

1 **TITLE: Diversity and spatial variability of shallow benthic macrofaunal assemblages in**  
2 **a high-Arctic fjord (Young Sound, North-East Greenland)**

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27 **Abstract**

28 Although knowledge of Arctic benthic biodiversity has increased considerably in recent  
29 decades, some regions, such as Northeast Greenland, remain poorly studied. The aim of this  
30 study was to complement a previous macrofaunal inventory carried out in Young Sound, a  
31 High-Arctic fjord in this region (74°N). We sampled shallow benthic assemblages along a small  
32 inner/outer fjord gradient, including one station previously prospected two decades ago and  
33 three new stations. This sampling strategy revealed highly diversified benthic assemblages (166  
34 species identified on a total sampling area of 1.32 m<sup>2</sup>), which considerably increases the number  
35 of species recorded for the fjord (i.e. 225 species vs 100 previously recorded). The outermost  
36 station was dominated in abundance by various assemblages of bivalves, while the middle  
37 stations showed greater species evenness, including numerous species of polychaetes, bivalves  
38 and crustaceans. The innermost station was dominated by ostracods, gammarid amphipods and  
39 tube-dwelling polychaetes. Overall, benthic assemblages varied little between the four stations  
40 and the transect as a whole exhibited characteristics typical of outer fjord habitats, reflecting  
41 the rather moderate impact of meltwater inputs in this part of the fjord. Finally, trophic plasticity  
42 and omnivory were observed in most of the recorded macrobenthic species, highlighting the  
43 adaptability of these species to low trophic availability in the ecosystem. Future biodiversity  
44 studies will need to explore the innermost and deepest areas to provide a more comprehensive  
45 inventory and understanding of the influence of environmental conditions on the structure and  
46 functioning of Young Sound benthic habitats.

47 **Key words**

48 **Benthic ecosystems • Macrofauna • Community structure • Spatial variability • Meltwater**  
49 **inputs • Arctic fjord**

## 50 **Introduction**

51 To date, Arctic benthic biodiversity remains poorly studied although species colonizing the  
52 seafloor represent around 90 % of the total estimated species richness of Arctic invertebrates  
53 (Sirenko 2001; Gradinger et al. 2010). While Arctic benthic ecosystems have long been  
54 considered as areas of low species diversity (e.g. Knox and Lowry 1977), there is increasing  
55 evidence that this assumption has likely been biased by the scarcity of surveys conducted at  
56 high latitudes (Piepenburg 2005; Piepenburg et al. 2011). Recent inventories estimate the  
57 diversity of Arctic benthic species to be over 4600 species, a much higher number than  
58 previously estimated, while several thousand species are likely to remain to be discovered  
59 (Bluhm et al. 2011; Josefson et al. 2013). Although considerable progress has been made in  
60 Arctic biodiversity in recent decades, some geographic areas remain understudied such as East  
61 Greenland (Piepenburg et al. 2011).

62 Under climate change, Arctic benthic ecosystems are expected to undergo major modifications  
63 driven by changes in sea ice cover, salinity and temperature, acidification, or sedimentation  
64 (AMAP 2017). It is very likely that these changes will impact benthic biodiversity, especially  
65 in coastal areas, which are expected to be the most exposed to future environmental changes  
66 (Węśławski et al. 2011). The expansion of boreal species into high latitudes is also expected to  
67 affect benthic biodiversity by bringing new species into Arctic areas (Węśławski et al. 2011;  
68 Kotwicki et al. 2021; Cottier-Cook et al. 2023), with likely consequences for species  
69 interactions and ecosystem functioning. Indeed, the recent expansion of several engineer  
70 species with boreal affinities into the Arctic (e.g. mussel species from the *Mytilus* spp. complex  
71 or the kelp *Laminaria hyperborea*) has had implications for the structure of shallow benthic  
72 ecosystems by increasing habitat heterogeneity and complexity, facilitating the arrival of  
73 species new to these ecosystems (Berge et al. 2005; Krause-Jensen and Duarte 2014; Mathiesen  
74 et al. 2017; Filbee-Dexter et al. 2019; Kotwicki et al. 2021). Many questions remain unresolved  
75 regarding how endemic benthic species will cope with new abiotic conditions (increases in  
76 temperature or sedimentation, decreases in seawater salinity) and associated changes on biotic  
77 interactions (e.g. the arrival of new predators or competitors, Renaud et al. 2015). Assessing  
78 the vulnerability of Arctic benthic communities experiencing such environmental and  
79 ecological changes thus remains challenging and highlights the strong need for initial references  
80 states in benthic biodiversity in all Arctic ecoregions.

81 In this context, any increase in knowledge of the benthic biodiversity of poorly characterized  
82 Arctic ecoregions is crucial, especially in shallow coastal ecosystems. For example, the only  
83 studies conducted on the North-Eastern coasts of Greenland have focused on epifauna and/or  
84 megafauna shelf communities (e.g. Piepenburg and Schmid 1996; Brandt and Schnack 1999;  
85 Starmans et al. 1999; Fredriksen et al. 2020). To our knowledge, for such real *Terra Incognita*  
86 of Arctic ecoregions (Piepenburg et al. 2011), only Sejr et al. (2000) provided quantitative  
87 datasets on shallow macrozoobenthos assemblages in a high-Arctic fjord (Young Sound, NE  
88 Greenland). Few studies have also investigated trophic linkages in macrobenthic ecosystems  
89 from the Northeast Water Polynya providing only limited details on their biodiversity (Hobson  
90 et al. 1995; Ambrose and Renaud 1997; Piepenburg et al. 1997). Finally, although many older  
91 studies from the mid-20<sup>th</sup> century have been conducted on the shallow habitats (e.g. Ockelmann  
92 1958), they have mostly focused on the occurrence of individual taxa than providing insights  
93 into the structure and diversity of macrobenthic communities.

94 The aim of the present study is to complement the inventory of benthic macrofaunal species  
95 carried out by Sejr et al. (2000) in Northeast Greenland (Young Sound, 74°N) with additional  
96 sampling sites. While Sejr et al. (2000) examined the macrobenthic communities in vertical  
97 zonation (from 20 to 85 m), this study prospects shallow benthic macrofaunal assemblages ( $\approx$   
98 20 m) distributed along a small inner/outer gradient located in the most marine part of the fjord.  
99 We resampled the shallowest Daneborg station from Sejr et al. (2000) and investigated three  
100 new stations, one towards the outer fjord (Kap Breusing) and two towards the inner fjord (Basalt  
101 Island, Pass Hytten). The description of the assemblages, coupled with those of Sejr et al.  
102 (2000), aims to provide a baseline for future studies investigating the temporal variability of  
103 Arctic shallow coastal soft-bottom communities and their response to ongoing climate change.

## 104 **Materials and Methods**

### 105 **Study sites**

106 Young Sound is a high Arctic fjord located in North-East Greenland (Fig. 1) of about 90 km  
107 long and 2-7 km wide (Rysgaard et al. 2003). A sill is located at its outer section ( $\approx$  45 m depth,  
108 see Fig. 1) limiting seawater exchanges between the inner parts of the fjord and the shelf  
109 (Bendtsen et al. 2007) where average and maximum depths reach 100 and 360 m, respectively  
110 (Rysgaard et al. 2003). This fjord is characterized by extreme seasonal variations, thick sea ice  
111 (> 150 cm) and snow cover (20 – 100 cm) covering the fjord for 9 to 10 months per year (Glud

112 et al. 2007) and strongly limiting the annual primary production, estimated to be about 10 g C  
113 m<sup>-2</sup> yr<sup>-1</sup> (Rysgaard et al. 1999). During the summer, glacier and snow melt generate strong river  
114 flows ranging annually from 0.9 to 1.4 km<sup>3</sup> (Bendtsen et al. 2014). Such freshwater inputs  
115 induce strong spatial salinity and temperature gradients in summer surface waters, with  
116 temperature decreasing from 9 to 2°C and salinity increasing from 8 to 30, from the inner to the  
117 outer part of the fjord, respectively (Bendtsen et al. 2007). In contrast, deeper waters (> 20 m)  
118 generally remain stable throughout the year (i.e. > 2° C) along the entire fjord length (Holding  
119 et al. 2019). Vertical variations in salinity induce a strong permanent haline stratification at 5  
120 to 10 m depth, which limits pelagic primary production by reducing nutrient exchange between  
121 surface and subsurface waters (Holding et al. 2019; Sejr et al. 2022). On a decadal scale, the  
122 sea ice conditions in the fjord have changed with a gradual increase in the duration of the ice-  
123 free season from about 80 days per year (1950-2000) to more than 90 days per year but  
124 combined with a significant increase in the year-to-year variability since 2000. At the same  
125 time, freshwater runoff (mostly glacial meltwater) is estimated to have increased 6-7 fold since  
126 the 1960s (Sejr et al. 2022). Annual monitoring of main Young Sound environmental  
127 parameters (physico-chemical, pelagic productivity and community structure) has been made  
128 for three decades by the Greenland Ecosystem Monitoring – Marine Basis program, making  
129 Young Sound ideal for studying the effects and feedback of climate change in the high Arctic.  
130 However, the macrozoobenthic benthic communities are not part of the monitoring program  
131 since the study by Sejr et al. (2000), highlighting the crucial need to reacquire biodiversity data.

