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# Distribution patterns of micro- and meso-zooplankton communities in sea ice regions of Lützow-Holm Bay, East Antarctica

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Abstract In the Southern Ocean, zooplankton research has focused on krill and macro-zooplankton despite the high densities of micro- and meso-zooplankton. We investigated their community structure in relation to different sea ice conditions around Japan's Syowa Station in Lützow-Holm Bay, in the summers of 2011 and 2012. Zooplankton samples were collected using vertical hauls (0-150 m), with a closing net of 100-µm mesh size. The results of cluster analysis showed that the communities in this region were separated into fast ice, pack ice, and open ocean fauna. The fast ice fauna had lower zooplankton abundance  $(393.8-958.9 \text{ inds. m}^{-3})$  and was dominated by cyclopoid copepods of Oncaea spp. (54.9-74.8 %) and Oithona similis (6.6-19.9 %). Deep-water calanoid copepods were also found at the fast ice stations. Pack ice and open ocean fauna had higher zooplankton abundance  $(943.6-2,639.8 \text{ inds. m}^{-3})$  and were characterized by a high density of foraminiferans in both years (6.6-61.9 %). Their test size distribution indicated that these organisms were possibly released from melting sea ice. The pteropod *Limacina* spp. was a major contributor to total abundance of zooplankton in the open ocean zone in 2012 (26.4 %).

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K. T. Takahashi · T. Iida · T. Odate · M. Fukuchi National Institute of Polar Research, 10-3 Midori-cho, Tachikawa, Tokyo 190-8518, Japan The physical and/or biological changes between 2 years may affect the abundance and distribution of the dominant zooplankton taxa such as cyclopoid copepods, foraminiferans, and pteropods. Information on the relationships between the different species associated with sea ice will help to infer the possible future impacts of climate change on the sea ice regions.

**Keywords** Southern Ocean · Sea ice · Micro- and meso-zooplankton · Foraminiferans

# Introduction

Zooplankton are secondary producers in marine ecosystems and play an important role as food for pelagic fish and air-breathing predators such as seals, whales, and penguins in the Southern Ocean (e.g. Hempel 1985a). While various factors defining zooplankton distribution in the Southern Ocean have been discussed, sea ice plays a crucial, highly dynamic, and variable role in the life cycles of organisms including zooplankton in the sea ice region (Massom and Stammerjohn 2010). In fact, Antarctic zooplankton communities are usually divided into three ice-associated faunal zones: (1) the northern ice-free zone, dominated by copepods and Salpa thompsoni; (2) the seasonal pack ice zone, dominated by Antarctic krill Euphausia superba; and (3) the fast ice zone, where the ice krill Euphausia crystallorophias is dominant (Hempel 1985b; Hosie 1994; Loeb 2007). However, these distributional data for macrozooplankton are mostly based on information obtained from sampling with 200-500 µm nets or coarser nets such as the RMT (Rectangular Mid-water Trawl) 8. Recently, the increasing use of finer plankton nets (60–200  $\mu$ m) has provided a more realistic view of the ecological

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significance of meso-  $(200-2,000 \ \mu\text{m})$  and micro-zooplankton  $(20-200 \ \mu\text{m})$ , such as small calanoid and cyclopoid copepods, foraminiferans, and appendicularians (Metz 1996; Atkinson 1998; Schnack-Schiel et al. 1998; Atkinson and Sinclair 2000; Dubischar et al. 2002). Micro- and meso-zooplankton have been estimated to have a high abundance in the seasonal ice zone (Chiba et al. 2001; Swadling et al. 2010). Although their roles in the pelagic food web under sea ice are being gradually identified, the dynamics of these smaller zooplankton communities within the sea ice regions are still poorly understood.

Lützow-Holm Bay, where Japan's Syowa Station is located, is a region where fast ice cover persists during summer, and some studies have been made of the zooplankton community structure around the shore-based stations of this bay (e.g. Fukuchi and Sasaki 1981; Fukuchi and Tanimura 1981; Fukuchi et al. 1985; Tanimura et al. 1986; Tanimura et al. 2008). The marine biological monitoring program of the sea ice region by ship-based sampling began with the 52nd Japanese Antarctic Research Expedition (JARE-52; 2010/2011 season). The aim of this program was to investigate biological production and mechanisms in relation to sea ice. The objectives of the present study were to examine the abundance and distribution patterns of microand meso-zooplankton communities in Lützow-Holm Bay

Fig. 1 Spatial and temporal variability in sea ice distribution around Syowa Station in the austral summers of 2011 (a) and 2012 (b). *Stars* indicate sampling stations

during the austral summer, and their relationship to sea ice distribution. To achieve this objective, it was assumed that sea ice distribution would affect the distribution of microand meso-zooplankton communities in the sea ice regions as well as that of macro-zooplankton.

