



Meiobenthic colonisation of soft sediments in arctic glacial Kongsfjorden (Svalbard)

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ABSTRACT

An *in situ* experiment on metazoan meiofauna colonisation and succession was carried out at Brandal (78°56.88'N, 11°51.63'E), situated in arctic glacial Kongsfjorden (Spitsbergen). 28 soft sediment containers were deployed at a depth of 20 m and sampled after a one-, two- and three-year immersion period. The main taxonomic groups, abundance of colonising meiofauna and sediment parameters are described and compared. Meiofauna communities at Brandal show the highest densities reported for inner Kongsfjorden to date. While the samples from the three experimental periods and the ambient sediments did not differ in total individual numbers, the treatments showed marked differences in community structure as ascertained by ANOSIM based on Bray Curtis and cosine similarity. An extended three-year period was required until the community stage comparable to ambient sediments was reached. This leads to the assumption that long recovery and colonisation times can be expected for polar shallow water meiofauna.

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1. Introduction

Polar shallow water habitats are frequently affected by natural environmental disturbances. Besides gradients that form in the course of sedimentation and freshwater run-off from tidal glaciers, another factor that may cause serious disturbance of benthic communities is iceberg scouring. A number of specific conditions, such as latitude, depth, local current regimes, geography and site exposure, influence such impacts (Barnes, 1999; Conlan et al., 1998; Dayton, 1990; Dayton et al., 1994; Dowdeswell and Forsberg, 1992; Gutt et al., 1996; Gutt and Piepenburg, 2003; Kotwicki et al., 2004). Physical disturbances both on spatial and temporal scales are thus highly variable. Laudien et al. (2007) assume that for macrobenthic infaunal communities at Kongsfjorden iceberg scouring may be the main structuring force, capable of producing up to 5 m deep plough marks (pers. observ.; Barrie, 1980) in water depths down to 40 m (Dowdeswell and Forsberg, 1992). On the other hand, increasing temperatures due to global warming lead to enhanced glacial retreat, thus creating new, ice-free areas providing a pristine habitat for the settlement of benthic organisms (Cook et al., 2005). Knowledge of colonisation by meiobenthic assemblages is crucial for an understanding of early community development when new substrate becomes available.

As community development is an extended process in polar environments (Barnes, 1996), a study of the successive colonisation should cover several years of monitoring. Therefore, the results of *in situ* subtidal

soft-bottom meiofauna colonisation experiments and recovery studies carried out in temperate regions (Chandler and Fleeger, 1983 and references therein; Schratzberger et al., 2002) cannot be applied to polar environments. Recently Lee et al. (2001) described recolonisation by meiofauna after iceberg scouring at Signy Island (Antarctica). However, no information on the recovery dynamics is available from the Arctic as yet.

Artificial substrata have commonly been used in studies of settlement, recruitment, colonisation and community development (Rumohr and Arntz, 1982; Sarnthein and Richter, 1974; Stanwell-Smith and Barnes, 1997). While reducing substratum heterogeneity, these substrata facilitate replication that is essential for analyses of variation (e.g. between treatments). The provision of artificial soft-sediments in our study at Brandal, Kongsfjorden (Spitsbergen) allowed for the simulation of natural conditions found in newly exposed areas. We therefore assumed that the sediments used in the experiment should not be enriched with organic material as compared to surrounding pristine sediments.

The field experiment was aimed at investigating the long-term colonisation by meiofauna assemblages in an arctic environment in terms of higher taxa abundance over a three-year period to improve the scarce knowledge on long-term community recovery in polar regions.

2. Material and methods

2.1. Study area

An *in situ* colonisation experiment was carried out at Brandal (78°56.88'N, 11°51.63'E, world geodetic system - WGS 84), a site located in central glacial Kongsfjorden (west coast of Spitsbergen, Svalbard,

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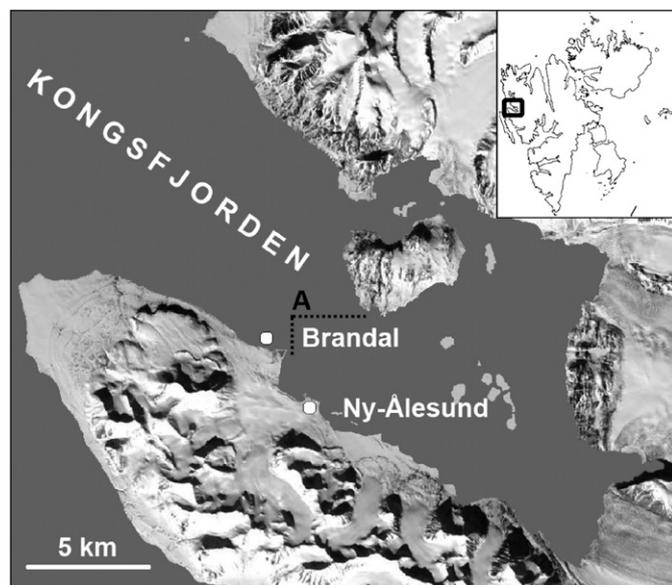


Fig. 1. Landsat TM image of Kongsfjorden at Spitsbergen, the largest island of the Svalbard archipelago (modified from Svendsen et al., 2002). The study site at Brandal, the village Ny-Ålesund and transect A of Dowdeswell and Forsberg (1992), where iceberg monitoring took place, are indicated.

