Diversity of larger consumers enhances interference competition effects on smaller competitors

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Abstract  Competition between large and small species for the same food is common in a number of ecosystems including aquatic ones. How diversity of larger consumers affects the access of smaller competitors to a limiting resource is not well understood. We tested experimentally how species richness (0–3 spp.) of benthic deposit-feeding macrofauna changes meiofaunal ostracods’ incorporation of fresh organic matter from a stable-isotope-labeled cyanobacterial bloom, using fauna from the species-poor Baltic Sea. Presence of macrofauna mostly decreased meiofaunal incorporation of bloom material, depending on the macrofauna species present. As expected, the species identity of macrofauna influenced the incorporation of organic matter by meiofauna. Interestingly, our results show that, in addition, species richness of the macrofauna significantly reduced meiofauna incorporation of freshly settled nitrogen and carbon. With more than one macrofauna species, the reduction was always greater than expected from the single-species treatments. Field data from the Baltic Sea showed a negative correlation between macrofauna diversity and meiofaunal ostracod abundance, as expected from the experimental results. We argue that this is caused by interference competition, due to spatial niche differentiation between macrofauna species reducing the sediment volume in which ostracods can feed undisturbed by larger competitors. Interference from macrofauna significantly reduces organic matter incorporation by meiofauna, indicating that diversity of larger consumers is an important factor controlling the access of smaller competitors to a limiting food resource.

Keywords  Asymmetrical competition · Biodiversity · Complementarity · Resource partitioning · Species richness

Introduction

Understanding how many species can coexist on a few limiting resources has long been a central issue in ecology. Interspecific competition is a primary ecological mechanism influencing the abundance, distribution and coexistence of species, and hence the diversity of ecological communities (Chase et al. 2002; Connell 1983; Schoener 1983). Competition between species can occur through exploitative competition, where negative interactions arise from the removal of a shared resource, or through interference competition, where foragers reduce the capacity of other species to utilize this resource through behavioral mechanisms (Amarasekare 2002; Begon et al. 1990). Although potentially as important as exploitative competition, interference competition is less well understood (Adler and Mosquera 2000; Valeix et al. 2007).

Body size is a trait that commonly differentiates coexisting species, often with important consequences for ecological relationships (Basset and Angelis 2007; De...
Roos et al. 2003). Larger animals tend to benefit from their size, which increases interference cost for smaller species, thus creating asymmetrical competition (Persson 1985), which may result in reduced access to resources and switching to alternative resources by the smaller species (Amarasekare 2003). As a result, interference competition by larger species can play an important role in structuring ecological communities (Lawton and Hassell 1981; Persson 1985). Although some studies have focused on the role of interference by large species in consumer–resource relationship (Basset and Angelis 2007; Valeix et al. 2007), to our knowledge, none have looked at how their species richness influences the intensity of interference competition with smaller species.

Body size-regulated interference competition is probably an important mechanism shaping soft sediment bottom ecosystems, one of the most extensive habitats on Earth (Aljetlawi and Leonardsson 2002; Wilson 1990). Metazoan assemblages in soft sediment bottoms often show a strong dichotomy in size, and have traditionally been divided into macro- and meiofauna, with organisms retained on 0.5- or 1-mm mesh classified as macrofauna, and those passing through this mesh, but retained on a 40-μm mesh, as meiofauna (Giere 2009). Since most soft-bottom sediments are situated below the photic zone, benthic assemblages are often critically dependent on the supply of organic matter from settling phytoplankton blooms (Graf 1992). How interspecific competition regulates the use of this resource by benthic deposit-feeding species is important for understanding their coexistence (Byrén et al. 2006).

The activities of animals in the sediment, such as bioturbation, can dramatically alter the habitat structure and resource availability of soft-bottom communities (Erwin 2008; Meysman et al. 2006). The size of macrofaunal species give them a greater capacity for bioturbation, and thus increase their potential to physically modulate trophic interactions with meiofauna (Tita et al. 2000), shape their habitat structure (Austen et al. 1998) alter their supply of oxygen (Meyers et al. 1987) and interfere with their access to fresh organic matter from settling phytoplankton blooms (Modig et al. 2000; Ólafsson et al. 2005). Additionally, macrofaunal assemblages are generally composed of species that differ in size, feeding and burrowing activity, mobility, and therefore in bioturbation activity. When macrofaunal species differ in their bioturbation activity, there is a potential for macrofaunal species richness to affect the intensity of interference competition on meiofaunal assemblages (Austen et al. 1998). Therefore, soft-bottom sediment assemblages are of particular interest for studies of the relationship between species richness and interference competition.