## Materials and methods

# Sampling

The survey was conducted during two cruises of the Japanese icebreaker *Shirase* through the sea ice zone in Lützow-Holm Bay. The first took place from February 9 to 24, 2011 (JARE-52) and the second from February 14 to March 4, 2012 (JARE-53; Fig. 1). Zooplankton samples were collected using a closing net (mouth diameter 0.75 m, mesh size 100  $\mu$ m; Table 1) at nine stations (five in 2011 and four in 2012) with contrasting sea ice environments (Fig. 1): fast ice (four stations: 52A, 52B, 53A, and 53B), pack ice (three stations: 52C, 52D, and 53C), and ice-free open ocean (two stations: 52BP and 53BP). To prevent the sea ice from being mixed into the net, an ice fence was employed and the net was closed as it reached the surface (Fig. 2; Takahashi et al. 2012). The net was equipped with



Table 1 Sampling locations and environmental information

Station	Location		Date (LMT)	Sampling	Sea-ice	Saverage	Taverage	Chl. $a$
	Lat. (S)	Long. (E)		depth (m)	distribution	[min, max] (PSU)	[min, max] (°C)	(mg m <sup>2</sup> )
52A	69°03′	39°20′	February 9, 2011	75-0	Fast ice	33.97 [31.76, 34.15]	-1.72 [-1.81, -1.64]	9.78
52B	68°59′	39°11′	February 12, 2011	135-0	Fast ice	34.12 [33.98, 34.22]	-1.72 [-1.86, -1.64]	1.08
52C	68°31′	38°42′	February 18, 2011	150-0	Pack ice	34.06 [33.71, 34.22]	-1.80 [-1.85, -1.76]	12.48
52D	67°43′	38°18′	February 23, 2011	150-0	Pack ice	34.07 [33.27, 34.43]	-1.51 [-1.78, -0.84]	30.01
52BP	66°50′	37°49′	February 24, 2011	150-0	Open ocean	34.45 [34.07, 34.68]	0.63 [-0.52, 1.44]	80.13
53A	68°57′	39°05′	February 14, 2012	150-0	Fast ice	34.00 [32.27, 34.26]	-1.76 [-1.85, -1.71]	1.46
53B	68°56′	39°05′	February 17, 2012	150-0	Fast ice	No data	-1.74 [-1.82, -1.71]	1.16
53C	68°34′	38°39′	March 1, 2012	150-0	Pack ice	No data	No data	12.26
53BP	66°50′	37°51′	March 4, 2012	150-0	Open ocean	34.33 [33.88, 34.77]	0.17 [-1.73, 1.31]	28.74

**Fig. 2** Zooplankton sampling in the pack ice using an "icefence" to guard against net closing due to sea ice



onset

recovery

a flow meter to estimate the volume of water filtered and was vertically hauled from a depth of 150 m to the surface at stations where the bottom was deeper than 150 m, or from 5 m above the bottom to the surface at stations where the bottom was shallower than 150 m (Table 1). All samples were fixed immediately with buffered 5 % formaldehyde and seawater solution.

A number of environmental variables were also measured during the cruise. Temperature and salinity were measured at 1-m intervals down to the seafloor using a CTD (SBE 55 ECO, Sea-Bird Electronics, Bellevue, Washington, DC, USA). Water samples for measurement of chlorophyll a (Chl a) concentration were collected from three to seven depths in the upper 150 m with a Niskin bottle mounted on the CTD. Chl a concentration was determined fluorometrically with a Turner Designs fluorometer (model 10AU). For the purpose of comparison with zooplankton data, temperature and salinity data were averaged across the sampling layers. And Chl a concentrations were integrated vertically (e.g. Suzuki et al. 1998; Uitz et al. 2006).

Daily values of sea ice concentration were obtained from the National Snow and Ice Data Center (NSIDC) in Boulder, Colorado, USA. These data were derived from passive microwave data collected by the Special Sensor Microwave Imager (SSMI) (Cavalieri et al. 1996) and the Advanced Microwave Scanning Radiometer for EOS (AMSR-E) (Cavalieri et al. 2004).