Arctic). The study area is situated on the northeastern fringe of the Brøgger Peninsula, which forms the southern coast of Kongsfjorden (Fig. 1). The latter is 20 km long with a width varying from 4 km to 10 km at the mouth between Kvadehuken and Kapp Guisez. The maximum depth is nearly 350 m with an average depth of 200 to 300 m. The fjord system is directly connected with the North Atlantic Ocean via the Kongsfjord-Renna trough (Bluhm et al., 2001; Jørgensen and Gulliksen, 2001; Svendsen et al., 2002).

Semidiurnal tides of the fjord system range between 1.5 and 2 m and only weak currents prevail. The mean sea surface temperature is just above 0 °C, while maximal values may reach 6 °C in summer. The temperature at the 20 m isobath is 3.6 °C (Bluhm et al., 2001). During summer the 34 isohaline is located at a depth of 5 m. During the winter Kongsfjorden is commonly covered with sea ice, while icebergs and growlers are present throughout the year (Dowdeswell and Forsberg, 1992). A comprehensive description of the physical environment can be found in Svendsen et al. (2002) and ecological information on Kongsfjorden is broadly compiled in Hop et al. (2002).

The widespread, apparently uniform soft bottom off Brandal (Fig. 1) declines gradually within the first 50 m from the coast, followed by a steeper slope. Sediments consist of a sand-clay mixture of quite homogeneous granulometric characteristics throughout the fjord (Włodarska-Kowalczyk and Pearson, 2004; Kotwicki et al., 2004). For a nearby station (<1 km distance) M. Zajackowski (unpublished data, fide Włodarska-Kowalczyk and Pearson, 2004) estimated a POC/PON value of 7.6, indicating the presence of fresh detritus. Irregular boulders transported via ice are present providing settlement substrate for hard-bottom fauna and flora (e.g., *Acrosiphonia* aff. *flagellata*, *Laminaria digitata* and *Phycodrys rubens*) in the soft-bottom habitat (Whittington et al., 1997; Lippert et al., 2001).

2.2. Experimental set-up

Conical terracotta containers with an upper diameter of 43 cm, a lower diameter of 40 cm and a height of 40 cm were installed by SCUBA divers at a depth of about 20 m in June 2002. Due to the adequate spacing between containers (>1.5 m), the replicates were considered to be independent (Conover, 1980). Round containers were chosen in order to keep edge effects equal (Foster, 1975). The bottoms of the 28 numbered containers were replaced by a glass-fibre reinforced plastic

mesh (1 mm) to allow vertical exchange processes. For the installation of the containers an open metal cylinder of the respective size was first pushed into the sediment in order to prevent sediment from sliding into the hole, which was then excavated with a small blade. The terracotta containers were placed in the pit and the cylinder was removed. A low 5 cm fringe of the containers remained above the surrounding sediment preventing it from entering into the experimental unit and avoiding a bias which would be caused by edges elevated higher over the benthic boundary layer. Since the walls of the containers provide a secondary hard substrate that may attract species not common on soft bottoms (Arntz, 1981), the containers were filled to the brim with pre-watered artificial sediment by the SCUBA diver. A 80:20 mixture of clean sand (fine and medium grain size) and clay (Calcium-Bentonit "Ibeco Agromont AWI", IKO Minerals GmbH, Marl, Germany) resembling the natural grain size distribution was used as it was difficult to sterilise large quantities of natural sediment and to remove the organic material. The remaining organic material could have caused unforeseeable effects on the settling fauna (Brunswig et al., 1976). A comparable artificial sediment has been successfully implemented in temperate waters (e.g. Sarnthein and Richter, 1974; Rumohr and Arntz, 1982). To avoid untimely initial colonisation while successively installing the containers, they were covered with plastic sheets until their simultaneous exposure (11 July 2002).

2.3. Sampling and sample treatment

In the following three years samples for meiofauna and sediment analyses were collected from randomly allotted containers (Table 1). Round plastic tube corers with a diameter of 4.5 cm (area: 15.90 cm²) were used to take one sample for meiofauna (in order to guarantee true replicates) and two to three samples for sediment analyses per container. Because the same experimental set-up was used for sampling macrofauna as well, the tube corers were not placed centrally in the containers. Nevertheless, a distance of at least 5 cm to the rim of the containers was always maintained. The uppermost 3 cm of the substrate provided were sampled.

For meiofauna analyses the sediment was preserved in a borax-buffered formaldehyde-seawater solution to a final concentration of 4%. The samples for sediment analyses were stored at -80 °C.