Our study was performed with Baltic Sea soft sediment communities, which due to the Baltic Sea’s low salinity and young age have a naturally low biodiversity (Elmgren and Hill 1997). The indigenous deposit-feeding macrofaunal community in much of the Baltic sediments below the photic zone is dominated by only three species. This naturally species-poor ecosystem provides an ideal platform for testing experimentally how interference competition by macrofauna influences meiofaunal feeding on a limiting resource, using realistic and ecologically relevant abundances and combinations of species.

We manipulated macrofaunal species richness and composition by testing all ecologically realistic combinations of the three main indigenous depositing-feeding macrofaunal species in the Baltic Sea, as well as a treatment without macrofauna. After experimental labeling of a typical Baltic Sea summer cyanobacterial bloom with stable isotopes of both carbon and nitrogen (Larsson et al. 2001), we measured incorporation of N and C of cyanobacterial origin by the three dominant species of ostracods, the most important meiofaunal group in terms of biomass in the Baltic Sea (Elmgren et al. 1984). We show that the presence of macrofauna greatly reduced the incorporation of fresh organic matter by all three ostracod species, and that this effect was most pronounced at the highest macrofaunal species richness tested (biomass kept constant), indicating that diversity of larger consumers can be an important factor controlling the access of smaller competitors to a limiting resource.

**Materials and methods**

**Experimental design**

To test how macrofaunal species richness and composition affect incorporation of N and C from a freshly settled phytoplankton bloom by meiofauna, we performed an experiment using the three macrofaunal species that dominate the indigenous deposit-feeding macrofauna guild in Baltic Sea sediments: the bivalve *Macoma balthica*, and the two amphipods *Monoporeia affinis* and *Pontoporeia femorata*. These macrofauna species differ in their feeding and bioturbation activities and therefore have the potential to interfere differently with how meiofaunal assemblages process fresh phytodetritus. *Macoma balthica* is semi-mobile, burrowing surface feeder, that uses its siphon to forage on the surface sediment (Ólafsson et al. 2005), while both amphipods are mobile bioturbators, with *M. affinis* predominantly a surface feeder and *P. femorata* a subsurface feeder (Byrén et al. 2006; Lopez and Elmgren 1989). Meiofaunal communities in the Baltic Sea proper are generally dominated in biomass by three ostracod species: *Candona neglecta, Paracyprideis fennica* and *Heterocyprideis sorbyana* (Elmgren et al. 1984). Two different
feeding ecologies are found among these species, with *C. neglecta* feeding preferentially on freshly deposited detritus while *P. feminica* and *H. sorbyana* rely more on older organic matter (Nascimento et al. 2008; Ólafsson et al. 1999). As these three ostracod species do not seem to compete strongly for freshly settled phytodetritus (Modig et al. 2000) and, since sorting live meiofaunal animals is extremely time consuming, we used natural abundances of *C. neglecta*, *P. feminica* and *H. sorbyana* in the experiment (*C. neglecta*: 1.5 ± 0.3 ind. 10 cm\(^{-2}\); *P. feminica*: 11 ± 1 ind. 10 cm\(^{-2}\); *H. sorbyana*: 7 ± 0.9 ind. 10 cm\(^{-2}\); \(n = 3\), mean ± standard deviation, SD).

Our experiment included seven treatments, each with eight replicates, using a substitutive design, where macrofaunal species richness varied (0, 1, 2 or 3) while density and biomass of animals was kept as constant as possible. Our experimental design included: (1) a treatment without macrofauna (NoMac), (2) three macrofauna monoculture treatments with 24 ind. per replicate, one for each deposit-feeding macrofaunal species *M. affinis* (Ma), *P. femorata* (Pf), and *M. balthica* (Mb), (3) a treatment containing a mixture of all three macrofaunal species together (Ma + Pf + Mb), with 8 ind. of each of the three macrofaunal species per replicate, and (4) two treatments with two macrofaunal species (Ma + Pf, and Ma + Mb) with 12 ind. of each species per replicate. We did not include a *P. femorata* plus *M. balthica* treatment since this community composition is rarely found in nature. This design allows the identification of potential interference competition effects on meiofaunal incorporation of fresh organic matter due to macrofaunal species composition and richness.