### 132 **Sampling and laboratory analyses**

133 Sampling was conducted in August 2016 at four stations distributed along an inner/outer marine  
134 fjord gradient: Pass Hytten, Basalt Island, Daneborg and Kap Breusing (Fig. 1, Table 1).  
135 Macrofauna samples were collected by scuba divers for each replicate using a modified Hiscock  
136 and Hoare suction dredge (Eleftheriou and McIntyre 2005), fitted with a suction pipe directly  
137 connected to a nylon mesh sample bag (square mesh = 1 mm). Each replicate was collected  
138 over an area of 0.09 m<sup>2</sup> (except for one replicate at Kap Breusing, which was collected over  
139 0.0625 m<sup>2</sup>) and then fixed directly in 10% buffered formalin. Sediment samples were also  
140 collected at both Pass Hytten and Kap Breusing using small sediment cores for grain size  
141 analysis. Due to logistical problems, we were unable to collect sediment samples at either  
142 Daneborg or Basalt Island. Once in the laboratory, the animals were sorted, transferred to 70%  
143 ethanol and then identified to the lowest taxonomic level. Sediment samples were analyzed by  
144 the private company “Eurofins – Analyses pour l’environnement” (Saverne, France) using laser

145 methods: samples were sieved through a 2 mm square mesh and then analyzed using a Malvern  
146 Mastersizer 2000 laser particle size analyzer to provide different size fractions ranging from 2  
147 to 2000  $\mu\text{m}$ .

#### 148 **Statistical analyses**

149 Spatial differences in benthic assemblage structure were visualized using Principal Component  
150 Analysis (PCA) performed on Hellinger-transformed fauna abundances. This transformation  
151 was selected to reduce the weight of rare species in the analysis (Legendre and Gallagher 2001).  
152 The first twenty species explaining the most spatial variation in the ordination (i.e. accounting  
153 for more than 60% of SIMPER dissimilarities) were plotted in a second graph to visualize which  
154 species contributed to the benthic structure of each station. Spatial differences in community  
155 structure were tested using one-way permutational multivariate analysis of variance  
156 (PERMANOVA) performed on Bray-Curtis transformed fauna abundances. Homoscedasticity  
157 was checked using a permutational analysis of multivariate dispersions (PERMDISP) test to  
158 ensure that data dispersion did not influence the PERMANOVA result (Anderson et al. 2008;  
159 Anderson and Walsh 2013). Species rarefaction curves were calculated for each replicate to  
160 compare trends in species richness between stations. Species richness was compared among  
161 replicates based on rarefaction curves by standardizing all data to the same abundance  
162 (corresponding to the number of individuals in the replicate with the lowest abundance) to  
163 remove any abundance bias on taxa richness (Gotelli and Colwell 2001). An additional  
164 rarefaction curve was plotted to estimate the specific richness for the entire Young Sound  
165 system and Hulbert's diversity index was calculated to estimate the expected number of species  
166 within a random sample of 35 individuals (based on the lowest number of individuals observed  
167 in a replicate). Jackknife1 Chao2 estimates were calculated and plotted as accumulation curves  
168 to estimate the total species richness in case species rarefaction curves did not reach their  
169 asymptotes (Colwell et al. 2004; Chao and Chiu 2016). Finally, we calculated the Pielou's  
170 evenness index to take into account the diversity observed at each station along with the  
171 associated species richness using the "vegan" R package (Oksanen et al. 2019). All other  
172 diversity indices were calculated using EstimateS v9.1 software.

## 173 Results

### 174 Structure of benthic community

175 Significant differences in benthic community structure were observed between sampling  
176 stations (degrees of freedom for treatment and error [ $df$ ] = 3 and 11, pseudo- $F$  = 8.2529,  $p$ -perm  
177 = 0.001). Pairwise analyses revealed that such differences were observed between all stations  
178 (the smallest significant difference was observed between Basalt Island and Kap Breusing,  $df$   
179 = 1 and 4, pseudo- $F$  = 7.4856,  $p$ -perm = 0.049). PCA analysis showed that more than half of  
180 the spatial variability was explained by the first three principal components (PC1, PC2 and PC3  
181 explained 24.36, 17.92 and 17.27 % of the variance, respectively). Low dispersion was  
182 observed among the replicates of each station (Fig. 2a, c). Significant differences in Pielou's  
183 evenness index were observed between stations ( $df$  = 3 and 14, pseudo- $F$  = 36.4600,  $p$ -perm =  
184 0.001). These differences were observed between all stations by pairwise PERMANOVA  
185 analyses (the smallest significant difference was observed between Daneborg and Kap  
186 Breusing,  $df$  = 1 and 4, pseudo- $F$  = 16.2496,  $p$ -perm = 0.048), except between Pass Hytten  
187 and Kap Breusing ( $df$  = 1 and 4, pseudo- $F$  = 0.0386,  $p$ -perm = 1).

188 Benthic assemblages at Pass Hytten were strongly dominated by crustaceans (relative  
189 abundance = 52.9 - 78.0%, Fig. 3a, b), mainly due to high densities of ostracods and the  
190 amphipod *Metopa glacialis* (Fig. 2b). The community structure at Daneborg and Basalt Island  
191 was more evenly balanced between taxa (Fig. 3a, b), although polychaetes were the most  
192 dominant group (relative abundance = 32.6 - 58.3% at Daneborg and 44.9 - 55.9% at Basalt  
193 Island). *Euchone analis*, *E. incolor* and *Laphania boeckii* were the most abundant polychaete  
194 species at Daneborg, while *Abyssoninoe* sp., *Laphania boeckii* and *Nereimyra aphroditoides*  
195 dominated the polychaete assemblage at Basalt Island (Fig. 2b, d). Finally, bivalves were the  
196 dominant taxonomic group at Kap Breusing (relative abundance = 67.5 - 68.5%, Fig. 3a, b),  
197 largely represented by high densities of the filter-feeding bivalves *Hiatella arctica* and  
198 *Musculus discors* (> 50%, Fig. 2b, d).

199 Total abundance showed large variation among stations and replicates (Fig. 3b). The highest  
200 abundances were observed in the second replicate from Kap Breusing (reaching 6044 ind. m<sup>-2</sup>,  
201 Fig. 2b). In contrast, the first replicate from Kap Breusing showed a much lower total abundance  
202 (1824 ind. m<sup>-2</sup>), which was in the same order of magnitude as the abundances observed at  
203 Daneborg (from 933 to 1733 ind. m<sup>-2</sup>) and Pass Hytten (from 1467 to 2667 ind. m<sup>-2</sup>, Fig. 2b).

204 Finally, the lowest total abundances were observed at Basalt Island (from 389 to 1611 ind. m<sup>-2</sup>,  
205 Fig. 2b).

