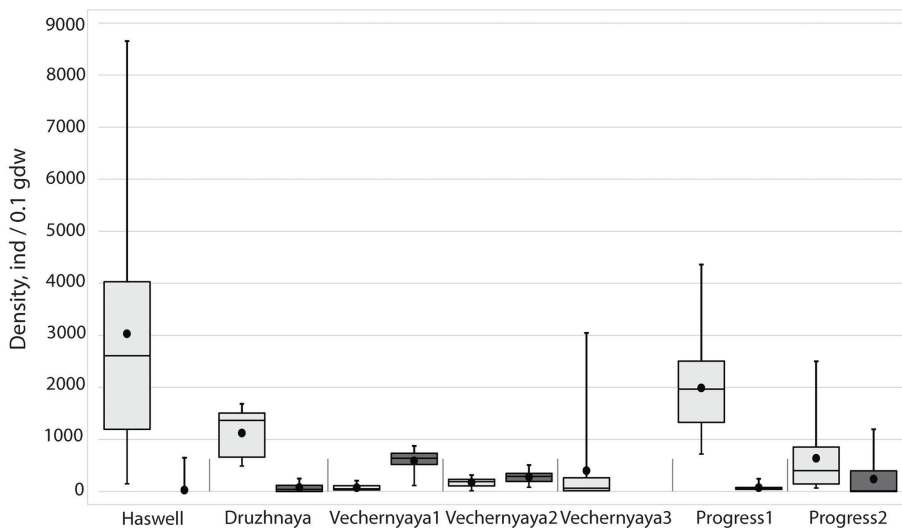
Fig. 7 Rotifer (light grey) and tardigrade (dark grey) density values. Boxes show interquartile ranges; box horizontal midlines are medians; whiskers are maxima and minima; black dots represent means.

micrometazoan taxa occur more frequently in mosses, and they avoid soil (Heatwole & Miller 2019). Unfortunately, our findings cannot be compared with micrometazoan density results that were assessed using other types of standardization, for example, individuals per area or individuals in a core. However, roughly extrapolating between the sample volumes analysed in other studies (e.g., the 3.0 × 3.5 cm cores used by Jennings [1976]) and the much smaller sample volumes analysed in the present study suggests values that are
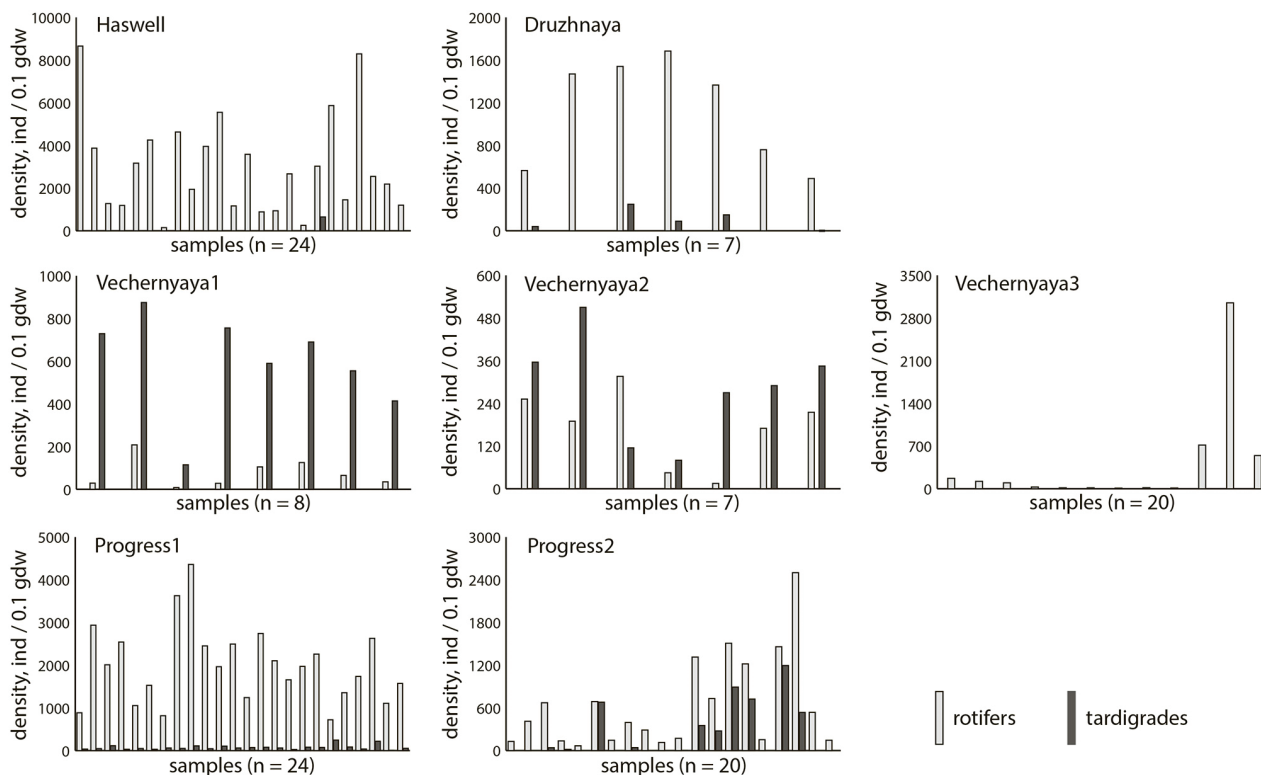
Fig. 8 Comparison of rotifer and tardigrade density values.