Phytoplankton collection and labeling

A cyanobacterial bloom dominated by *Nodularia spumigena* was collected at 1–1.5 m depth in the open Baltic proper on 6 July 2006, using a 100-μm plankton net. The cyanobacteria were separated from mesozooplankton using a light trap, as described in Nascimento et al. (2009). After zooplankton removal, the cyanobacterial suspension was incubated for 7 days in the laboratory under constant shaking and illumination at 19°C in f/2 medium, where the inorganic C and N sources were entirely composed of NaH\(^{13}\)CO\(_3\) and \(^{15}\)NH\(_4\)Cl (Cambridge isotopes, 99% heavy isotope), respectively. The \(^{13}\)C- and \(^{15}\)N-labelled cyanobacteria were then harvested by creating a sharp pressure shock to break their gas vacuoles (Walsby 1975), causing them to settle to the bottom, after which they were sieved through a 27-μm sieve and rinsed with filtered brackish seawater to remove non-incorporated \(^{13}\)C and \(^{15}\)N. The harvested cyanobacteria were concentrated to a dense stock suspension and samples of this suspension were taken for stable isotope analysis (SIA).

Sediment sampling

Surface sediment was collected in May 2006 from a depth of 27 m in the northern Baltic proper (58°49′N, 17°31′E) using an epibenthic sled (Blomqvist and Lundgren 1996) set to a depth of 3 cm. The sediment in this area is characterized as silty clay with high water content (87% in the top centimetre) and a C/N ratio of 7.6 ± 0.1 (\(n = 3\)). The sediment was sieved through a 1-mm sieve to remove the macrofauna while retaining the natural abundance of meiofaunal ostracods. After homogenization, 500 cm\(^3\) of the sieved sediment were carefully transferred to each of 56 Plexiglas microcosms (surface area 50 cm\(^2\)), followed by the addition of 450 mL of brackish seawater (6 psu) pumped in from a depth of 15 m near where the sediment was collected, and filtered through a sand filter and a 40-μm sieve.

The sediment was left to settle for 1 day, after which each microcosm was supplied with gentle aeration to avoid anoxic conditions, and left to equilibrate for 4 weeks in the dark at 5 ± 1°C, a temperature similar to that in the field at that depth at this time of year. Before the start of the experiment, individuals of *M. affinis*, *P. femorata* and *M. balthica* were collected from the same location as the sediment using the epibenthic sled, separated using a 1-mm mesh sieve, and stored in the dark in sediment at 5 ± 1°C with aeration until the start of the experiment.

Addition of macrofauna and simulation of the settling of the cyanobacterial bloom

Active individuals of *M. affinis*, *P. femorata* and *M. balthica* of similar shell-free dry mass (*M. affinis*: 1.9 ± 0.4; *P. femorata*: 1.8 ± 0.2; *M. balthica*: 2.1 ± 0.8 mg; \(n = 30\), mean ± standard deviation, SD) were added to the experimental microcosms, to achieve a total density of 24 individual per replicate or 4,800 ind. m\(^{-2}\) which is within the range of abundances usually found in the field (Ankar and Elmgren 1976). Five days after the addition of macrofauna, the experiment was started by adding approximately 16 mL of the harvested cyanobacterial liquid suspension, corresponding to 156 ± 16 mg dry mass (mean ± SD) or 13.7 g C m\(^{-2}\) to the water column of each microcosm. This is equivalent to about 1 month of sedimentation of phytoplankton material during a summer bloom in the area (Heiskanen et al. 1998). The experimental incubation ran for 4 weeks in the dark at 5°C.

Termination of the experiment

Sediment samples were taken from all replicates with cut-off syringes (0.15 cm), which were immediately frozen at –20°C and later sliced in 0.5-cm layers down to 5 cm
depth. Each layer was homogenized, weighed before and after drying at 60°C and sub sampled (15 mg dry mass) for SIA. We did not sample below the top 5 cm of sediment because the amounts of label there are expected to be negligible since no macrofauna and very little meiofauna live in sediments below 5 cm. Without bioturbation, the transport of label to this depth would be insignificant in the timescale of this experiment.

After the sampling with the cut-off syringes, the remaining surface sediment layer (0–1 cm) where 95% of Baltic ostracods normally are found (Ólafsson et al. 1999) in each microcosm was sliced off and sieved sequentially through 1,000- and 40-µm sieves, and the contents of the 40-µm sieve preserved in 4% formalin. The macrofauna retained in the 1,000-µm sieve were put in filtered sea water for 24 h to empty their gut, rinsed in distilled water, dried at 60°C, weighed (bivalve without shell) and put in tin capsules for SIA. The differences in macrofaunal isotope values and incorporation of organic matter among treatments are described and discussed in Karlson et al. (2010).