## 206 **Abiotic and biotic description of benthic habitats**

### 207 *Pass Hytten*

208 The sediment texture is dominated by fine silt particles (55% of the total size fraction for  
209 particles < 63 µm) mixed with fine sands (25%) and gravels (20%, Fig. 1 in Online Resource  
210 1). Significant bioturbation activity probably occurs due to high abundances of epifauna (e.g.  
211 *Ophiocten sericeum*, Fig. 4a, Table 2) and burrowing shrimps (e.g. *Sclerocrangon boreas*, L.  
212 Chauvaud pers. obs.). The associated macrozoobenthic assemblage shows a low species  
213 evenness with ten taxa representing 83.5% of the total abundance (Table 2), including  
214 crustaceans such as ostracods, the amphipod *Metopa glacialis* and the cumacean *Diastylis*  
215 *scorpioides* (59 % of the total abundance). The strictly benthic species (epifauna and endofauna)  
216 were dominated by both tube-dwelling (*Euchone incolor* [see high abundances in Fig. 3a],  
217 *Lyssipe labiata*, *Maldane sarsi* and *Owenia borealis*) and errant polychaetes (*Abyssoninoe* sp.  
218 and *Leitoscoloplos mammosus*). Bivalves *Astarte moerchi*, *Hiatella arctica*, *Mya truncata* and  
219 *Musculus discors*) also occur but in much lower proportions (< 15 % of the total abundance).

### 220 *Basalt Island*

221 Here, the sediment is a mixture of stone, sand and small gravel (2-5 mm) associated with detrital  
222 organic matter (mainly macroalgae and shell debris) accumulated on the seafloor (Fig. 4b).  
223 However, we were unable to sample the sediment at this station due to the dominance of stones.  
224 The steep slope of the seafloor (slope gradient ≈ 80 – 100 %, see Fig. 4b) probably makes the  
225 bottom unstable due to the downward export of mineral material towards the deep fjord basins.  
226 The species assemblage at Basalt Island shows greater species evenness than Pass Hytten and  
227 Kap Breusing, with the ten most dominant species accounting for only 51.3% of total abundance  
228 (Table 2). Polychaetes dominate the species assemblage with *Abyssoninoe* sp., *Clymenura* sp.,  
229 *Laphania boeckii*, *Nereimyra aphroditoides* and *Pholoe* sp. representing 26.4 % of the total  
230 abundance. Several bivalves such as *A. moerchi*, *H. arctica* or *M. truncata* were also found at  
231 the site but in low abundances, as well as the brittle star *Ophiocten sericeum* (Table 2, Table S1  
232 in Online Resource 1). Among crustaceans, ostracods show a high abundance (5.9 %, Table 2).

### 233 *Daneborg*

234 Based on underwater pictures and previous data on particle size distribution (Sejr et al. 2000),  
235 the seabed was characterized by a mixture of fine and very fine sand (particles [63 - 125 µm] =



236 24.6 % and particles > 125  $\mu\text{m}$  = 44.2 %). The species evenness of the species assemblage was  
237 close to that observed at Basalt Island (Table 2). Daneborg is mainly dominated by two taxa,  
238 bivalves and in particular the species *Astarte moerchi* (18.6 %) and crustaceans (ostracods and  
239 *D. scorpioides* representing 11.8 % and 3.1 % of the total abundance, respectively). Polychaetes  
240 complete the assemblage, including *Euchone incolor*, *Laphania boeckii*, *Euchone analis*, *Chone*  
241 sp. and *Clymenura* sp., which all account for 23.6% of the total abundance.

#### 242 *Kap Breusing*

243 The grain size distribution of Kap Breusing was variable between replicates, but overall it was  
244 a poorly sorted sandy sediment (the 63-2000  $\mu\text{m}$  grain size range varied from 53 to 80 %, Fig.  
245 S1 in Online Resource 1). Heterogeneous medium/coarse sand (38 to 55 % of the particles  
246 between 250 and 2000  $\mu\text{m}$ ) was found in two of the three replicates, the other was largely  
247 composed of fine sand (61 % of particles between 63 and 250  $\mu\text{m}$ ).

248 The species assemblage here was associated with a rock wall close to the location where the  
249 sediment samples were collected and included Sertulariidae hydrozoans and *Musculus discors*  
250 bivalves (L. Chauvaud, pers. obs., see background of Fig. 4d). Both species form a complex  
251 three-dimensional structure hosting a highly diversified assemblage including the gastropod  
252 *Alvania scrobiculata* and the bivalves *A. moerchi*, *H. arctica*, *M. discors* and *M. truncata*,  
253 representing more than 68% of the total abundance (Table 2). Four vagile polychaete species  
254 were also found as *Harmothoe* sp., *Nereimyra aphroditoides*, *Nereis zonata* and *Syllis* sp. (11.3  
255 % of total abundance) and the sedentary terebellid *Polycirrus medusa* (2.8 %; Table 2).

#### 256 **Biodiversity estimates of benthic habitats**

257 Based on a total area of 1.32 m<sup>2</sup> sampled by 15 suction dredge replicates, we collected a total  
258 of 2322 individuals belonging to 166 species (Table S1 in Online Resource 1). The polychaete  
259 phylum was by far the most diverse (67 species), followed by mollusks (33 species) and  
260 crustaceans (31 species). Echinoderms, nemerteans, sipunculans, pycnogonids, tunicates,  
261 nematodes, brachiopods and platyhelminths were also found but always displayed with much  
262 lower species richness. Macroalgae randomly collected in our samples were also identified, and  
263 the species list is included in the supplementary material as complementary observations (Table  
264 S2 in Online Resource 1).

265 Species accumulation curves did not reach their asymptotes for all stations (Fig 5). The initial  
266 slope of the species accumulation curves was highly variable between stations, but more  
267 consistent between intra-site replicates. Based on the lowest number of individuals (i.e. 35 in  
268 the 5<sup>th</sup> Basalt Island replicate), Hulbert's expected species richness within a random sample of

269 35 individuals ( $ES_{35}$ ) displayed the lowest levels in Pass Hytten and Kap Breusing (i.e. range  
270  $ES_{35} = 9 - 13$  and  $12 - 13$  for Pass Hytten and Kap Breusing, respectively). Expected species  
271 richness showed intermediate values in Daneborg (range  $ES_{35} = 15 - 21$ ) and the highest levels  
272 in Basalt Island (range  $ES_{35} = 19 - 22$ ). Based on the total number of species per replicate ( $S_{tot}$ ),  
273 species richness was also lower at Pass Hytten (range  $S_{tot} = 17 - 31$ ) than at Basalt Island (range  
274  $S_{tot} = 19 - 43$ ) and Daneborg (range  $S_{tot} = 32 - 47$ ), while Kap Breusing had the highest intra-  
275 station variability (range  $S_{tot} = 27 - 59$ ). A rarefaction curve based on the total set of fauna  
276 samples (i.e. all replicates from the four sampled stations combined) also does not reach an  
277 asymptote either, despite the much higher number of individuals (i.e.  $> 2000$ ).

278 The rarefaction curves of Chao2 and Jackknife1 estimates slightly differ from the species  
279 accumulation curves (Fig. 6a, b). Based on the lowest number of replicates (i.e. 2 sampled  
280 stations at Kap Breusing), biodiversity estimates were lowest at Pass Hytten (Chao2 = 48,  
281 Jackknife1 = 51), intermediate at Basalt Island (Chao2 = 62, Jackknife1 = 64) and Daneborg  
282 (Chao2 = 77, Jackknife1 = 78), and highest at Kap Breusing (Chao2 = 95, Jackknife1 = 91).  
283 Based on the entire dataset, Chao2 and Jackknife1 estimates predicted a total species richness  
284 of 204 and 221 species, respectively.

## 285 **Discussion**

### 286 **Benthic species richness of the study area: historical and recent data comparison**

287 The present study complements the previous work of Sejr et al. (2000) conducted twenty years  
288 ago in Young Sound, which described gradual, spatially limited changes in species assemblages  
289 along a depth gradient (from 20 to 85 m) in the vicinity of Daneborg station (Fig. 1). Our study  
290 conducted along a small inner/outer fjord gradient in the most marine part of Young Sound  
291 shows little spatial variation in shallow (above 20 m depth) benthic community structure. This  
292 pattern suggests that our four sampling stations cover a relatively continuous and uniform  
293 habitat. The sampled habitat is mostly dominated by large tubicolous polychaetes (e.g.  
294 Maldanidae, *Euchone*) and some specific motile polychaetes (*Leitoscoloplos*, *Lumbrineris*),  
295 and has a high abundance of suspension-feeding bivalves (e.g. *Astarte*, *Bathyarca*, *Hiatella*).  
296 Turbidity and meltwater inputs are generally factors that control community structure in Arctic  
297 fjords, decreasing the presence of filter feeders towards the inner areas of fjords exposed to  
298 glacial melt (Włodarska-Kowalczyk and Pearson 2004; Sejr et al. 2010; Włodarska-Kowalczyk  
299 et al. 2012; Udalov et al. 2021). However, we doubt that turbidity and meltwater inputs account  
300 for the spatial variations observed between our four stations, as weak environmental forcing is  
301 observed along the small inner/outer fjord gradient studied in our survey. In fact, although

302 turbidity reaches more than 50 NTU in the innermost parts of Young Sound, it only varies from  
303 2-3 to less than 1 NTU between Pass Hytten and Kap Breusing (Meire et al. 2017). It is therefore  
304 more likely that the observed spatial variability in community structure reflects small-scale  
305 habitat variation (bottom varying abundance in bare rock, fine sand, mixed sand and gravel or  
306 mud) rather than large-scale environmental variation (i.e. differences in exposure to meltwater  
307 inputs).