Table 3 Correlations between rotifer and tardigrade densities. The significant correlation is in boldface.

No.	Site	Correlation indices	
		R^a	P^b
1	Haswell	0.18	0.39
2	Druzhnaya	0.65	0.12
3	Vechernyaya1	0.41	0.33
4	Vechernyaya2	0.29	0.56
5	Vechernyaya3	-	-
6	Progress1	0.16	0.46
7	Progress2	**0.82**	**0.000012**

[a]Spearman's rank correlation coefficient. [b]Significance.

similar or at least of the same order of magnitude. Unfortunately, there were no quantitative data in Murray's work (1910a, b), which was the first to report high rotifer abundances in Antarctic algae.

Four of the six species of rotifers found in the present study are Antarctic endemics. *Philodina gregaria* and *Adineta grandis*, found on the continent during the earliest zoological surveys (Murray 1910a), have large sizes (up to 700 μm), are viviparous and are conspicuous because of their pigmentation: *A. grandis* is bright red and

Ph. gregaria is dark red. The correct *A. grandis* identification was possible only by applying mtDNA sequencing, as *A. fontanetoi* (Iakovenko 2015) is a morphological twin of *A. grandis* and these can only be distinguished from one another genetically. *Adineta emsliei* and *Habrotocha antarctica* were recently described as a result of an in-depth study of Antarctic bdelloids (Iakovenko et al. 2015). Despite the fact that both *Habrotocha constricta* and *Adineta vaga* had been formally noted in the Antarctic (Segers 2007), identifying them on the basis of morphology alone is problematic. Application of DNA taxonomy methods, along with analysis of the trophi microstructure, has demonstrated that Antarctic representatives are distinct from their relatives elsewhere in the world (Iakovenko et al. 2015). The situation in relation to *A. vaga* is particularly complicated as this name has been assigned to a species that should either be re-described as a species group or considered unidentifiable (Örstan 2020). The difference between the supposed member of the *A. vaga* complex in the present study and *A. emslei*—the Antarctic variant of *A. vaga* (Iakovenko et al. 2015)—underscores the need for modern molecular approaches.

As tardigrades and rotifers in the Antarctic often co-exist with nematodes (e.g., Sohlenius et al. 2004), the utter absence of the latter is a very unexpected result of the study. These organisms—the most abundant and

Table 4 Correlations between micrometazoan densities and nutrient concentrations across all sites.