After each sampling, the remaining substrate in the corresponding container was completely removed by a SCUBA diver and replaced by fresh artificial sediment in order to start a new exposure series. Differences in the first-year primary succession between 2003 and 2004 were able to be tested by means of this procedure (Table 1). For comparison with the natural situation 5 sediment cores were taken outside the containers in 2005.

The fixed meiofauna samples were washed with tap water through a 40 µm mesh sieve. Meiofauna and organic material were extracted from remaining sand particles by centrifugation with a colloidal silica polymer (H. C. Stark, Levasil 200/40%, $\rho = 1.17$) as the flotation medium and kaolin to cover heavier particles (McIntyre and Warwick, 1984). Centrifugation was repeated three times at 4000 rpm for five minutes respectively. After each centrifugation the floating matter was decanted and rinsed with tap water. The supernatant containing the meiobenthic organisms was thereafter stained with Rose Bengal before manually sorting to a higher taxon level using a Leica MZ 12.5 stereo microscope. While specimens of most groups were stored in 70% ethanol, copepods were subsequently transferred to glycerine. Nematodes and nauplii were quantified as well but were left in the remaining sample. The

Table 1

Meiofauna samples from soft sediment containers at Brandal, Kongsfjorden: sampling date and number of replicates for different treatments and ambient samples

Treatment	No. of replicates	Date
one-year exposure	2	18./19.06.2003
	3	27./30./31.08.2004
two-year exposure	3	01./02./03.09.2004
three-year exposure	3	02./06./07.09.2005
ambient	5	14.10.2005

Table 2
Median (Med) total organic matter (TOM; % of sediment dry mass), chlorophyll a and phaeopigment content of sediments from three different treatments from Brandal, Kongsfjorden, Spitsbergen (minima (Min) and maxima (Max) included)

Exposure	1-year			2-year			3-year		
	Med	Min	Max	Med	Min	Max	Med	Min	Max
Sediment parameters									
TOM [% dm]	2.034	1.496	2.592	1.895	1.603	2.721	3.069	2.539	3.189
Chlorophyll a [$\mu\text{g mg}^{-1}$]	0.161	0.144	0.266	0.223	0.168	0.318	0.239	0.226	0.287
Phaeopigments [$\mu\text{g mg}^{-1}$]	0.101	0.068	0.390	0.159	0.063	0.210	0.400	0.288	0.409

All data from experimental units sampled in 2004 and 2005.

Table 3
Median density (Med) of major taxa and total individual numbers (Ind. 10 cm^{-2}) for three different treatments and the ambient control samples from Brandal, Kongsfjorden, Spitsbergen (minima (Min) and maxima (Max) included)

Exposure	1-year			2-year			3-year			ambient		
	Med	Min	Max	Med	Min	Max	Med	Min	Max	Med	Min	Max
Individuals [10 cm^{-2}]												
Nematoda	693.7	277.4	869.8	550.3	386.8	588.1	991.2	675.5	2087.4	1325.8	539.6	2158.5
Copepoda	477.4	244.7	556.6	431.4	341.5	544	287.4	234.6	298.7	120.8	31.4	184.3
Copepod nauplii	291.8	73.6	976.7	192.5	157.2	199.4	83.0	79.2	89.9	87.4	31.4	106.3
Annelida	77.4	35.8	214.5	38.4	28.9	45.3	25.8	2.5	44.0	25.2	9.4	43.4
Halacarida	11.9	5.7	49.1	13.8	9.4	23.3	3.8	3.1	3.8	3.8	1.3	12.6
Cumacea	0.6	0	17.6	0	0	3.1	0.6	0	2.5	1.9	0	3.1
Bivalvia	9.4	3.1	23.9	11.9	2.5	11.9	23.3	17.0	29.6	21.4	6.9	92.5
Amphipoda	0.6	0	0.6	0	0	0	0	0	1.3	0.6	0	3.1
Loricifera	0	0	1.3	0	0	0	0	0	0	0	0	0
Ostracoda	6.9	1.9	12.6	1.9	1.9	3.1	2.5	1.9	6.9	2.5	1.3	6.3
Rotifera	0	0	0.6	0	0	0	0	0	0	0.6	0.6	1.9
Tardigrada	0	0	0	0	0	0	0	0	0	0	0	1.3
Priapulida	3.8	1.3	27.7	6.9	5	6.9	0.6	0.6	5.0	1.3	0	5.7
Indet.	3.8	0	48.4	0	0	15.1	0	0	0	0	0	5
Total individuals	1576.7	764.2	2544	1220.8	1081.8	1318.2	1383	1088.1	2530.8	1613.2	645.9	2452.8

individual numbers were counted and abundance values taken for further analyses and interpretation.

The frozen sediment samples were sliced (0–1 cm, 1–2 cm, 2–3 cm) and dried (50 °C until constant weight). The slices were analysed as to their content of organic material (combustion in a muffle furnace, 500 °C for 7 h) and chloroplastic pigments (fluorometric determination after acetone extraction, Turner Fluorometer TD 700). The results for the first centimetre of the sediment were used for data analysis.