308 The present study provides a complementary census of 166 benthic species, including  
309 numerically dominant polychaetes (67 species), crustaceans (46 species) and mollusks (33  
310 species). It complements the previous work by Sejr et al. (2000) which listed a total of 100  
311 species despite a higher sampling effort (total sampled surface = 2.24 m<sup>2</sup> vs 1.3225 m<sup>2</sup> here).  
312 With only 41 species in common between both studies, the total number of benthic invertebrate  
313 species recorded from the Young Sound now reaches a total of 225. The relatively low number  
314 of taxa shared between the two inventories mainly reflects the different sampling strategies as  
315 Sejr et al. (2000) targeted species assemblages from increasing depths while our sampling  
316 focused on a limited spatial fjord gradient that covers contrasting shallow habitats (i.e. silt,  
317 sand/gravel, rocky wall) and their associated species assemblages.

318 It should also be noted that identification bias cannot be excluded as a single species may be  
319 recorded under two different names over time, as different identification experts were involved  
320 and taxonomical descriptions have been produced since 2000. As an example, *Scoloplos*  
321 *armiger* (only in Sejr et al. 2000) and *Leitoscolopos mammosus* (only here) have been reported  
322 to taxonomic confusion (Mackie 1987; Renaud et al. 2007). Further census efforts will include  
323 harmonization of the species list with the involvement of benthic taxonomy experts. It could be  
324 assumed that the relatively low number of species common to this inventory and the study by  
325 Sejr et al. (2000) is due to long-term variability in benthic community structure. However, this  
326 seems unlikely as previous studies conducted in Arctic silled-fjord show a high stability of  
327 benthic community structure over several decades (Renaud et al. 2007), in contrast to open  
328 fjords (Beuchel et al. 2006; Kortsch et al. 2012).

329 As indicated by the slopes of the species accumulation curves, the inventory of benthic  
330 biodiversity in this fjord is far from being complete, with many rare species likely yet to be  
331 recorded at all study sites. The expected number of species per 35 individuals (ES<sub>35</sub>) varies  
332 between 9 and 22 species, underlining the high biodiversity but also the heterogeneity across  
333 stations and replicates. Based on biodiversity estimates, total species richness of the outer fjord

334 shallow sites should reach around 204-221 species, much higher than previous estimates based  
335 on Sejr et al. (2000) and this should be much higher as only a very limited part of the shallow  
336 Young Sound has been visited and a wide variety of habitats remains unexplored. Inventory  
337 studies on shallow and deep areas of both outer and inner fjord usually find much higher species  
338 diversity than in the present work. For example, Kędra et al. (2010) recorded more than 300  
339 taxa after extensive multi-year sampling of all soft-bottom habitats in Hornsund (SW Svalbard,  
340 from 30 m to 250 m), and Voronkov et al. (2013) found 403 benthic taxa in shallow hard-bottom  
341 communities from Kongsfjorden (NW Svalbard). We thus think that the actual benthic species  
342 inventory of the Young Sound is far from complete, as it is limited to the only most marine  
343 section.

#### 344 **Ecological characteristics and insights about the functioning of the benthic fauna**

##### 345 *Ecological characteristics of benthic assemblages*

346 The current inventory identifies four major benthic assemblages, all belonging to the Young  
347 Sound outer fjord benthic community: 1) the rocky outer fjord assemblage (Kap Breusing), 2)  
348 the sandy outer fjord assemblage (Daneborg), 3) the sand-gravelly outer fjord assemblage  
349 (Basalt Island) and 4) the silty outer fjord assemblage (Pass Hytten).

350 The macrobenthic assemblage collected at Kap Breusing was composed of a unique and poorly  
351 described species assemblage dominated by two engineer species, the bivalve *Musculus discors*  
352 and the sertulariid colonial hydroids. Several studies have highlighted the ability of some  
353 bivalve species to secrete byssus threads to construct nests that host a high diversity of other  
354 benthic invertebrate species (e.g. *Arcuatula senhousia*, *Limaria hians*; Morton 1974; Hall-  
355 Spencer and Moore 2000). Byssus threads produced by these species transform the physical  
356 habitat into a complex three-dimensional structure that is then colonized by hydroids (Merrill  
357 and Turner 1963). To our knowledge, only one study has reported this ability for *M. discors*  
358 (Merrill and Turner 1963). The high biodiversity observed at Kap Breusing (where the highest  
359 number of species was observed within a single replicate) reflects the structuring effect of the  
360 *M. discors* / Sertulariidae assemblage on the community structure. The high abundance and  
361 diversity of motile polychaetes in this assemblage, especially for Nereididae, Polynoidae or  
362 Syllidae, is similar to assemblages observed in complex three-dimensional habitats such as  
363 maerl beds, kelp forests or artificial structures (Grall et al. 2006; Yakovis et al. 2007; Pabis and  
364 Sicinski 2010). The *M. discors* / Sertulariidae assemblage was also dominated by large  
365 suspension-feeding bivalves (*Hiatella arctica*, *Musculus discors* and *Mya truncata*), which is

366 more in line with common species assemblages observed on seafloors protected from sediment  
367 disturbance caused either by freshwater inputs or by burrowing species (Włodarska-Kowalczyk  
368 2007).

369 The outer fjord sandy assemblage of Daneborg was dominated by the bivalve *Astarte moerchi*,  
370 ostracods and polychaetes of the genus *Euchone*. The dominance of *A. moerchi* among the  
371 bivalves collected at Daneborg is similar to that observed by Sejr et al. (2000) at the same  
372 station but community structure differs with our data: several abundant polychaete species  
373 found in 2016 (e.g. *Clymenura* sp., *Euchone* spp., *Laphania boeckii*) were not observed by Sejr  
374 et al. (2000) while other species abundant in 1995 (e.g. *Dipolydora quadrilobate* and  
375 *Lumbriclymene minor*) were not collected in 2016. Such long-term differences could be  
376 explained either by long-term changes in the benthic assemblages but also to our weak sampling  
377 effort that is submitted to small scale habitat heterogeneity.

378 The Basalt Island sand-gravel outer fjord assemblage is dominated by an association of motile  
379 polychaetes (*Abyssoninoe* sp., *Clymenura* sp., *L. boeckii*, *Nereimyra aphroditoides*, *Pholoe* sp.),  
380 ostracods and the bivalve *A. moerchi*. Although the associated community structure differs with  
381 that of Daneborg, both display several common dominant species (e.g. *A. moerchi*, *Clymenura*  
382 sp., *L. boeckii*) indicating common ecological characteristics potentially related to their vicinity  
383 in the fjord and similar environmental constraints.