Taxa	Na		P		K		N		C	
	R^a	p^b	R^a	p^b	R^a	p^b	R^a	p^b	R^a	p^b
Rotifers	0.50	0.27	−0.42	0.35	−0.39	0.39	0.04	0.96	−0.09	0.92
Tardigrades	−0.57	0.20	0.14	0.78	0.46	0.30	0.35	0.44	0.43	0.42
All	0.28	0.56	−0.50	0.27	−0.25	0.59	0.32	0.49	0.37	0.49

[a]Spearman's rank correlation coefficient. [b]Significance.

diverse group of metazoans on Earth (Poinar 2015; van den Hoogen et al. 2019)—are widespread in the Antarctic (e.g., Velasco-Castrillón, Gibson et al. 2014; Velasco-Castrillón & Stevens 2014), including hostile Antarctic deserts, where nematodes can be the only metazoan taxon present (Zawierucha et al. 2021). Nematodes can dominate in algae (e.g., Sohlenius et al. 2004) and in bacterial mats, playing a key role as top consumers (Almela et al. 2019). Other than the present study, micrometazoan communities without nematodes are exemplified only in cryoconite holes, which are unique glacier habitats formed when dark debris on the ice leads to melting. The absence of nematodes in cryoconite holes can be explained by their unsuitableness for reproduction: low temperatures and rapid shifts from oxic to anoxic conditions (Zawierucha et al. 2021). Explanations for the absence of nematodes in samples of terrestrial alga *Prasiola crispa*, as we found, require further investigation.

The extremely high abundance of rotifers and tardigrades shown in the present study can be considered as a particular case of the patchy spatial distribution of Antarctic micrometazoans (Miller, Miller et al. 1994; Adams et al. 2006). Microsites with high abundance values, such as samples in the present study, may be interspersed with those where micrometazoan abundance is low to moderate or micrometazoans may be absent altogether, even within a homogeneous biotope. As the data set in the present study is limited, we need further research in this direction to elucidate this pattern. However, some preliminary ideas regarding to micrometazoan extreme abundances in *Prasiola crispa* can be outlined.

The specific conditions of terrestrial algae may be facilitating an increase in micrometazoans. The chemical composition of the substrate seems to be an important factor affecting the distribution of animals dwelling there. The organic matter content and C/N ratio do not influence the micrometazoan community across several types of habitats (Sohlenius et al. 1996). In Antarctic soils, strong nutrient contamination by bird faeces (high N content) inhibits tardigrades (Sohlenius et al. 2004; Smykla et al. 2012); likewise, nematodes have a limited presence in ornithogenic soils in active penguin rookeries (Porazińska et al. 2002a).

However, rotifers appear to be associated with areas of higher organic content (Sinclair & Sjursen 2001; Sohlenius et al. 2004). Bokhorst et al. (2019) demonstrate increases in the abundance and diversity of moss- and lichen-dwelling tardigrades and micro-arthropods with nitrogen input from Antarctic marine vertebrates. In the Arctic, the picture seems to be clearer: invertebrate communities are more abundant in habitats enriched by seabird guano (Zmudczyńska et al. 2012; Zmudczyńska-Skarbek et al. 2015; Zawierucha et al. 2016; Zmudczyńska-Skarbek et al. 2017). However, all these studies neglected rotifers. In our study, most of sites presented the places Adélie penguin (*Pygoscelis adeliae*) moulting. We expected to find more micrometazoans at the sites where there were more expressed signs of 'ornithogeneity,' but this factor was not estimated quantitatively. Nonetheless, we demonstrated a lack of correlations between nutrient content and animal abundance (Table 4). It may be that the microscopic animal communities are mostly determined by completely different chemistry-independent factors, while the observed variability of element concentration in the samples, caused by the presence of inconspicuous mineral particles (sand grains), does not influence the abundance of animals that live in algae.

The explanation may possibly lie in the algal surface, which is structurally complicated, offering many microhabitats for tiny animals. Another important aspect that may have a positive influence on abundance is the water content around the algae. The existence of Antarctic invertebrates is limited by liquid water, which is the main driver, critically influencing their dispersal (Block et al. 2009; Convey et al. 2014). All samples were collected during the summer season (December–February), when bryosystems experienced their highest saturation with thawing water. We suggest that the water film that forms on and around the algal thalli may serve as a spacious environment for micrometazoans. This may help account for the predominance of bdelloid rotifers over tardigrades. These animals, unlike tardigrades, filter sediment and can easily feed on bacteria and other microscopic food in a liquid environment. Moreover, many rotifers are semi-attached animals that alternate between swimming while feeding and crawling or sitting on a firm surface.

Table 5 Micrometazoan density values (ind·0.1 gdw⁻¹), comparing data obtained in the present study to the literature. Some values taken from the literature were recalculated to ind·0.1 gdw⁻¹ from individuals per other weight units (g, 100 g, etc.).