## Analysis

In the laboratory, zooplankton samples were split using a Motoda box splitter (Motoda 1959) so that approximately 1,000 individuals were counted per sample. The samples were identified to the lowest possible taxonomic level (generally species or genus) with a stereomicroscope. Members of the Ostracoda, Polychaeta, Salpa, Foraminifera, Chaetognatha, and Echinodermata were not identified to species level (Table 2). The copepods Oithona similis and Oncaea spp. were classed as adult or copepodite, with adult Oncaea spp. further identified to species level and classified as male or female. The maximum diameter of the test was measured for individual foraminiferans because the test size gives an indication of the growth stage. The test sizes of approximately 100 foraminiferans were measured per sample, except in cases where there were <100individuals. Individual counts were converted to the number of individuals per 1 m<sup>3</sup> for each station.

Abundance data were log-transformed ( $\log_{10}[n + 1]$ ) to decrease any bias in abundance, and a similarity matrix was constructed using the Bray–Curtis similarity index (Field et al. 1982). For grouping the samples, group average linkage was performed based on similarity. The SIM-PER (similarity percentage) routine identified those species contributing to the similarity within the observed clusters. These statistical analyses were conducted using the PRI-MER v6 software package (Clarke and Gorley 2006). The Shannon index (H') was chosen to show the species diversity in each cluster group.

## Results

# Environmental variables

The average temperature and salinity ranged from -1.80 to 0.63 °C and 33.97 to 34.45 PSU, respectively (Table 1). A few stations contributed no data as the CTD did not work. Vertically integrated Chl *a* concentrations ranged from 1.08 to 80.13 mg m<sup>-2</sup> (Table 1). Both temperature and salinity tended to increase from the fast ice stations to the open ocean stations each year, Chl *a* concentration showed a similar trend, with the exception of station 52A.

# Community structure

A total of 42 zooplankton species/taxa were identified from the nine stations (Table 2). Zooplankton abundance ranged from 393.8 to 2,639.8 inds.  $m^{-3}$  (Fig. 3). Values were lower at the stations under fast ice and greater at those under pack ice or in the open ocean. Copepods (including nauplius stages) were generally the dominant zooplankton

Table 2 Zooplankton	taxa/species	recorded	in	this	stud
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Copepoda	Aetideidae
	A. antarctica
	Amallothrix sp.
	Calanoida
	C. acutus
	Calanus propinquus
	Clausocalanus laticeps
	C. citer
	Euchaetidae
	Euchirella rostromagna
	Haloptilus oxycephalus
	Harpacticoida
	Heterorhabdidae
	Heterorhabdus austrinus
	Metridia gerlachei
	M. pygmaeus
	Oithona frigida
	O. similis
	Oncaea spp.
	Paraeuchaeta antarctica
	Paraeuchaeta biloba
	Racovitzanus antarcticus
	Rhincalanus gigas
	S. minor
	Scolecitrichidae
	Spinocalanidae
Copepod nauplii	
Amphipoda	Themisto spp.
Euphausiacea	Euphausiidae
	E. superba
	Thysanoessa macrura
Ostracoda	
Polychaeta	
Appendicularia	Fritillaria spp.
	Oikopleura spp.
Salpa	Salpa spp.
Chaetognatha	
Siphonophorae	
Echinodermata	
Pteropoda	Limacina spp.
Foraminifera	
Fish egg	

component, with the copepods *O. similis* and *Oncaea* spp. being the most abundant copepod taxa at all stations. For example, at the fast ice stations (52A, 52B, 53A, and 53B), *Oncaea* spp. accounted for 54.9–74.8 % of the total zooplankton abundance. Foraminiferans were also highly abundant at pack ice and open ocean stations, comprising 6.6–61.9 % of the total zooplankton community. *Limacina* 

Fig. 3 Abundance (inds.  $m^{-3}$ ) and species composition of

zooplankton at all stations in 2011 and 2012





Fig. 4 Dendrogram of the cluster analysis based on the Bray–Curtis similarity index with UPGMA linkage

spp. occurred in high numbers at pack ice and open ocean stations in 2012 (10.3 and 26.4 %, respectively). Copepod nauplii were also abundant, accounting for 4.3–21.8 % under pack ice and in the open ocean (Fig. 3).