Ostracod extraction from the sediment

The meiofauna was extracted from the 40-µm sediment fraction using Ludox colloidal silica at a specific gravity of 1.15 (Ólafsson et al. 1999). Ostracods were sorted, counted and identified to species level. After three extractions, the remaining sediment was again sieved through a 160-µm sieve and checked for ostracods under a ×50 binocular stereomicroscope. Extracted ostracods were rinsed in distilled water, sorted, and identified to species level under a ×50 binocular stereomicroscope. Individuals of three ostracod species, Candona neglecta, Heterocyprideis sorbyana, and Paracyprideis fennica, were picked out into a watch glass containing 1 M HCl to remove inorganic C. After 24 h in acid- the ostracods were rinsed in distilled water, placed in tin capsules and dried at 60°C for 24 h in preparation for SIA.

Analyses of elemental content of C and N and their stable isotope ratios in samples of cyanobacteria, ostracods, and homogenized sediment were made at the UC Davis Stable Isotope Facility, USA. For P. fennica and H. sorbyana, 35–45 individuals per replicate of each species were pooled in order to achieve sufficient biomass for reliable SIA (sample range: 0.193–0.680 mg dry mass). For the less abundant C. neglecta, individuals from two replicates sometimes had to be pooled to achieve the biomass needed for SIA (sample range: 0.106–0.283 mg dry mass).

All stable-isotope values are given in the δ notation where:

\[
\delta^{15}N \text{ or } \delta^{13}C \text{ (‰)} = \left[ \frac{R_{\text{sample}}}{R_{\text{standard}}} - 1 \right] \times 10^3
\]

where \( R \) is \((^{15}N/^{14}N \text{ or } ^{13}C/^{12}C)\).

To quantify amounts of cyanobacterial N and C incorporated by the three ostracod species and stored in the sediment, a linear 2-source mixing model was used for N and C separately (Karlson et al. 2010):

\[
f_1 + f_2 = 1; \quad f_1 = (\delta_{\text{sample}} - \delta_{\text{source2}})/(\delta_{\text{source1}} - \delta_{\text{source2}})
\]

where \( f_1 \) is the proportion of new N or C (of cyanobacterial origin) in the sample (animal or sediment), \( f_2 \) is the proportion of aged N or C (N and C sources in the initial sediment), \( \delta_{\text{source1}} \) is the isotope value of the added cyanobacteria and \( \delta_{\text{source2}} \) the isotope value of the experimental sediment. Uncorrected isotope data were used in the mixing model, since species-specific differences in fractionation (Goedkoop et al. 2006) and fat content (Post et al. 2007) were negligible compared to the strong labeling. This model was used to calculate the proportion of N and C of cyanobacterial origin in each ostracod sample. This value was then extrapolated to the total biomass of each ostracod species in each replicate, to obtain the total of N and C incorporated by the ostracods as a community.

The amount of N and C of cyanobacterial origin in the sediment was also calculated from this mixing model, by summing the N and C content of all layers (Karlson et al. 2010).

Expected incorporation values (\( I_{\text{exp}} \)) of cyanobacterial N and C by the ostracod community in the multi-macrofauna species treatments (Ma + Pf, Ma + Mb and Ma + Pf + Mb) were calculated by subtracting the incorporation by ostracods in each replicate of the macrofauna monocultures (\( I_{\text{a}} \)) from the average incorporation in the treatment without macrofauna (\( I_{\text{NoMac}} \)). This enables the calculation of how much incorporation by ostracods was reduced by the presence of each macrofauna species (\( T \)). Assuming that each of the macrofauna individual of the same species causes approximately the same \( T \) on ostracod incorporation, it is possible to calculate \( T \) per macrofauna individual for each macrofauna species (\( T_{\text{ind}^{-1}} \)) by dividing \( T \) by the number of macrofauna individuals present in the corresponding replicate of the macrofauna monocultures. Multiplying \( T_{\text{ind}^{-1}} \) by the number of corresponding macrofauna individuals in the mixed treatments, will give the expected \( T \) of each macrofauna species within the mixed treatment (\( T_{\text{a}} \)). Then \( T \) for the a mixed treatment would be expressed as:

\[
T_{\text{mixed treatment}} = T_{\text{species a}} + T_{\text{species b}}
\]

It is then possible to determine expected incorporation in the mixed treatments (\( I_{\text{exp}} \)) by subtracting \( T_{\text{mixed treatment}} \) from \( I_{\text{NoMac}} \).

\[
I_{\text{exp}} = I_{\text{NoMac}} - T_{\text{mixed treatment}}.
\]
Net effects of macrofaunal species richness on incorporation of cyanobacterial C and N were calculated as the differences between observed and expected incorporation values in multi-macrofaunal species treatments.