384 Finally, the silty outer fjord assemblage collected at Pass Hytten is dominated by ostracods, the  
385 *Metopa glacialis* amphipod as well as tube-dwelling polychaete species (e.g. *Maldane sarsi*,  
386 *Owenia borealis*, *Euchone* sp.) and crustaceans. Surprisingly, several species described in the  
387 literature as typical of either outer or inner fjord assemblages are found in association in Pass  
388 Hytten. For instance, the sub-surface deposit feeder *M. sarsi* is the most dominant polychaete  
389 although it is reported to be typical of outer fjord assemblages (Gulliksen et al. 1985; Renaud  
390 et al. 2007; Kędra et al. 2013), whereas it is absent from inner fjord areas (Włodarska-  
391 Kowalczyk et al. 2012). Surface deposit-feeding thyasirids and nuculanids are also present,  
392 albeit in moderate abundance, at Pass Hytten, although they are generally representative of  
393 benthic assemblages from inner fjord areas (Włodarska-Kowalczyk and Pearson 2004). The  
394 presence of *Euchone* sp. or *Leitoscoloplos mammosus* polychaetes in Pass Hytten is also  
395 consistent with previous observations made in shallow and deep areas exposed to high  
396 sedimentation rates (Sejr et al. 2010; Kędra et al. 2013; McGovern et al. 2020). Therefore, the  
397 coexistence of several species typical of both outer and middle/inner fjord assemblages at Pass

398 Hytten suggests that although this site is more similar to outer fjord habitats, the presence of a  
399 handful of species adapted to high sediment loads would indicate that this station is approaching  
400 the transition zone between the outer and inner parts of the fjord. This could be explained by  
401 the more interior location of Pass Hytten compared to the other three stations sampled, as well  
402 as its location closer to the Zackenberg River delta. We hypothesize that future sampling, either  
403 in the innermost part of the fjord or closer to the Zackenberg delta, will identify new species  
404 assemblages typical of benthic habitats from areas exposed to glacial and freshwater inputs.

405 *New insights about the functioning of the shallow benthic outer fjord community*

406 The new data from our study provide a better understanding of the importance of trophic  
407 adaptations for benthic community functioning in outer fjord habitats. A complete analysis of  
408 the trophic structure of benthic assemblages has already been conducted at Pass Hytten and Kap  
409 Breusing by Bridier et al. (2021). It appears that several dominant species in our study are  
410 known to have high trophic plasticity (i.e. feeding on several carbon sources, De Cesare et al.  
411 2017; Bridier et al. 2021). The bivalves *Astarte moerchi*, *Astarte elliptica*, *Hiatella arctica*, and  
412 *Musculus discors* in Young Sound appear to feed on a range of carbon sources such as  
413 phytoplankton, benthic organic matter (i.e. macroalgae and/or microphytobenthos) or  
414 sedimentary organic matter (detritus, Bridier et al. 2021). For some species (e.g. the astartidae  
415 *A. moerchi* and *A. elliptica*), the proportion of alternative organic matter resources can even  
416 reach almost 50% of their diet (Bridier et al. 2021). In addition, other species abundant in Young  
417 Sound are known to be omnivores (i.e. feeding on several trophic levels, McMeans et al. 2015;  
418 Bridier et al. 2021). This is particularly true for the brittle star *Ophiocten sericeum*, the  
419 amphipod *Metopa glacialis* and the polychaete *Nereis zonata*, which were observed in high  
420 abundance at all stations. Although the brittle star *O. sericeum* is generally considered a surface  
421 feeder, it falls between the primary and the secondary consumer levels in Young Sound (Bridier  
422 et al. 2021). Conversely, the polychaete *N. zonata* and the amphipod *M. glacialis* exhibit trophic  
423 levels below those of the strict predators with which they are generally associated (Bridier et  
424 al. 2021). These variations in trophic position show that these species are capable of feeding at  
425 multiple trophic levels, feeding on sources of primary organic matter when these are available  
426 in the ecosystem, or feeding at higher trophic levels when the availability of primary production  
427 is low (McMeans et al. 2015; Bridier et al. 2021). All these trophic adaptations likely reflect  
428 the low pelagic productivity of Young Sound, which ranks among the least productive coastal  
429 ecosystems in the world, with pelagic primary production estimated at 10 g C m<sup>-2</sup> yr<sup>-1</sup> (Rysgaard  
430 et al. 1999; Cloern et al. 2014). Trophic plasticity might therefore allow species to adapt to this

431 low pelagic primary production by optimizing carbon flows through food webs through the  
432 exploitation of all carbon sources available in the ecosystem (e.g. microphytobenthos,  
433 macroalgae, detritus, Mincks et al. 2008; Renaud et al. 2015). Omnivory may also help benthic  
434 species to adapt to low pelagic primary production by reusing carbon that has already passed  
435 through the food web a second time, thus participating in the recycling of carbon circulating in  
436 the community.

437 The present study complements previous work by Bridier et al. (2021) on the functioning of  
438 benthic communities in the shallow habitats of Young Sound. Although Bridier et al. (2021)  
439 highlighted the existence of trophic adaptations for several benthic species in Young Sound,  
440 the paucity of information on benthic community structure prevents drawing conclusions on  
441 the functioning of benthic communities in outer fjord habitats. The new data presented here  
442 show that the observations on community trophic adaptations involve species that are  
443 numerically dominant in the inventoried benthic habitats (i.e. the rocky, sandy and silty outer  
444 fjord assemblages). For example, the species *H. arctica*, *M. discors* (characterized by high  
445 trophic plasticity) and *N. zonata* (characterized by high levels of omnivory) alone account for  
446 66.5% of the total abundance at Kap Breusing. It could be argued that biomass data would have  
447 been more appropriate than abundance data to discuss the role of these latter species in  
448 community carbon fluxes. However, considering that all species mentioned as omnivorous or  
449 trophically plastic are larger than the rest of the species identified in this study, the biomass  
450 dominance of these individuals is likely greater than their numerical dominance, suggesting  
451 then an even greater role for community carbon fluxes. Trophic adaptations thus play a  
452 predominant role in the functioning of the outer fjord benthic communities, which feed only  
453 partially on fresh organic matter from pelagic primary production. These results also highlight  
454 the crucial role of benthic primary producers in the functioning of shallow Arctic coastal  
455 ecosystems, whose importance is often underestimated, although they represent a significant  
456 carbon source (Attard et al. 2016; Gaillard et al. 2017; Ørberg et al. 2022). With climate change,  
457 increased meltwater inputs will have multiple consequences for Young Sound's pelagic  
458 environments, including impacts on pelagic primary production, organic matter quality and  
459 organic matter transfers to zooplankton (Meire et al. 2017; Middelbo et al. 2018; Bridier et al.  
460 2019). The existence of trophic adaptations in several species that dominate the benthic  
461 communities of the outer fjord (and therefore potentially represent major carbon transfer  
462 pathways) could thus be a crucial advantage for their resilience in the face of upcoming  
463 environmental changes. In the future, it would be important to investigate the community and

464 trophic structures of the inner fjord to observe how the dominance of these trophic adaptations  
465 evolves in habitats exposed to stronger meltwater inputs and higher trophic constraints. In  
466 addition, it would also be interesting to describe the structure of benthic communities and food  
467 webs in the deeper basins of the fjord (up to 350 m), which do not benefit from local benthic  
468 primary production.

#### 469 **Conclusion and further directions of research**

470 The species list obtained in this study is a substantial addition to the inventory carried out two  
471 decades ago, adding 66 new species to the 100 already listed by Sejr et al. (2000). The sampling  
472 of three new stations (Pass Hytten, Basalt Island, Kap Breusing) in addition to the station  
473 previously described by Sejr et al. (2000) (i.e. Daneborg) also provides new insights into the  
474 spatial variability of benthic communities in this fjord. Although the community structure  
475 differs significantly between the four stations, these differences remain small overall,  
476 suggesting that these stations all belong to the same habitat type. Comparison of the community  
477 structure with that of the literature suggests that the four stations are characteristic of outer fjord  
478 assemblages.

479 Young Sound is a unique location in the Arctic where annual monitoring has been conducted  
480 for over 20 years by the Greenland Ecosystem Monitoring - MarineBasis Program. This fjord  
481 is therefore an ideal candidate for future biodiversity inventories in northeast Greenland which  
482 is one of the Arctic regions with the least studied benthic communities (Piepenburg et al. 2011).  
483 Additional sampling will undoubtedly be needed to improve our knowledge of the structure,  
484 dynamics and functioning of the Young Sound benthic habitats and associated biodiversity.  
485 This sampling should be prioritized in the deeper silty basins (up to 350 m), but also in the inner  
486 (i.e. Tyrolerfjord) and intermediate sections of the fjord, which are more exposed to strong  
487 freshwater inputs, turbidity and sedimentation. On the opposite, the fjord mouth should also be  
488 explored to sample various habitats made up of coarse sands, gravels or potentially biogenic  
489 habitats. According to the literature, the innermost parts of the fjord should be dominated by  
490 opportunistic species (e.g. the cirratulid polychaete *Chaetozone* spp.) and mollusks feeding on  
491 surface deposits such as small thyasirid and nuculanid bivalves (e.g. Włodarska-Kowalczyk  
492 and Pearson 2004; Sejr et al. 2010; Włodarska-Kowalczyk et al. 2012), which seem to be  
493 underrepresented in the samples analyzed in this study. Finally, the more oceanic part of the  
494 fjord should contain a wide variety of benthic habitats characterized by a high and original



495 biodiversity, including biogenic habitats (e.g. maerl/rhodolith or horse mussel beds could be  
496 typical in such an environment).