2.4. Data analysis

The data were not distributed normally, therefore no arithmetic means and standard deviations could be calculated for the different treatments. If outliers are presumed and in the event of skewed distributions, these values are replaced by the median and the median deviation (Sachs, 1993). In this case we added minima and maxima to the median values in order to demonstrate the distribution ranges of our meiofauna and environmental data.

Non-parametric Kruskal-Wallis one-way analysis by rank was applied to individual numbers and dominance D-values in order to test whether there were significant differences between the samples of the four groups (one-year, two-year, three-year and ambient sediments).

Multivariate meiofauna community analyses were carried out on square root transformed abundance data of the higher taxa recorded using the PRIMER v6 package (Clarke and Gorley, 2006) and the Paleontological Statistics Software Package PAST (Hammer et al., 2001).

Bray-Curtis and cosine similarity analyses were conducted for the square root transformed abundance data and visualised with the aid of non-metric multi-dimensional scaling MDS. The cosine similarity is a more “qualitative” measure as compared to the Bray-Curtis similarity, which discriminates more “quantitatively” between samples (Pfeifer et al., 1998). Overall, the cosine similarity reveals the structural properties of communities more than the Bray-Curtis similarity, which has the major drawback that it separates “similar” samples if the difference is mainly due to multiples in abundance (Pfeifer et al., 1998). For this reason we selected the cosine similarity as a counterpart to the widely used Bray-Curtis similarity for our analyses.

Different factors (i.e. exposure time, sampling year) were applied for the analyses of similarities ANOSIM (one-way analysis) of the outcomes of both similarity indices in order to have a control tool for unexpected events that might occur in one year but not in the following. Thus differences could be detected independently from the starting year of the experiment.

For the interpretation of the relation of meiofauna with sediment factors a canonical correspondence analysis (CCA) was chosen, as this analysis enables comparison of animal data and environmental data in the same plot. CCA was carried out according to Ter Braak (1986) using the statistical package MVSP 3.1. No data transformations were carried out because every transformation bears the risk of bias.

3. Results

3.1. Sediment parameters

Sediment factors differed slightly between the three experimental periods (Table 2). The given minima and maxima indicate a high

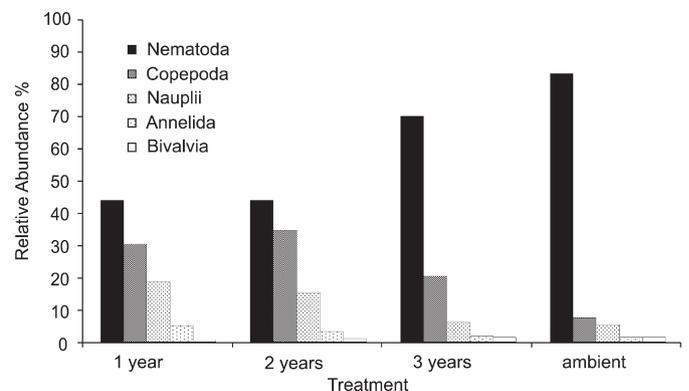


Fig. 2. Relative median abundances of meiofauna higher taxa and nauplii from sediments exposed for one, two and three years as well as ambient sediments from a succession experiment at Brandal, Kongsfjorden.

variability of total organic matter, chlorophyll a and phaeopigment contents. Nevertheless, increased organic matter and the accumulation of phaeopigments in the third year of sediment exposure showed that there were detectable changes in the food supply for meiofauna throughout the duration of the experiment.

3.2. Meiofauna abundances

Animals belonging to twelve higher taxa colonised the ambient sediment as well as the sediment provided independently of exposure time (Table 3; for complete original data set see: Veit-Köhler et al., 2006). Total meiofauna abundance varied only slightly between treatments (minimum in two-year exposure, median: 1220.8 ind./10 cm²; maximum in ambient samples, median: 1613.2 ind./10 cm²). This is confirmed by the non-parametric Kruskal-Wallis test by ranks, which

did not show any significant differences in total individual numbers between the samples ($H=1.735$, $p=0.6291$). The differences between the dominance D measures for the meiofauna higher taxa in the samples, on the other hand, are highly significant ($H=13.61$; $p=0.003484$), thus emphasising a shift in taxon composition with exposure time.