Taxa	HI[a]	LH[b]	TH[c]	VH[d]	DM[e]	DM[f]	DM[g]	DM[h]	DM[i]	RI[j]	EL[k]	VL[l]	VL[m]	DV[n]	DV[o]	DV[p]	DG[q]
	Algae					Other bryosystem substrata (e.g., moss)					Soil						
Rotifers																	
Mean	3030	1251	216	ca. 16[s]	87	1–15[r]	5	5	2–51[r]	0–<1[r]	nd[t]	1	3	nd[t]	0–<1[r]	<1	–
Max	8656	4360	3049	ca. 40[s]	272	ca. 45[s]	75	75	425	<1	nd[t]	8	34	nd[t]	<1	n/d	–
Tardigrades																	
Mean	29	129	290	ca. 39[s]	2	<1–2[r]	4	1	<1–4[r]	0–<1[r]	<1	<1	<1	nd[t]	0–<1[r]	<1	–
Max	647	1195	875	ca. 49[s]	3	ca. 20	58	36	240	<1	<1	2	3	nd[t]	<1	<1	–
Nematodes																	
Mean	–	–	–	nd[t]	123	<1–3[r]	3	3	<1–50[r]	0–<1[r]	nd[t]	<1	<1	0–<1[r]	<1	<1	<1
Max	–	–	–	nd[t]	506	ca. 20[s]	21	59	506	<1	nd[t]	2	3	<1	<1	nd[t]	<1
Totals																	
Mean	3059	1380	506	nd[t]	nd[t]	1–20[r]	nd[t]	nd[t]	3–102[r]	0–<1[r]	nd[t]	nd[t]	4	nd[t]	<1	<1	<1
Max	8656	4470	3049	nd[t]	nd[t]	nd[t]	nd[t]	nd[t]	781	<1	nd[t]	nd[t]	35	nd[t]	<1	nd[t]	<1

[a]Haswell Island; present study. [b]Larsemann Hills & Sanderfjord Bay; present study. [c]Thala Hills; present study. [d]Vestfold Hills; Everitt 1981. [e]Dronning Maud Land; Sohlenius et al. 2004. [f]Dronning Maud Land; Sohlenius et al. 1995. [g]Dronning Maud Land; Sohlenius et al. 1996. [h]Dronning Maud Land; Sohlenius et al. 2002a. [i]Ross Island; Porazińska et al. 2002a. [k]Ellsworth Land Convey & McInnes 2005. [l]Victoria Land; Smykla et al. 2010. [m]Victoria Land; Smykla et al. 2018. [n]Antarctic Dry Valleys; Porazińska et al. 2002b. [o]Antarctic Dry Valleys; Andriuzzi et al. 2018. [p]Antarctic Dry Valleys; Shaw et al. 2018. [q]Darwin Glacier; Zawierucha et al. 2019. [r]Range reflects the several types of estimated habitats/areas. [s]Approximate values (taken from figures). [t]No data.

Hydration, together with the availability of food, creates a superior environment for them, causing the population to increase rapidly in terms of number and biomass. The fact that rotifers and tardigrades require different conditions is illustrated by our results: almost no correlations between these two were found. However, others have found correlations between rotifer and tardigrade abundances (Sohlenius & Boström 2005, 2008).

Another consideration is the bryosystem characteristics on a broader scale. The sites Progress1 and Progress2 have voluminous conglomerations of algae in deep hollows, as opposed to the thin algae layers on ground or rock surfaces that we commonly encountered during our collecting. Deep hollows can foster the continuous accumulation of animals, as new algal layers grow. The cumulative effect of this process over years, allowing for periods of abundance (summer) interspersed with dormancy periods (winter), has resulted in millions of rotifers.

Conclusion

We observed remarkably high abundances of bdelloid rotifers, followed by tardigrades, in algal samples collected at different East Antarctic oases. These values far exceeded those available in the literature for algae and other bryosystems components and soil. Counter to our expectations, we found no nematodes. We believe that the irregular distribution of micrometazoans in our samples does not relate to nutrient content but, instead, is likely to be associated with algal surface characteristics, moisture content and/or the history of the habitat formation.

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Disclosure statement

The authors report no conflict of interest.

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