Statistics

The effects of both species richness and species composition on interference were tested in a nested ANOVA. Macrofaunal species composition was set as a factor nested under species richness with interference on ostracod community incorporation of N or C as the dependent variable. The differences among treatments in N or C content of cyanobacterial origin within and below the first centimetre of sediment at the end of the experiment were tested with one-way ANOVA. Cochran’s test was used to check the assumption of homoscedasticity and when necessary the data were log-transformed to obtain homogeneity of variance. Paired a posteriori comparisons were carried out using Tukey HSD. To test if the observed interference on ostracod community incorporation of N and C was significantly different from the predicted, a single mean $t$ test was performed for each of the multi-macrofaunal treatments, for both N and C. Potential differences in survival of macrofaunal species were tested using non-parametric Kruskal–Wallis.

Results

Cyanobacterial bloom composition and labeling

The cyanobacterium *N. spumigena* made up 97% of the phytoplankton biovolume, with diatoms *Amphora* spp. and resting cells of the cyanobacterium *Anabaena lemmermannii* accounting for the rest. After labeling with NaH$^{13}$CO$_3$ and $^{15}$NH$_4$Cl, the isotope values (mean ± SD) of the cyanobacterial suspension increased from $-1.4 ± 0.4$ to $906 ± 32$ for $\delta^{15}$N and from $-23.7 ± 0.2$ to $157 ± 6$ for $\delta^{13}$C.

| Treatment | Macrofaunal final abundance (ind. 50 cm$^{-2}$)/biomass (g dw$^{-1}$) | Ostracod final abundance (ind. 50 cm$^{-2}$)/biomass (mg dw$^{-1}$) |
|-----------|-------------------------------------------------|-------------------------------------------------|
| NoMac     | $-4$                                           | $97 ± 9/0.85 ± 0.2$                              |
| Ma        | $14 ± 0.8/28 ± 2$                              | $81 ± 7/1.0 ± 0.1$                               |
| Pf        | $13 ± 0.2/26 ± 1$                              | $96 ± 7/1.3 ± 0.2$                               |
| Mb        | $21 ± 0.5/32 ± 0.6$                            | $89 ± 8/1.1 ± 0.2$                               |
| Ma + Pf   | $15 ± 0.4/32 ± 0.7$                            | $107 ± 8/1.3 ± 0.1$                              |
| Ma + Mb   | $18 ± 0.9/32 ± 1$                              | $94 ± 8/1.8 ± 0.3$                               |
| Ma + Pf + Mb | $17 ± 0.7/30 ± 1$                           | $100 ± 7/1.3 ± 0.1$                              |

Values represent average ± SE

Isotope values of ostracods and sediment before the experiment

Sediment isotope values before the cyanobacterial addition were $5.0 ± 0.1$ for $\delta^{15}$N and $-22.9 ± 0.1$ for $\delta^{13}$C ($n = 3$), while initial $\delta^{15}$N and the $\delta^{13}$C values were $6.5 ± 0.5$ and $-21.9 ± 0.2$ for *C. neglecta*, $8.4 ± 0.7$, $-21.3 ± 0.3$ for *P. femorata*, and $8.2 ± 0.8$ and $-21.8 ± 0.1$ for *H. sorbyana* ($n = 3$ in all cases).

Macrofaunal survival and ostracod abundance and biomass at the beginning and end of the experiment

Survival of *M. balthica* was high in all treatments ($93 ± 2$%), while the amphipods had lower survival of $65 ± 7$% for *M. affinis* and $57 ± 5$% for *P. femorata*, with no significant difference in survival among the treatments for any of the species [*M. affinis*; $H \ (3, n = 32) = 6.547$, $P = 0.088$; *P. femorata*; $H \ (2, n = 23) = 2.594$, $P = 0.273$; *M. balthica*; $H \ (2, n = 24) = 3.290$, $P = 0.193$]. The observed amphipod mortality reflects their higher sensitivity to handling and oxygen stress and is within the range for summertime experiments of similar duration (Bianchi et al. 2000). Final macrofaunal biomass (total biomass of all surviving individuals) was similar among treatments except that the Pf treatment had slightly lower community biomass than the the Ma + Pf + Mb treatment ($F_{5,41} = 3.03$, $P = 0.007$) (see Table 1).