497 These data will be crucial for future predictions on the evolution of benthic communities in  
498 response to future global changes, and in particular increased freshwater inputs to coastal areas  
499 in response to the accelerated melting of the Greenland Ice Sheet (Mouginot et al. 2019; Höning  
500 et al. 2023). Although the structure of benthic communities from sill fjords is generally  
501 considered to be stable on multi-decadal scales (Renaud et al. 2007), the long-term dynamics  
502 of benthic communities in Arctic fjords are still poorly understood. To date, few long-term  
503 monitoring studies have been carried out in the Arctic, and to our knowledge none in East  
504 Greenland. It would therefore be important to carry out such monitoring to ensure community  
505 stability over longer time scales, especially when warmer water masses of Atlantic origin are  
506 increasing along the Northeast Greenland Shelf (Gjelstrup et al. 2022) and could drive an  
507 expansion of boreal species into the region. Complementary sampling methods, including  
508 scientific SCUBA diving, have been used in other Arctic regions (e.g., Svalbard) to study  
509 shallow benthic communities (i.e., < 30 m, e.g., Jørgensen and Gulliksen 2001). For example,  
510 image analysis based on photographs taken by SCUBA divers has been used to study small-  
511 scale spatial variability in benthic communities (Jørgensen and Gulliksen 2001) or long-term  
512 successional dynamics following episodes of disturbance or climate change (Beuchel et al.  
513 2006; Beuchel and Gulliksen 2008; Al-Habahbeh et al. 2020). Applying these methods to future  
514 studies in Young Sound could be an interesting way to obtain complementary data on benthic  
515 communities and would also allow comparative studies with Svalbard fjords, which have been  
516 studied for several decades.

517

## 518 **Author contributions**

519 FO, LC, MKS conducted the field sampling campaign. GB, LP, JJ, TB, MLD and JG conducted  
520 species identifications. GB carried out the statistical analyses, interpreted the results and wrote  
521 the original draft of the manuscript. FO, LP, JJ, LC, MKS, TB, MLD, and JG helped to interpret  
522 the results and write the manuscript. FO, LC, MKS, and JG obtained the necessary funding for  
523 the study. FO, LC, and JG developed the sampling design. All authors read and approved the  
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## 536 **Statements and Declarations**

537 **Competing Interests:** The authors declare that they have no known competing financial  
538 interests or personal relationships that could have appeared to influence the work reported in  
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540 **Data availability:** All data supporting the results and analyses in this article are available in  
541 the tables included in Online Resource 1.

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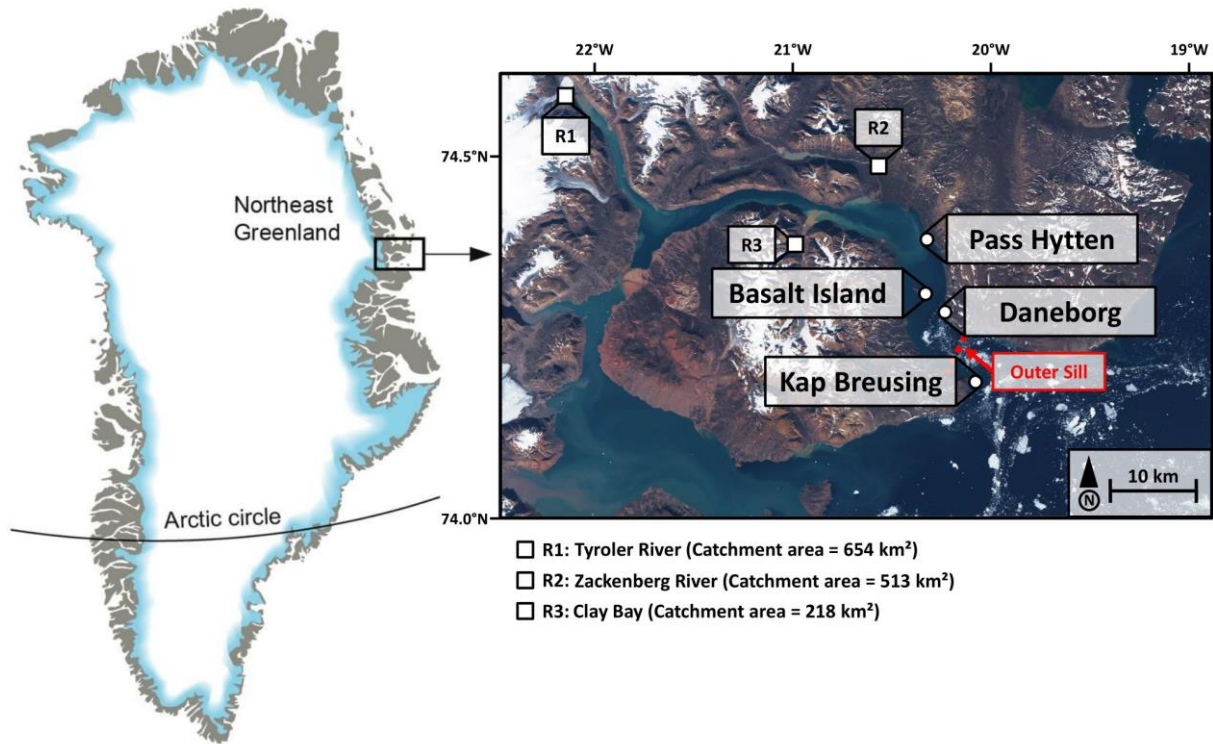
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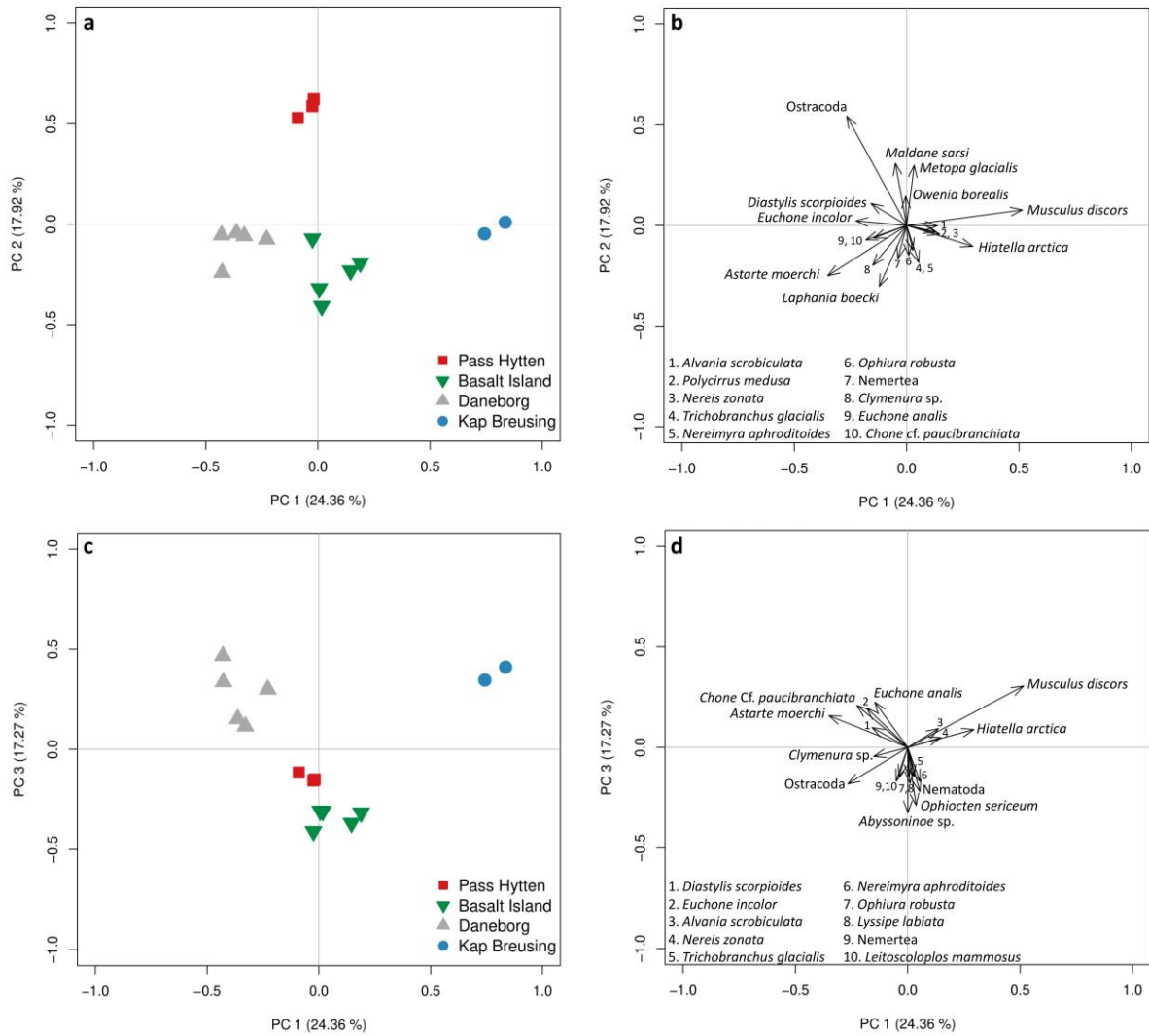
785 **Figures**