Throughout the experimental time (Fig. 2) it became obvious that nematode abundances were clearly higher than those of the other meiofauna groups in the three-year and the ambient samples. They comprised 44% of the total individuals in the sediment exposed for one year, 44.1% and 69.9% in the sediments exposed for two and three years respectively and 83.3% in the ambient samples (Fig. 2). The numerical importance of copepods and their nauplii diminished with exposure time as well as the relative abundances of annelids. Only bivalves showed slightly increasing values. The rise in relative abundances

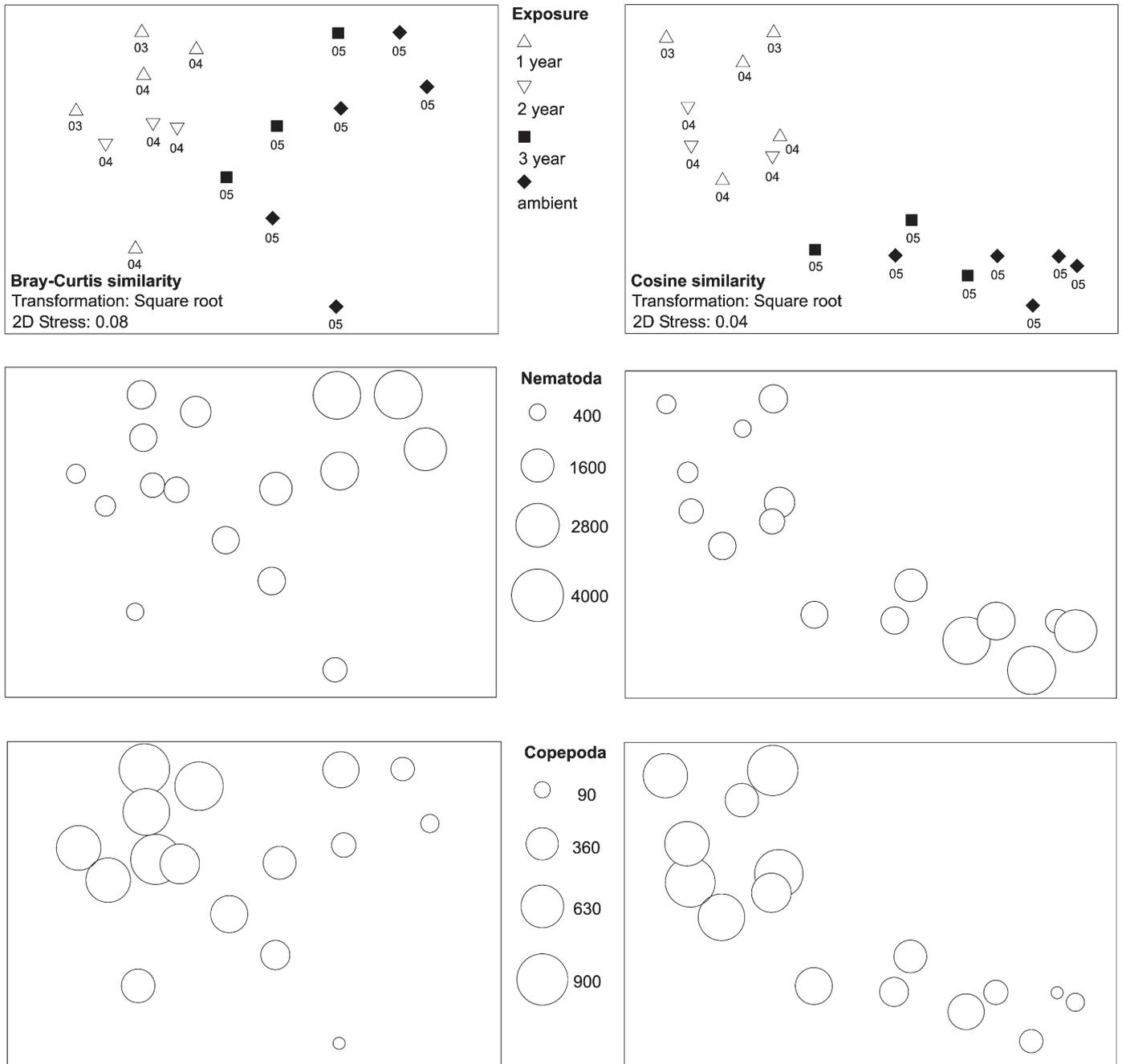


Fig. 3. MDS of the Bray Curtis (left side) and cosine similarities (right side) of the square root transformed meiofauna community data from sediments exposed for one, two and three years as well as ambient sediment samples. Numbers close to station labels indicate sampling year. Nematode and copepod densities are shown as bubble plots.

of nematodes was not only caused by the substantial increase in absolute nematode numbers, but also by the decreases in individual numbers of copepods, nauplii, and annelids.

Nematodes were numerically dominant in all but two samples. Their median abundances reached from 550.3 ind./10 cm² (two-year exposure) to 991.2 ind./10 cm² (three-year exposure) and ambient median values of 1325.8 ind./10 cm². Copepods including adults and copepodids were numerically the second important group (one-year exposure, median: 477.4 ind./10 cm²; ambient, median: 120.8 ind./10 cm²) followed by copepod nauplii, which outnumbered copepods only in three samples. More than 99% of the benthic copepods were Harpacticoida. The total percentage of all remaining taxa did not exceed 7.1%; Annelida were still fairly abundant whereas Halacarida, Bivalvia, Ostracoda, Cumacea and Priapulida (larvae) showed lower individual numbers. Amphipoda, Loricifera, Rotifera, and Tardigrada were represented only rarely and in low numbers in our samples.