The abundance and biomass of ostracods before the beginning of the experiment were $94 ± 7$ ind. 50 cm$^{-2}$ and $1.25 ± 0.2$ mg dw$^{-1}$, respectively. The values for ostracod abundance and biomass at the end of the experiment are presented in Table 1. There were no significant differences among treatments in ostracod abundance at the end of the experiment or between the initial abundance of ostracods and the abundance in all the other treatments at the end of the experience (ANOVA, $F_{6,23} = 1.2$, $P = 0.4$ for *C. neglecta*; $F_{6,31} = 1.8$, $P = 0.1$ for *P. femorata*; $F_{6,31} = 0.4$, $P = 0.8$ for *H. sorbyana*). There were no significant differences in biomass among the treatments at
the end of the experiment or between initial biomass of ostracods and all the other treatments at the end of the experiment (ANOVA, $F_{1,6} = 2.3$, $P = 0.07$ for C. neglecta; $F_{1,6} = 0.8$, $P = 0.59$ for P. fennica; $F_{1,6} = 1.9$, $P = 0.2$ for H. sorbyana). Furthermore, no significant differences were found in initial N or C content for any of the ostracods (data not shown).

Ostracod isotope values after the experiment

The ostracod isotope values at the end of the experiment showed that all three species had incorporated cyanobacterial N and C, with significant differences among treatments (Fig. 1). Ostracod stable N isotope values often decreased markedly, indicating decreased incorporation of cyanobacterial N, when a macrofaunal species was present in the experimental sediments, but this depended greatly on identity of the species (nested ANOVA, composition: $F_{3,23} = 9.34$, $P = 0.0003$ for C. neglecta; $F_{3,42} = 16.7$, $P < 0.00001$ for P. fennica; $F_{3,42} = 8.6$, $P = 0.0001$ for H. sorbyana). In addition, increased macrofaunal species richness significantly decreased the nitrogen isotope values for all ostracod species, again indicating lower incorporation of cyanobacterial N ($F_{3,23} = 21.1$, $P < 0.00001$, for C. neglecta; $F_{3,42} = 23.0$, $P < 0.00001$ for P. fennica; $F_{3,42} = 27.6$, $P < 0.00001$ for H. sorbyana). Similar decreases were found for carbon isotope values, but while macrofaunal species richness significantly lowered the carbon isotope values (nested ANOVA, $F_{3,23} = 0.21$, $P = 0.0001$ for C. neglecta; $F_{3,42} = 36.4$, $P < 0.00001$ for P. fennica; $F_{3,42} = 16.3$, $P < 0.00001$ for H. sorbyana), the effect of species composition on $\delta^{13}C$ values was significant only for P. fennica (nested ANOVA, $F_{3,42} = 21.3$, $P < 0.00001$).

*Candona neglecta* had the highest N and C isotope values in the absence of macrofauna, indicating a more rapid incorporation of freshly deposited N and C (Fig. 1a), but also showed the greatest reduction in incorporation when together with macrofauna. A similar but lower decrease in incorporation of N and C of cyanobacterial origin in the presence of macrofauna was seen for *P. fennica* and *H. sorbyana* (Fig. 1b and c, respectively), which are naturally less dependent on fresh organic matter.

Ostracod community incorporation of cyanobacterial N and C

There were marked differences among treatments in the degree to which macrofauna reduced the ostracod community’s incorporation of cyanobacterial N and C (uncrossed bars in Fig. 3a for N and Fig. 3b for C). Macrofaunal species composition significantly affected the incorporation of cyanobacterial N and C by ostracod assemblages (nested ANOVA, $F_{3,34} = 33.7$, $P < 0.00001$ for N; $F_{3,34} = 11.9$, $P < 0.00001$ for C), and increased macrofauna species richness significantly reduced the amounts of labelled N and C incorporated by the ostracod community (nested ANOVA, $F_{3,34} = 96.6$, $P < 0.00001$ for N; $F_{3,34} = 71.8$, $P < 0.00001$ for C). Importantly, our results show significantly reduced incorporation of both N and C in all treatments with macrofaunal species, with observed incorporation values (crossed bars in Fig. 2) always being lower than expected in the treatments with
more than one macrofaunal species (see Fig. 2a, b for P values), and the greatest reduction found where all three macrofauna species were present.

Sediment content of cyanobacterial N and C after the experiment

There were no significant differences among treatments in content of labelled N or C in the upper centimetre of sediment (Fig. 3a, b), where ostracods and most other meiofaunal groups live (ANOVA, $F_{6,48} = 0.9$, $P = 0.49$ for N; $F_{6,48} = 0.7$, $P = 0.69$, for C). We were not able to account for all the label added to our experimental system. It is reasonable to assume that part was lost to the atmosphere as respired CO$_2$ or as NH$_4$. In addition, due to the burrows in the sediment created by the infauna, one can expect the concentration label in the sediment to be patchy and thus hard to sample difficult to subsample accurately.