786 **Figure 1**



787

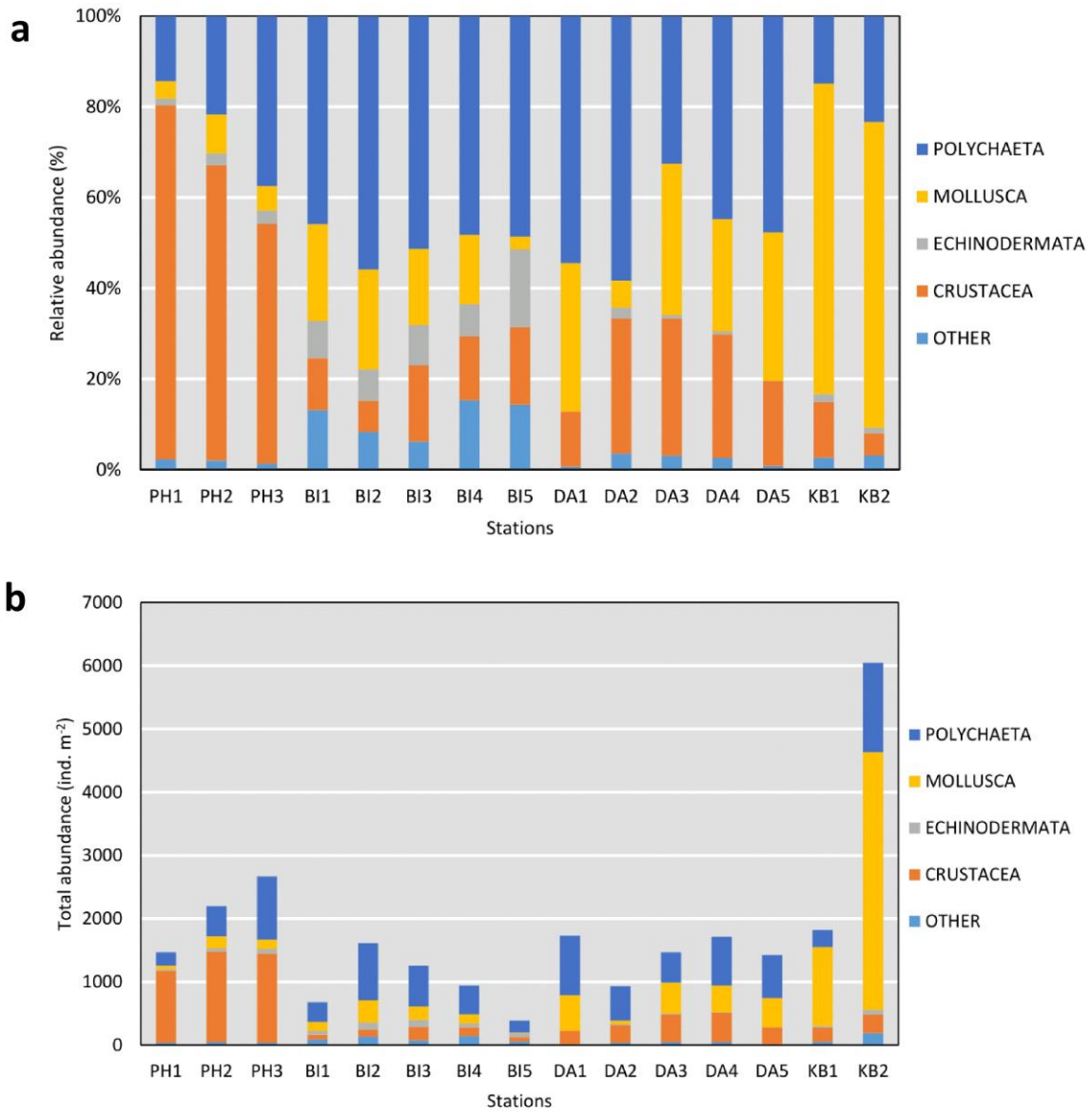
788 **Figure 1** Map showing the four sampling stations (white dots) and the three main rivers (white  
789 squares) in Young Sound. River catchment areas are from Bendtsen et al. (2014)



791

792 **Figure 2** Principal component analyses based on Hellinger-transformed fauna abundances.  
 793 Upper plots (a, b) show the first two principal components while lower plots show the first and  
 794 third principal components. Left plots (a, c) represent the position of each replicate while right  
 795 plots (b, d) represent the position of the first twenty species that explain at least 60 % of the  
 796 spatial variation

797



799

800 **Figure 3** Relative abundance (% , a) and total abundance (ind. m<sup>-2</sup>, b) of major taxonomic groups  
 801 from Pass Hytten (from PH 1 to PH 3), Basalt Island (from BI 1 to BI 5), Daneborg (from DA  
 802 1 to DA 2) and Kap Breusing (KB 1 & KB 2) stations

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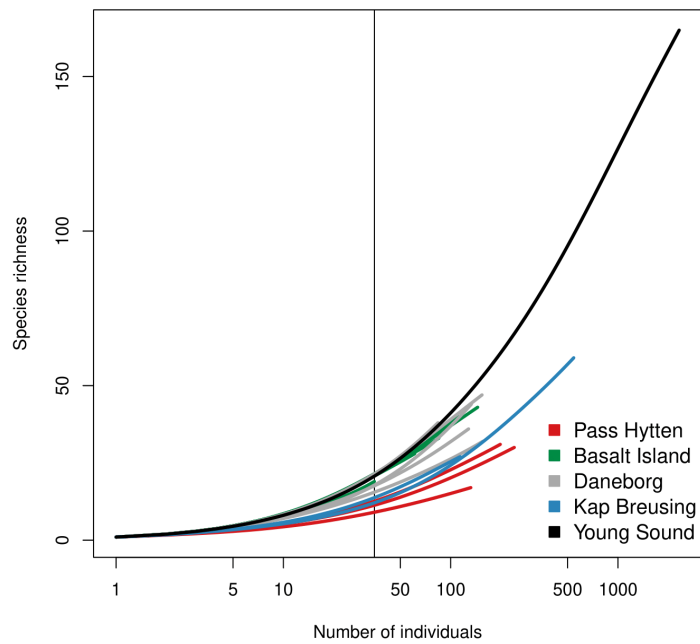


805

806 **Figure 4** Submarine landscapes of Pass Hytten (a, depth = 18 m), Basalt Island (b, depth = 21.5  
807 m), Daneborg (c, depth = 20 m) and Kap Breusing stations (d, depth = 20 m). Photo credits:  
808 Erwan Amice