Cluster analysis revealed three groups at  $\geq$ 72.2 % similarity (Fig. 4). The groups were clearly separated by

the interface between fast ice and pack ice. Group 1 comprised all four stations located under fast ice in 2011 and 2012. Groups 2 and 3 comprised the five stations located under pack ice or in the open ocean, Group 2 with the two stations in 2012 and Group 3 with the three stations in 2011.

The species contributing most to the similarity within Group 1 was *Oncaea* spp., followed by *O. similis*, *Microcalanus pygmaeus*, and *Ctenocalanus citer* (Table 3). Group 2 was characterized by *O. similis*, copepod nauplii, *Oncaea* spp., *Limacina* spp., and foraminiferans. Group 3 was mainly composed of foraminiferans, *Oncaea* spp., *O. similis*, copepod nauplii, and *C. citer*. The values for average abundance and species diversity were the highest in samples within Group 2, followed by Group 3 and then Group 1 (Table 3; Fig. 4). The average abundance of zooplankton in samples within Group 2 was more than triple that of samples within Group 1, while the species richness of samples within each group was almost the same (Table 3). The organisms contributing most to the difference between Groups 1 and 3 were foraminiferans, followed by copepod nauplii, *Fritillaria* spp., *M. pygmaeus*, *O. similis*, and echinoderms (Table 4). For Groups 1 and 2, *Limacina* spp. and foraminiferans were the most influential contributors to the dissimilarity. The species causing most of the dissimilarity between Groups 2 and 3 was *Limacina* spp., followed by *Fritillaria* spp., *Calanoides acutus*, *Scolecithricella minor*, copepod nauplii, and polychaetes (Table 4).

## Population structure

The abundance of O. similis ranged from 47.9 to 491.4 inds.  $m^{-3}$  (Fig. 5a), and their pattern of abundance follows a similar pattern to that of total zooplankton abundance in this study (Fig. 3). Copepodite stages were dominant at all stations, accounting for 65.5-94.2 % of total abundance (Fig. 5a). The abundance of Oncaea spp. varied from 96.0 to 670.0 inds.  $m^{-3}$  (Fig. 5b). It was highest at pack ice station 52C in 2011 (472.0 inds.  $m^{-3}$ ), while in 2012, the highest value occurred at fast ice station 53A (670.0 inds.  $m^{-3}$ ). The species distribution and stage composition at each station were similar between 2011 and 2012: adults were dominant under fast ice, contributing 50.4-76.5 %, and adult abundance gradually decreased toward the open ocean. In contrast, copepodites were most abundant at the open ocean stations (79.4 % in 2011 and 83.7 % in 2012) and decreased in abundance toward the fast ice. Adult Oncaea antarctica occurred in abundance only at the northern pack ice and open ocean stations, while adult Oncaea curvata occurred at the southern pack ice and fast ice stations. Adult male O. curvata tended to be dominant in the fast ice area.

The test size distribution patterns for foraminiferans in 2011 and 2012 were similar (Fig. 6). At the open ocean stations, 52BP and 53BP, size ranges were 77.8–320.4 and 86.4–330.2  $\mu$ m, respectively, and a peak in abundance occurred at approximately 175  $\mu$ m. The peak at pack ice station 52D was the same as that found at the open ocean stations. However, at this station, the size range was 110–354.8  $\mu$ m. The smaller individuals found at the open ocean stations were not present at station 52 D, and the largest sizes found were greater than those found at open ocean stations. At pack ice stations 52C and 53C, larger individuals were even more abundant in the samples (105.7–419.4 and 103.1–407.0  $\mu$ m, respectively) and two peaks in size occurred, at approximately 200 and 300  $\mu$ m (Fig. 6).