3.3. Similarity analyses

Numbers of nauplii may be more subject to seasonal changes than the rest of the copepods represented by copepodids and adults due to the shorter time span spent in this stage. As sampling dates inevitably varied throughout the years (Table 1), we removed nauplii from the data set before statistical tests were carried out. Especially the samples collected in June 2003 had very high naupliar abundances as compared to the samples of the other years taken later in the season.

The MDS resulting from the more “quantitative” Bray-Curtis and the more “qualitative” cosine similarity analyses are presented in Fig. 3. Although the Bray-Curtis similarity separates one- and two-year samples from three-year and ambient samples spatially, it is obvious that scatter within both groups is high (Fig. 3, graphs on the left). There are distances between ambient samples that are of a similar magnitude as distances between ambient and one-year samples. The same is true for the MDS of the cosine similarity, though the separation in this ordination plot is clearer and distances within the groups are smaller (Fig. 3, graphs on the right). Regarding the bubble plots for Nematoda and Copepoda of both MDS plots, the difference between the two similarity indices becomes obvious: the Bray-Curtis similarity only ordines according to individual numbers, which is shown by the continuous change in bubble size over the array of the plot. The more variable distribution of bubbles of different sizes in the cosine similarity plots follows the characteristic of the index in grouping samples with comparable relative individual numbers rather than samples with similar absolute individual numbers.

As expected, ANOSIM of the Bray-Curtis and the cosine similarity led to different outcomes (Table 4). According to an analysis of the Bray-Curtis results, the global R value for the tests for exposure time (one-, two-, and three-year treatments and ambient sediments) and the factor sampling year (2003, 2004, 2005) independently of treatment were not significant. ANOSIM of the cosine similarity values, by contrast, showed clearly higher global R values. The grouping for exposure time showed that the different treatments and the ambient data are separated quite well. Additionally, the cosine similarity data of grouped samples from different sampling years are clearly dis-

Table 4

Results of ANOSIM for the Bray-Curtis and cosine similarity applied to meiofauna data from three different treatments and ambient control samples from Brandal, Kongsfjorden, Spitsbergen (*groups well separated; **groups clearly distinguishable)

ANOSIM	Bray Curtis	Cosine
Test for factor grouping	Global R	Global R
Exposure time (1-, 2-, 3-year, and ambient)	0.37	0.67*
Sampling year (2003, 2004, 2005)	0.46	0.83**
1-year exposure 2003 vs 2004	-0.17	0.42
1-year exposure 2004 vs 2-year exposure 2004	0.22	-0.07
3-year exposure 2005 vs ambient samples 2005	-0.03	0.26

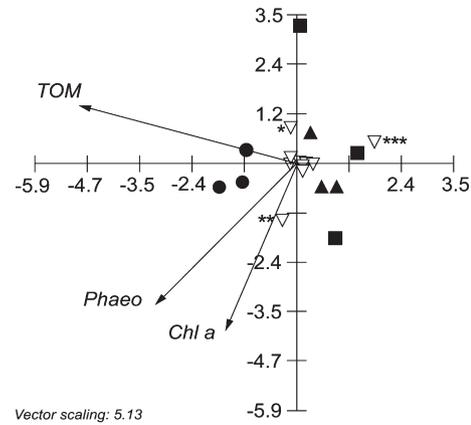


Fig. 4. Canonical correspondence analysis of samples from sediments exposed for one (black squares), two (black triangles) and three years (black dots), and meiofauna taxa (white triangles; *Cumacea, **Amphipoda, ***Rotifera) according to environmental data (vectors: total organic matter – TOM; chlorophyll a – Chl a; phaeopigments – Phaeo).

tinguishable with global R=0.83. Both similarity indices indicated that the grouping of samples by sampling year represents reality better than the grouping by exposure time.

Detailed analyses of part of the data groups show that there are no inter-annual differences in first-year primary succession between sediments exposed for one year from 2003 and 2004. Additionally, sediments exposed for one and two years collected in the same year (2004) do not show a group affiliation at all, nor is there any difference between the three-year and ambient samples from 2005.

3.4. Canonical correspondence analysis

A canonical correspondence analysis (CCA; Fig. 4) for 12 variables (meiofauna taxa) and 9 cases (3 replicates per treatment) was performed for the environmental data with 3 variables (total organic matter – TOM; chlorophyll a – Chl a; phaeopigments – Phaeo). The ordination diagram of CCA displays samples and animal data as points and environmental variables as vectors (Fig. 4). The vector for TOM decreases with the first axis (x), whereas Chl a and Phaeo decrease with the y axis. For data interpretation the vectors can virtually be produced through the intersection of the axis. The ordination of data points in relation to environmental vectors is determined by projecting the data points at a right angle to the corresponding vector (Ter Braak, 1987).

The CCA shows that the longer the exposure time, the more organic matter and chlorophyll degradation products, as represented by the phaeopigments, are accumulated on the sediment. While the replicates from the one-year exposure deviate very significantly from each other, two- and three-year replicates are more homogeneous within their groups. The CCA indicates that there is no significant correlation of most of the animal taxa with any of the environmental factors, though the very rare Cumacea, Amphipoda and Rotifera seem to have habitat preferences that are distinct from most of the other taxa.