Discussion

We found that the incorporation of fresh organic matter by ostracods was influenced both by the species of macrofauna, and by the number of macrofauna species present in the treatment (Fig. 1).

Regarding the effects of the macrofaunal species composition, even though $P$. femorata itself incorporated large amounts of fresh organic matter (Karlson et al. 2010) the isotope values of the three ostracod species in the treatment with $P$. femorata alone show that it did not significantly alter incorporation by any of the three ostracod species. This is probably due to the activity of $P$. femorata being mainly below the surface, away from the first centimetre of sediment where ostracods live and feed. The effects of the surface-feeding $M$. affinis tested alone were intermediate,
while *M. balthica* tested singly reduced incorporation by all three ostracod species to a greater extent than any of the amphipods. Even though this bivalve is the least mobile of the tested macrofauna species, it uses its long siphon to forage through surface sediment for phytodetritus several times a day (Ólafsson et al. 2005). The overall activity of *M. balthica*, with both the siphon and the bivalve itself moving about in the sediment, clearly result in a disturbance that interferes greatly with ostracod access to food freshly deposited on the sediment surface.

Incorporation of cyanobacterial N and C by all three ostracods was clearly reduced in treatments with macrofauna, indicating reduced feeding on freshly deposited cyanobacteria. Incorporation of label was lowest for all the three ostracod species in the Ma + Pf + Mb treatment (Fig. 1), even though there was no significant difference in the surface sediment content of cyanobacterial N and C among treatments at the end of the experiment (Fig. 3). Furthermore, there was no significant difference between the treatments with macrofauna in the amounts of cyanobacterial N and C buried beneath the top centimeter of the sediment. This indicates that the lower incorporation of cyanobacterial N and C by meiofauna was not due to burial of phytodetritus by macrofauna. In addition, the low incorporation values for ostracods in the *M. balthica* monoculture, the macrofauna species with lowest incorporation values, also indicate that interference rather than exploitative competition by macrofauna for freshly deposited detritus was the main mechanism behind the differences among the treatments in incorporation of N and C by the ostracods. This is further supported by the weak correlations between incorporation of freshly deposited N and C by ostracods and macrofauna (data from Karlson et al. 2010) \( r^2 = 0.02, P = 0.4 \) for N; \( r^2 = 0.009, P = 0.89 \) for C and between meiofauna incorporation and surface sediment content of cyanobacterial N and C \( r^2 = 0.03, P = 0.3 \) for N; \( r^2 = 0.06, P = 0.6 \) for C).

Our results show that the reduction in incorporation of labeled N and C increased with macrofaunal species richness. In treatments with more than one macrofauna species, the reduction in incorporation of cyanobacterial N and C by ostracods was consistently significantly greater (Fig. 2) than expected from the effects of macrofauna species tested singly. This increased interference with higher macrofaunal species richness may be a result of the different feeding and bioturbation activities of the three macrofaunal species hindering ostracod feeding on freshly deposited N and C in complementary ways. Several studies have found spatial niche segregation between the two studied amphipods when in sympatry (Byrén et al. 2006; Hill and Elmgren 1987). This increased macrofaunal spatial segregation with higher diversity probably reduces the sediment volume in which meiofauna can avoid the disturbance caused by macrofaunal activity. That the highest interference was found in the Ma + Pf + Mb treatment was possibly due to the sediment reworking activities of *M. balthica* increasing the physical disruption of the sediment caused by the bioturbation of the mobile amphipod *M. affinis*. A way for ostracods to avoid this surface disturbance would be to move deeper in the sediment. Indeed, Modig et al. (2000) suggested that fine scale stratification within the first 2 cm of sediment is possible in ostracods. However, the presence of the subsurface feeder *P. femorata* is likely to reduce the success of this strategy, by increasing disturbance and sediment disruption in deeper sediment layers. Thus, it seems likely that ostracods are forced to reduce their feeding activity on the added cyanobacteria when faced with high frequency amphipod and bivalve disturbance both at and below the surface. It is also possible that ostracods change their feeding strategy to consumer other unlabeled food sources when faced with high macrofauna interference. This would require ostracod species to actively avoid fresh organic matter from settling phytoplankton blooms while they forage in the sediments, which seems less plausible. However, our experimental design cannot distinguish these possibilities.