Table 3	Average	abundance	(inds.	m <sup>-3</sup> ),	species	richness	( <i>r</i> ),	and
diversity	(H') for	the three clu	isters i	dentifie	ed by clu	uster anal	ysis	

Species/taxa	Group 1	Group 2	Group 3
C. acutus	0.67	42.66	2.78
C. propinquus	0.59	4.99	1.77
Chaetognatha	0.94	2.33	2.70
Copepod nauplii	12.69	393.53	74.74
C. citer	38.86	182.88	59.89
Euchaetidae	2.83	0.10	1.82
Foraminifera	5.01	484.16	342.89
Fritillaria spp.	0.48	23.44	21.43
H. oxycephalus	0.03	1.39	0.20
Harpacticoida	1.98	1.50	1.89
Limacina spp.	0.93	466.04	1.11
M. gerlachei	4.17	28.17	5.24
M. pygmaeus	46.91	43.72	47.64
O. frigida	8.97	5.30	10.54
O. similis	72.30	396.85	227.77
Oncaea spp.	420.73	350.72	281.82
Ostracoda	0.26	0.24	1.07
Polychaeta	4.74	9.49	1.71
R. gigas	0.02	3.89	0.87
S. minor	0.53	16.40	1.92
Siphonophorae	1.81	1.62	1.52
Themisto spp.	0.02	0.05	0.06
Aetideidae	0.02	2.31	
P. antarctica	0.37	1.18	
Salpa spp.	0.02	1.59	
Heterorhabdidae	0.51		0.02
Oikopleura spp.	0.03		0.14
C. laticeps		2.27	2.51
Euphausiidae		0.05	0.01
T. macrura		0.05	0.09
A. antarctica	0.48		
Calanoida	0.06		
Echinodermata	2.12		
Spinocalanidae	0.48		
E. rostromagna		0.05	
E. superba		1.44	
Fish egg		0.05	
H. austrinus		0.19	
Amallothrix sp.			0.02
P. biloba			0.01
R. antarcticus			0.53
Scolecitrichidae			0.28
Total abundance (ind $m^{-3}$ )	629.53	2,468.56	1,094.99
Species richness (r)	31	32	31
Diversity $(H')$	1.84	2.98	2.60

Abundance of major contributor species/taxa are in bold

Table 4 Average abundance (inds.  $m^{-3}$ ) and contribution (%) within each cluster group as a result of SIMPER

Species/taxon	Av. Abund Group 1	Av. Abund Group 3	Contrib. (%)	Cum. (%)	
Groups 1 and 3 (average	ge dissimilarity 28.75)				
Foraminifera	5.01	342.89	18.35	18.35	
Copepod nauplii	12.69	74.74	8.57	26.92	
Fritillaria spp.	0.48	21.43	6.77	33.69	
M. pygmaeus	46.91	47.64	5.56	39.25	
O. similis	72.30	227.77	4.65	43.90	
Echinodermata	2.12	0.00	4.17	48.07	
	Group 1	Group 2			
Groups 1 and 2 (average	ge dissimilarity 38.94)				
Limacina spp.	0.93	466.04	14.61	14.61	
Foraminifera	5.01	484.16	11.76	26.37	
Copepod nauplii	Copepod nauplii 12.69		9.43	35.80	
C. acutus	0.67	42.66	6.44	42.25	
S. minor	0.53	16.40	6.30	48.54	
	Group 2	Group 3			
Groups 2 and 3 (average	ge dissimilarity 27.82)				
Limacina spp.	466.04	1.11	17.89	17.89	
Fritillaria spp.	23.44	21.43	7.19	25.08	
C. acutus	<i>acutus</i> 42.66		6.16	31.23	
<i>S. minor</i> 16.40		1.92	5.99	37.23	
Copepod nauplii	393.53	74.74	5.58	42.81	
Polychaeta	9.49	1.71	4.84	47.64	

# Discussion

Community structure of micro- and meso-zooplankton in sea ice regions

Information on micro- and meso-zooplankton community structure in sea ice regions, including those of the Southern Ocean, is sparse. The present study is one of few to try to assess variations in distribution and abundance of zooplankton under different sea ice conditions using shipbased sampling. This study indicates that micro- and mesozooplankton community structures vary according to sea ice distribution. Two characteristic distribution patterns of zooplankton occurred in both years: (1) the ubiquitous distribution of O. similis and Oncaea spp.; and (2) the regional distribution of foraminiferans which contributed highly to the separate characterization of zooplankton communities in this sea ice region.

In addition, the pteropod Limacina spp. was the major contributor to total zooplankton abundance in the open ocean zone in 2012 (Fig. 3). Pteropods are ubiquitous components of Southern Ocean zooplankton communities and are extremely abundant regionally in the meso-zooplankton size fractions (Hunt et al. 2008). Regional and inter-annual variation in primary production is probably the major determinant of spatial and temporal variability in pteropod densities (Seibel and Dierssen 2003). In the present study, no relationship between pteropod density and Chl a concentration could be found. Thus, factors other than primary production may also play a role in determining the occurrence of pteropods in sea ice regions.