4. Discussion

4.1. Meiofauna and sediment factors

The individual densities of the meiofauna higher taxa found at the 20 m deep experimental site off Brandal are not in line with the abundances reported by the only previous meiofauna study on subtidal sites in Kongsfjorden by Kotwicki et al. (2004). Although the first 5 cm of the sediment were sampled in their study (Van Veen grab; three subsampled cores with 3.8 cm² surface each unified as a single sample per station, no information on replicates and calculation

of median or mean per station) as compared to our experiment where only the first 3 cm were analysed, they only recorded two stations with meiofauna densities above 1000 ind./10 cm². Both stations are located in the part of the fjord defined as the “outer” region at 75 m and 265 m depth, respectively. All shallower stations and all stations from the “inner” part of the fjord do not exceed the numbers found off Brandal at a depth of 20 m. Brandal is situated at the western boundary of the inner fjord, which according to their findings is very poor in meiofauna densities, and opposite Blomstrandoya island, from which they report comparably high abundances. The discrepancy found is most certainly due to the fact that Kotwicki et al. (2004) sampled from a depth of 44 to 350 m and their stations always showed a silt and clay content of at least 86%. This is contrary to Brandal, where we found 80% fine and medium sand and only 20% silt and clay. Additionally, the different sampling techniques (Van Veen grab versus corers placed by scuba diver) might play a role. On the other hand, it is obvious from their data that the content of organic matter cannot play a crucial role in meiofauna abundances on a larger scale for Kongsfjorden. Our results support this in that the TOM in our case is only about a half to a third of their highest measurements. Therefore the contrasting densities of Brandal and the deeper regions are most certainly due to the differences in sediment grain size.

When comparing the meiofauna abundances encountered in Kongsfjorden and off Brandal to Antarctic polar shallow water soft-bottom areas, the special low-density situation of the Arctic sites becomes obvious: de Skowronski and de Corbisier (2002) report up to threefold higher total meiofauna abundances from King George Island (South Shetland Islands, Antarctica) and Veit-Köhler et al. (2008) even reported a maximum value of 16845 ind./10 cm² for the same island (0–5 cm sediment depth). Vanhove et al. (2000) report very high but highly variable meiofauna densities from Signy Island (South Orkney Islands, Antarctica) from a monitoring site over a period of 19 months. Only Lee et al. (2001) encountered densities at the control sites of an iceberg scour time series that are comparable to our values.

The very high copepod relative abundances (20.3–34.6%) within the experimental containers are exceptional for polar shallow areas even in the samples of the three-year treatment. In our experiment only the ambient sediments revealed relative abundances, e.g. 7.6% for copepods, comparable to Kotwicki et al. (2004) who found values between 0 and 10%. This supports our findings that the developing meiofaunal communities in our experiment are not similar to ambient values. The same is true for the Antarctic communities: de Skowronski and Corbisier (2002) found an average of only 9.5% Copepoda in total meiofauna. Veit-Köhler et al. (2008) report copepod percentages that never exceeded 2.8% of the total soft-bottom meiofauna in 24 single core samples. The figures determined by Lee et al. (2001) with 2.3% copepods in their control samples are also comparably low.

Overall, we found high meiofauna densities at Brandal as compared to other sites in Kongsfjorden that can be explained only partly by food availability, but most certainly by water depth, sediment stability, sedimentation processes near the glacier (Kotwicki et al., 2004) and therefore sediment grain size. But as compared to Antarctic soft bottom shallow water sites, the densities found at Brandal are quite low. This is most probably due to differences in total organic matter, which at Brandal is low for polar shallow waters, i.e. 1.49% to 3.19% of the sediment dry mass.

4.2. Colonisation and community development

Colonisation of sediments by meiofauna is generally a rapid procedure. This has been shown in several studies (reviewed by Coull and Palmer, 1984), but the remaining question for the Arctic is whether the newly establishing communities are comparable to the naturally existing ones in the surrounding sediment.

Settlement itself is controlled by several processes and three major mechanisms have been identified: locomotion, dispersal by water

currents and larval colonisation. The three methods act on different time scales (Peck et al., 1999).

While species with highly motile stages may invade the substrate actively, other groups depend on external forces like currents or recruitment via meroplanktic larvae. Colonisation by recruits in our case is unlikely as meiofauna taxa do not have planktonic larval stages. The observed increase in nematode abundance paralleled by a decrease in copepod abundance (Fig. 2) can thus be explained by the behaviour of the taxonomic groups: While the benthic Harpacticoida include several taxa capable of leaving the sediment and actively swimming small distances (Armonies, 2000; Thistle and Sedlacek, 2004), nematodes are exclusively infauna and mainly colonise sediments by active lateral migration (Schratzberger et al., 2004b). A laboratory experiment revealed that all specimens actively leaving the Brandal sediments when exposed to darkness were crustaceans (own unpublished data). Therefore, it must have been easier for actively swimming benthic copepods to surmount the container walls and settle inside. Adult sediment-dwelling copepods and nematodes may have colonised the sediment after resuspension and settlement events (Thistle et al., 1995; Schratzberger et al., 2004a) and then increased in abundance by reproduction.