The three ostracod species were differently affected by the increased interference caused by macrofauna, with *C. neglecta* showing the greatest reduction in isotope values when compared to the treatment without macrofauna (Fig. 1). Other studies have found *C. neglecta* to incorporate freshly deposited organic matter from phytoplankton blooms at much higher rates than the other two ostracods (Nascimento et al. 2008; Ólafsson et al. 1999). Modig et al. (2000) indicated that the different feeding ecologies of these three ostracod species might be connected to their life-history characteristics. In the Baltic Sea, *P. fennica* and *H. sorbyana* have a 2-year life cycle (Ankar and Elmgren 1976), while *C. neglecta* is able to reach the adult stage in 4 months (Savolainen and Valtonen 1983). The shorter generation time of *C. neglecta* probably requires it to feed and grow more quickly, and to use pulses of higher quality organic matter with greater efficiency. Nevertheless, in the multi-macrofaunal species treatments, this difference in assimilation of freshly deposited organic matter between *C. neglecta, H. sorbyana* and *P. fennica* was reduced, as incorporation of fresh bloom material by *C. neglecta* suffered the greatest decrease (Fig. 1). The more time an ostracod spends with its shells closed to avoid contact with macrofauna, the less time it will have available for feeding. It is, therefore, not unexpected that *C. neglecta* showed a greater reduction in incorporation. Our results suggest that outcomes of the effects of interference by larger competitors will depend on the life history traits and characteristics of the impacted species.
Our results have implications for understanding mechanisms that underlie species coexistence in aquatic sediments. Few papers have studied both macrofaunal diversity and meiofauna abundance and distribution in systems below the photic zone with a taxonomic resolution that allows testing of meaningful hypotheses about their relationship. In one of these studies, Ankar and Elmgren (1976) present data on both macrofaunal and meiofaunal distributions at 35 stations sampled in a stratified random sampling design in the geographical area where we performed our experiment. Analyzing their data on the abundance and species composition of benthic communities below the photic zone (≥15 m) we found that both abundance and biomass of meiofauna are negatively correlated with macrofaunal biodiversity, measured as Shannon index (Fig. 4; \( r^2 = 0.27, P = 0.0018 \) for abundance; \( r^2 = 0.12, P = 0.034 \) for biomass). Even though data from other geographical regions are needed, the negative correlation between macrofaunal diversity and meiofaunal abundance, and the increase in interference with higher species richness found in our study, indicate that interference competition with meiofauna by macrofauna is increased at greater macrofauna diversity.

As benthic communities living below the photic zone are critically dependent on pulsed inputs of settling organic matter for food (Graf 1992), most benthic deposit-feeders, like C. neglecta and to a lesser degree P. fennica and H. sorbyana, are sufficiently opportunistic in their diet to use an infrequently available resource. The ability of a consumer to increase consumption of a normally limiting resource when it becomes available in abundance has been shown to drive complex patterns of community dynamics in a number of ecological systems (Ostfeld and Keesing 2000; Yang et al. 2008). Our data show that an increase in species richness can intensify interference competition, and reduce the access of an impacted species to a limiting resource, potentially playing an important role in community dynamics. No effects of macrofaunal diversity on the abundance or biomass of ostracods were observed in our short study. However, the effect on the incorporation of freshly deposited cyanobacteria was clear. As incorporation of organic matter is the first step in somatic growth, it is a promising indicator in studies of factors controlling the secondary production of slow-growing species (Karlson et al. 2010).

Interference competition can be a decisive factor in community dynamics (Amarasekare 2002, 2003). Indeed, one of the most striking patterns in benthic communities is the decrease in abundance and biomass of benthic species with water depth, which is more rapid for macrofauna than for meiofauna (Rex et al. 2006; Thiel 1975). This transition to a meiofaunal dominated system in low energy habitats is also seen in the Baltic Sea with meiofaunal biomass exceeding that of macrofauna in the Bothnian Sea (Elmgren 1978). This transition is connected to lower food availability in this ecosystems, which do not provide enough energy to sustain high standing stocks of large organisms (Rex et al. 2006; Thiel 1975). As meiofauna have higher feeding efficiency than macrofauna (Elmgren 1978; Thiel 1975), their biomass and abundance decreases at a slower rate with depth. Our results suggest that another mechanism contributes to this global pattern of an increased proportion of smaller organisms with depth. We show that macrofauna interferes with the access of meiofauna to settled organic matter. With lower abundance and diversity of macrofauna in low energy systems, meiofauna may be released from interference competition and have better access to the food available, which may to some extent counterbalance the effect of lower food availability, thus contributing to the slower rate of decrease in meiofauna numbers and biomass and to the consequent decreasing size of metazoans with depth.

In conclusion, our study shows that interference competition from macrofauna can reduce incorporation of
freshly deposited phytoplankton material by meiofaunal ostracods, and that this interference is aggravated when macrofauna species richness is increased, even at constant macrofauna biomass.

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