#### Mesh size effects on zooplankton sampling

The mesh size used for plankton sampling is a major factor affecting plankton selection and thus sample composition. In the Southern Ocean, several studies using plankton nets with 100-µm mesh size have indicated that small copepods exceed the abundance, and sometimes the biomass, of larger species (Metz 1996; Atkinson and Sinclair 2000; Dubischar et al. 2002; Schnack-Schiel et al. 2008). Makabe et al. (2012) noted that a 100-µm mesh net is suitable sampling gear to determine meso-zooplankton (200-2,000 µm) community structure in the northern region of Lützow-Holm Bay, although it was not appropriate for copepod nauplii. In their study, the average abundance of zooplankton was 2,664  $\pm$  1,991 inds.  $m^{-3}\!,$  and our data were within this range. In addition, Fukuchi and Tanimura (1981) observed a meso-zooplankton abundance of  $2,495.7 \pm 1,935.2$  inds.  $m^{-3}$  at a fast ice station near St. 52A, a value also similar to Fig. 5 Abundance (inds.  $m^{-3}$ ) and life stage composition of *O*. *similis* (**a**) and *Oncaea* spp. (**b**) at all stations in 2011 and 2012



our own for the fast ice stations. Therefore, the present data seem appropriate for the sea ice regions of the Southern Ocean.

Nonetheless, it is likely that the abundance of some taxa has been underestimated. Over 80 % of planktonic foraminiferans are smaller than  $100 \mu m$ , and only adults

**Fig. 6** Size frequency distribution of foraminifera at northern stations in 2011 and 2012



are larger than 200  $\mu$ m (Berger 1971; Brummer et al. 1986; Spindler and Dieckmann 1986). Similarly, for *O. similis* and *Oncaea* spp., it is difficult to assess the population structure for copepodite stages I(CI) and II (CII) or for copepod nauplius stages, as the body size of these juveniles is often <100  $\mu$ m. Thus, the sampling regime used in the present study cannot fully represent micro-zooplankton and may underestimate the juvenile stages of meso-zooplankton. However, foraminiferans and copepods, including individuals smaller than 100  $\mu$ m, were clearly abundant within the samples at pack ice and open ocean stations where Chl *a* concentrations were higher in both years (Table 3). Thus, despite their underestimate, a clear distribution pattern could be discerned.

The ubiquitous distribution of *Oithona similis* and *Oncaea* spp.

*Oithona similis* and *Oncaea* spp. are considered to be key components of the planktonic food web of the Southern Ocean (Atkinson 1998). They are the most numerically

abundant copepod genera (Hunt and Hosie 2006a, b) and can form a significant proportion of the zooplankton biomass despite their small size (<1 mm). For example, they form up to 20 % of the total copepod biomass in the Weddell Sea (Schnack-Schiel et al. 1998). In the present study, they were the most abundant organisms in all cluster groups (Table 3).

Oithona spp. and Oncaea spp. are known to be omnivorous (Lampitt and Gamble 1982; Turner 1986). O. antarctica has been observed feeding on small copepods (Hopkins et al. 1993). O. similis are able to adapt to the low phytoplankton densities of the permanent open ocean zone (Takahashi et al. 2010). In the present study, Oncaea spp. occurred in high densities in the fast ice area where the Chl a concentration was the lowest. The omnivorous nature of these species may thus allow them to exploit areas of relatively low primary productivity, unavailable to purely herbivorous species. In addition to omnivory, these species have a behavioral strategy which may increase their intake of available microalgae. These copepods have been found to exhibit diel vertical migration synchronized with the sinking of microalgae from melting sea ice (Tanimura et al. 2008). The range of possible food sources and the feeding behavior of these copepods make them highly adaptable and thus lead to their ubiquitous distribution throughout the sea ice region.

Oncaea curvata is regarded as a deep-water species (Hardy and Gunther 1935; Seno et al. 1963). In ice-covered surface waters, Ainley et al. (1986) noted an abundance of crustacean species thought to occur only below 300 m. They suggested that the physical environment, in particular light intensity and quality, immediately beneath the ice was reminiscent of a mesopelagic environment. The pelagic environment in the fast ice area of our study would be similar to that of the deeper water where Oncaea spp. is generally found. The deep-water calanoid copepods, Ae-tideopsis antarctica and Spinocalanus sp. (Schnack-Schiel et al. 2008), were also found only at the fast ice stations in our study (Table 3).