Nematoda and Copepoda, the two most abundant taxa in meiofauna samples, are strongly correlated with food supply, among other environmental factors. Throughout our experiment food in terms of amount of organic matter and sedimented primary production accumulated in the sediment and likewise changed communities. Since the fluxes to the sediment may differ from year to year, the restart of the experiment in 2003 and the introduction of “sampling year” as a factor in our statistical analyses helped to interpret our findings.

While exposure time does not seem to be the most important factor for the state of the meiofauna assemblage, sampling year definitely is. However, the discrepancies detected by the similarity analyses using the different grouping factors “exposure time” versus “sampling year” did not contradict the idea of a temporally maturing meiofauna assemblage on a scale of years. A certain dependence of “exposure time” and “sampling year” in our data set was technically unavoidable. Nevertheless, we can affirm that the meiofauna assemblages of our experiment had reached a state closely related to ambient communities only after three years of exposure. The resemblance of three-year and ambient samples contrasts with observations on the establishment of macrozoobenthic communities within the same containers (Laudien and Sahade, unpubl. data) that did not reach the mature stage after three years of development. It has been shown that colonisation of settlement substrates by macrofauna in Antarctica is at least an order of magnitude slower than at temperate or tropical sites (Barnes, 1996; Stanwell-Smith and Barnes, 1997).

With regard to one-off studies our experiment underlines that generalisations from spatially or temporally single-spot samplings should be made cautiously and an appropriate number of replicates should be taken. Firstly, the sampled spots might not be representative for the whole area and secondly they might be in a transitional non-reproducible stage.

4.3. Disturbances and recovery

Kotwicki et al. (2004) discuss the influence of environmental disturbance on meiofauna communities in the inner Kongsfjorden. They find glacial runoff in terms of freshwater input and enhanced sedimentation rates to be one reason for low individual numbers. Obviously the Brandal experimental site does not fit into their scheme since it is as diverse and populated as their station 20 at a depth of 265 m in the outer bay. Although Brandal represents a very shallow depth not sampled by them (20 m), it is even more astonishing that melt water and glacial runoff should not negatively influence the meiofauna assemblages at this site.

Polar shallow water systems are affected by environmental changes first and act as small-scale laboratories (Dayton, 1990). Iceberg scouring is likely to be one of the most common disturbances in those benthic habitats. In fact, Laudien et al. (2007) related macrobenthic community differences of the same habitat to scouring. The only existing study recording shallow water meiofauna recovery after iceberg impact stems from the Antarctic (Lee et al., 2001). The latter found the data difficult to interpret because a rapid return of the meiofauna after the impact was then followed by a decrease in abundances. As the distribution of our data indicates a high variability between replicates (Table 3), we can only again stress the importance of enhanced replicative sampling.

Meiofauna may be less sensitive to sediment disturbance and may react with a more rapid establishment of mature natural communities than macrofauna (Warwick et al., 1990), whose recovery requires many years (Picken, 1985). On the other hand, the lacking structuring effect of macrofauna in our case could be a reason for the skewed colonisation of the provided substrates by meiofauna since the macrofauna itself was far from reaching a stage comparable to the situation outside of the containers (Laudien and Sahade, unpubl. data).

Furthermore, it is clear that barriers like the container walls in our experiment hinder nematodes in their active lateral invasion of newly exposed areas. Schratzberger et al. (2004a, 2004b) showed that lateral migration takes time and does not necessarily culminate in the establishment of assemblages similar to the source communities. According to Chandler and Fleeger (1983), large-scale defaunation of meiobenthic communities recovers more slowly than small-scale disturbances. Nematode colonisation in our case was restricted to a passive water-borne process of dispersal, which would also be necessary for colonisation of larger areas.

Obviously even three years of exposure were scarcely enough to establish a community with relative abundances comparable to the ambient assemblage. This leads us to the assumption that the colonisation of large newly ice-free or defaunated areas by meiofauna, at least in arctic Kongsfjorden, does not occur on the same time scale as in temperate regions.

5. Conclusions

- Polar meiofauna communities show extremely long colonisation and recovery times. This is especially relevant for newly ice-free areas and defaunated zones in iceberg scours.
- Brandal represents the highest meiofauna densities at the local scale for inner Kongsfjorden, though compared to other polar shallow water soft sediments, the site is exceptional in its low total meiofauna abundances.
- The experiment revealed total meiofauna individual numbers comparable to ambient sediments after one year of exposure, but high copepod relative abundances characterised the colonising communities even after three years of exposure.
- In addition, the analyses of our data set stress the importance of replicative sampling and the use of adequate similarity indices.

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