809

810 **Figure 5**



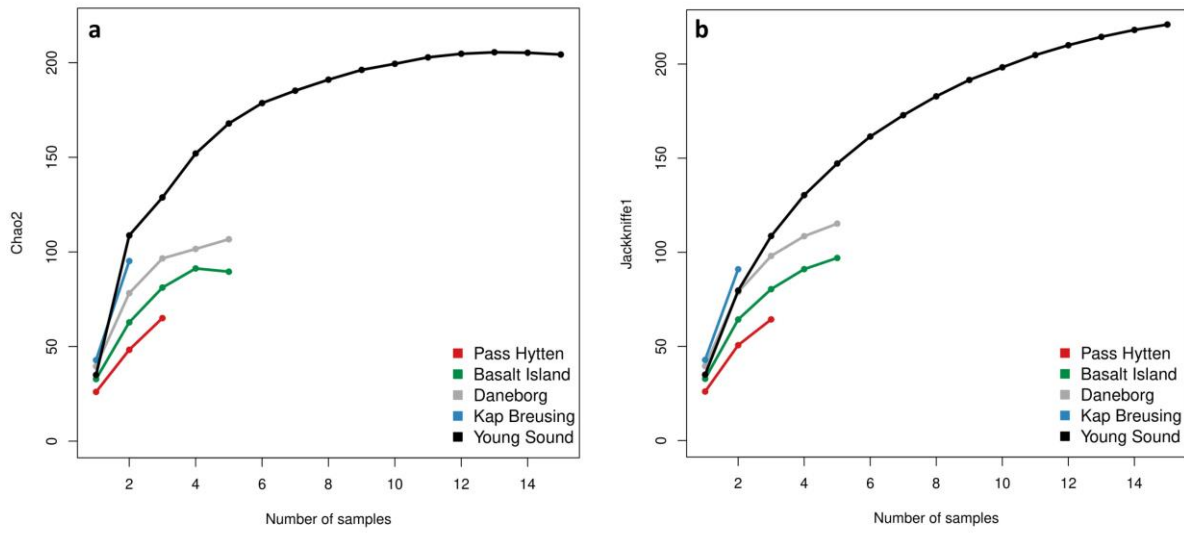
811

812 **Figure 5** Accumulation curves for each replicate from Pass Hytten, Basalt Island, Daneborg  
813 and Kap Breusing stations and the entire Young Sound site (combining 15 replicates). The  
814 vertical line represents the number of individuals selected for comparisons between replicates  
815 (35 individuals)

816



817 **Figure 6**



818

819 **Figure 6** Chao2 (a) and Jackknife1 (b) accumulation curves calculated for Pass Hytten, Basalt  
820 Island, Daneborg, Kap Breusing and the four stations combined.

821

822 **Tables**

823 **Table 1** Geographic coordinates, depth, total area sampled, and material collected (with number  
824 of replicates in brackets) for each station

Station	Latitude	Longitude	Depth (m)	Total sampled area (m <sup>2</sup> )	Collected material
Pass Hytten	N 74.41'	W 20.33'	18.0	0.2700	Macrofauna ( <i>n</i> = 3) and sediment ( <i>n</i> = 1)
Basalt Island	N 74.33'	W 20.36'	21.5	0.4500	Macrofauna ( <i>n</i> = 5)
Daneborg	N 74.31'	W 20.24'	20.0	0.4500	Macrofauna ( <i>n</i> = 5)
Kap Breusing	N 74.21'	W 20.11'	20.0	0.1525	Macrofauna ( <i>n</i> = 2) and sediment ( <i>n</i> = 3)

825

826 **Table 2** List of the ten most abundant taxa for each station, with their relative abundance (%),  
 827 their cumulative relative abundance (% Cum.), their abundance (Abund., number of individuals  
 828 per square meter [ind. m<sup>-2</sup>]) and the Pielou index (mean ± standard error of the mean) of the  
 829 corresponding station (calculated at the scale of the whole community)

Species	Phylum	Abund.	%	Cum. %
<b>Pass Hytten</b>		[Pielou index = 0.622 ± 0.027]		
Ostracoda Latreille, 1802	Arthropoda	996.3	47.2	47.2
<i>Metopa glacialis</i> (Krøyer, 1842)	Arthropoda	185.2	8.8	56.0
<i>Maldane sarsi</i> Malmgren, 1865	Polychaeta	166.7	7.9	63.9
<i>Owenia borealis</i> Koh, Bhaud & Jirkov, 2003	Polychaeta	114.8	5.4	69.3
<i>Leitoscoloplos mammosus</i> Mackie, 1987	Polychaeta	63.0	3.0	72.3
<i>Diastylis scorpioides</i> (Lepechin, 1780)	Arthropoda	63.0	3.0	75.3
<i>Abyssoninoe</i> sp. Orensanz, 1990	Polychaeta	48.1	2.3	77.5
<i>Lysippe labiata</i> Malmgren, 1866	Polychaeta	44.4	2.1	79.6
<i>Euchone incolor</i> Hartman, 1965	Polychaeta	40.7	1.9	81.6
<i>Ophiocten sericeum</i> (Forbes, 1852)	Polychaeta	37.0	1.8	83.3
<b>Basalt Island</b>		[Pielou index = 0.916 ± 0.004]		
<i>Laphania boeckii</i> Malmgren, 1866	Polychaeta	64.4	6.6	6.6
Ostracoda Latreille, 1802	Arthropoda	57.8	5.9	12.5
<i>Abyssoninoe</i> sp. Orensanz, 1990	Polychaeta	55.6	5.7	18.2
<i>Nereimyra aphroditoides</i> (O. Fabricius, 1780)	Polychaeta	55.6	5.7	23.9
Nematoda	Nematoda	55.6	5.7	29.6
<i>Ophiocten sericeum</i> (Forbes, 1852)	Echinodermata	51.1	5.2	34.9
<i>Astarte moerchi</i> Hopner Petersen, 2001	Mollusca	46.7	4.8	39.6
<i>Clymenura</i> sp. Verrill, 1900	Polychaeta	42.2	4.3	44.0
<i>Harmothoe</i> cf. <i>imbricata</i> (Linnaeus, 1767)	Polychaeta	42.2	4.3	48.3
<i>Pholoe assimilis</i> Örsted, 1845	Polychaeta	40.0	4.1	52.4
<b>Daneborg</b>		[Pielou index = 0.834 ± 0.022]		
<i>Astarte moerchi</i> Hopner Petersen, 2001	Mollusca	271.1	18.6	18.6
Ostracoda Latreille, 1802	Arthropoda	171.1	11.8	30.4
<i>Euchone incolor</i> Hartman, 1965	Polychaeta	91.1	6.3	36.6
<i>Laphania boeckii</i> Malmgren, 1866	Polychaeta	75.6	5.2	41.8
<i>Euchone analis</i> (Krøyer, 1856)	Polychaeta	62.2	4.3	46.1
<i>Chone</i> cf. <i>paucibranchiata</i> (Krøyer, 1856)	Polychaeta	57.8	4.0	50.1
<i>Clymenura</i> sp. Verrill, 1900	Polychaeta	57.8	4.0	54.0
<i>Diastylis scorpioides</i> (Lepechin, 1780)	Arthropoda	51.1	3.5	57.6
<i>Chaetozone</i> sp. 1 Malmgren, 1867	Polychaeta	31.1	2.1	59.7
<i>Harmothoe</i> cf. <i>imbricata</i> (Linnaeus, 1767)	Polychaeta	24.4	1.7	61.4
<b>Kap Breusing</b>		[Pielou index = 0.634 ± 0.062]		
<i>Musculus discors</i> (Linnaeus, 1767)	Mollusca	1875.4	43.7	43.7
<i>Hiatella arctica</i> (Linnaeus, 1767)	Mollusca	800.0	18.7	62.4
<i>Harmothoe</i> cf. <i>imbricata</i> (Linnaeus, 1767)	Polychaeta	242.6	5.7	68.0
<i>Nereis zonata</i> Malmgren, 1867	Polychaeta	177.0	4.1	72.2
<i>Polycirrus medusa</i> Grube, 1850	Polychaeta	85.2	2.0	74.2
<i>Alvania scrobiculata</i> (Møller, 1842)	Mollusca	78.7	1.8	76.0
<i>Mya truncata</i> Linnaeus, 1758	Polychaeta	65.6	1.5	77.5
<i>Polycirrus latidens</i> Eliason, 1962	Polychaeta	52.5	1.2	78.7
<i>Nereimyra aphroditoides</i> (O. Fabricius, 1780)	Polychaeta	45.9	1.1	79.8
<i>Apherusa sarsi</i> Shoemaker, 1930	Arthropoda	45.9	1.1	80.9