#### The regional distribution of foraminiferans

Cluster analysis clearly separated groups according to the presence or absence of foraminiferans, which were in high abundance at the northern stations, in particular, the pack ice stations. Only one species of planktonic foraminiferans, Neogloboquadrina pachyderma Ehrenberg, lives in Antarctic waters, and the occurrence of this species in Lützow-Holm Bay was noted in the first JARE (Uchio 1960). Lipps and Krebs (1974) reported a high density of N. pachyderma in Antarctic floating ice, although it was not certain whether they were alive or merely preserved in the ice. Further studies by Spindler and Dieckmann (1986) revealed that only juveniles and sub-adults were found in high densities and that they were alive in the sea ice. Subsequently, Dieckmann et al. (1991) showed that foraminiferans in ice cores were mainly found in granular ice and that their numbers were much higher in the pack ice. Foraminiferans attach to frazil ice whose growth result in granular ice using their rhizopods and stay in the sea ice as part of their life cycle in the Southern Ocean (Eicken 1992). The ice community with its high density of foraminiferans is released into the water column as the sea ice melts. In the present study, the northern stations were areas where sea ice was melting or had recently melted. The high densities of foraminiferans found at these stations seem likely to reflect the release of foraminiferans from the ice.

Analysis of the size distribution of foraminiferans tests in sea ice cores has indicated that large living individuals (200–300  $\mu$ m) are found in the lower part of the core (Spindler and Dieckmann 1986). Thus, it seems likely that in areas where the sea ice is in the process of melting the size of foraminiferans would be larger. Indeed, at the pack ice stations (52C and 53C), where the ice was just melting, a greater proportion of larger individuals was recorded. In addition, a trend in foraminiferans size distribution seemed apparent among stations with a greater proportion of smaller individuals at the open ocean stations (Fig. 6). If foraminiferans were released from melting sea ice, it might also be expected that their abundance would increase from the pack ice stations toward the open ocean stations. However, the densities at open ocean stations varied between years, and in 2012, the density at station 53BP was less than half of that at the pack ice station 53C. In addition, the numbers of larger individuals at open ocean stations were markedly less than those at the pack ice stations in both years. Adult specimens are mainly found below 200 m (Spindler and Dieckmann 1986); therefore, larger individuals may have sunk to deeper levels than the 150 m depth from which hauls were taken at the open ocean stations at the time of sampling. Another possibility is that the distribution of foraminiferans was related to Chl a concentration. The polar species N. pachyderma is known to be omnivorous but to exhibit a strong preference for phytoplankton (Hembleben et al. 1989; Lee and Anderson 1991). Bergami et al. (2009) and Takahashi et al. (2010) found correlations between foraminiferans abundance and Chl a concentration. In the present study, the Chl a concentration generally increased from fast ice to pack ice to open ocean stations. Thus, although melting ice and increased phytoplankton abundance may reflect the occurrence of foraminiferans at pack ice and open ocean stations compared with the fast ice stations, it does not explain variations between pack ice and open water stations. Foraminiferans did not show a clear distribution pattern in response to environmental conditions as identified by Hunt and Hosie (2006a) and Swadling et al. (2010).

The dynamics of sea ice around Antarctica is a major factor affecting the distribution and composition of zooplankton communities in these areas. This preliminary study indicates that, like macro-zooplankton, the distribution of micro- and meso-zooplankton varies between fast ice, pack ice and open ocean areas. While in general, O. similis and Oncaea spp. were ubiquitous, being able to adapt to fast ice, pack ice or open ocean, differences were observed in the distributions of copepodite stages compared with adults. Foraminiferans were only found in pack ice and open ocean stations. Foraminiferans have been found in the guts of higher trophic level organisms, such as polychaetes, fish, and tunicates (Hembleben et al. 1989), and they feed on phytoplankton, ciliates, and some copepods (Hembleben et al. 1989; Lee and Anderson 1991). The life cycle of these foraminiferans may be dependent on sea ice. Changes in sea ice dynamics and coverage because of climate change have major implications for the distribution of micro- and meso-zooplankton and for the ecology of the sea ice regions as a whole. Given this and the scarcity of studies of micro- and meso-zooplankton communities in sea ice regions, much more extensive and ongoing research is required in this field in the future.

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