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Mesozooplankton in the Southern Ocean: spatial and temporal patterns from
Discovery Investigations

Peter Ward, Geraint A. Tarling, Sally E. Thorpe

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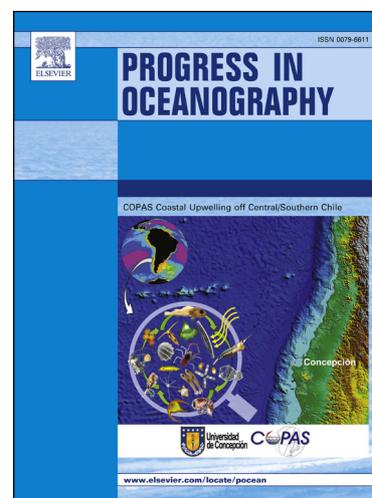
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1 **Mesozooplankton in the Southern Ocean: spatial and temporal patterns from *Discovery***

2 ***Investigations***

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4 Peter Ward, Geraint A. Tarling, Sally E. Thorpe

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7 British Antarctic Survey,

8 Natural Environment Research Council,

9 High Cross,

10 Madingley Road,

11 Cambridge,

12 CB3 0ET,

13 UK

14 pwar@bas.ac.uk

15 Tel: +44 (0)1223 221564

16

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24 Running head: Mesozooplankton from *Discovery Investigations*

25 ABSTRACT

26 Mesozooplankton samples taken during the *Discovery Investigations* in the Southern
27 Ocean in the 1930's were analysed from a series of 5 transects along 80°W. The samples
28 provide a unique level of depth-discrete resolution across large spatial scales, over most of
29 the productive austral season. Stratified net hauls were taken between 0 and 1000 m within
30 the period December 1933 to November 1934. Within the epipelagic (0-100 m), median
31 zooplankton abundance (278 ind. m⁻³) was ~ 22 times greater than at 1000 m. A 3-4 fold
32 variability of abundance in the epipelagic contrasted with depths > 250 m where variability
33 was <1 fold. Depth was the strongest factor separating samples (ANOSIM, R =0.66
34 $p=0.1%$), with a clear biological distinction between epipelagic and upper and lower
35 mesopelagic horizons. Results from multi-dimensional scaling indicated that, when plankton
36 abundance was integrated over all depth horizons, 3 different groups could be identified.
37 These 'communities' were consistent with the spatial extents of Antarctic, Polar Frontal
38 Zone, and sub-Antarctic water-mass regimes. Such groupings became less distinct when
39 considering only deeper horizons (500-1000 m) and excluding seasonal migrants. Seasonal
40 signals across all data became less distinct with depth. Rarefaction analysis indicated that
41 diversity increased with depth. Although depth alone was the most important influence on
42 sample diversity, ($r^2=0.60$), water mass regime and month improved the fit ($r^2=0.71$). Overall
43 plankton diversity was highest in the sub-Antarctic zone. Following atmospheric and ocean
44 warming that has taken place close to the study area in the last 80 years we hypothesise that
45 species richness may increase in the Antarctic water masses as sub-Antarctic species
46 increasingly encroach south.

47

48 **1. Introduction**

49 Within the world's oceans far more is known about the plankton occupying the epipelagic
50 than the deeper horizons. This is particularly true of the Southern Ocean where the vast
51 majority of investigations have taken place within the surface 200 m. Fewer studies have
52 taken into account the deeper water column and have largely emphasised taxonomy (e.g.
53 Park, 1978, 1983, 1993, Bradford-Grieve, 1994, 1999), or have focused on descriptions of
54 planktonic life cycles and vertical distribution (Andrews, 1966; Voronina, 1972; Atkinson,
55 1991, Marrari et al., 2011; Wiebe et al., 2011).

56 Fundamental knowledge of the distribution of plankton abundance and biomass within
57 the Southern Ocean has largely come from the *Discovery Investigations* carried out during
58 the early to middle part of the 20th century. Plankton samples taken with the N70V net have
59 variously been used for regional comparisons of standing stock in the Southern Ocean
60 (Foxton 1956), understanding the circulation of macroplankton (Mackintosh, 1937),
61 elucidating life cycles and distribution of copepods (Andrews, 1966; Ommaney, 1936;
62 Atkinson, 1991) as well as chaetognaths (David, 1955, 1958). Foxton (1956) used over 2,100
63 stratified N70V net samples (0-1000 m), from 366 *Discovery* stations within the Southern
64 Ocean to describe plankton volume on a regional and seasonal basis. Hopkins (1971)
65 similarly sampled to 2000 m in the Pacific sector (~75°W-160°W), obtaining 375 Bé net
66 samples and found biomass distributions to be broadly comparable, both regionally and
67 seasonally with Foxton's work. Important findings from both of these studies were that
68 within the top 1000 m, seasonal variation in plankton biomass is marginal and also that
69 increases in biomass were observed in the region of the Polar Front (PF). However, Foxton
70 (1956) recognised that the N70V was poor in terms of catching krill which are particularly
71 abundant south of the Polar Front, in the Antarctic Zone (AZ). Important studies by Hopkins
72 and co-workers (e.g. Hopkins, 1985; Hopkins and Torres, 1988; Hopkins et al., 1993a) have
73 investigated the biology and trophic ecology of the water column, including the deeper water

74 horizons in the Weddell Sea and adjacent marginal ice-zones (MIZ), as well as providing
75 community descriptions and insights into plankton diversity down the water column.
76 Atkinson and Sinclair (2000) used *Discovery* samples to assess the extent of seasonal
77 migration among the plankton and to clarify zonal distributions. Ward et al. (1995, 2006) and
78 Ward and Shreeve (1999) documented seasonal changes in community distribution and
79 biomass down to 1000 m around South Georgia and Marrari et al. (2011) investigated the
80 vertical distribution of zooplankton and habitat partitioning in the deeper water column in the
81 Marguerite Bay region of the Western Antarctic Peninsula (WAP).

82 Over the last 20 or so years, numerous studies have defined Southern Ocean plankton
83 community structure (e.g. Hosie, 1994; Errhif et al., 1997; Pakhomov et al., 2000, Ward et al.
84 2003, 2006). This has given us a near consistent view of epipelagic communities which are
85 bounded by the physical gradients and discontinuities often found at frontal zones. What is
86 presently unclear is whether such distinctions between plankton communities exist within the
87 mesopelagic and what patterns of diversity are discernible with respect to water mass and
88 depth. Studies of the mesopelagic (200-1000 m) elsewhere have emphasised that it is here
89 that processes govern the efficiency by which particulate organic carbon is transported to the
90 sea floor (Tréguer et al., 2003; Steinberg et al., 2008; Robinson et al. 2010) although we
91 generally know little regarding the abundance, biomass and vertical structure of the major
92 taxa involved. In the Southern Ocean Hopkins and Torres (1988) found diversity increased
93 below the epipelagic in the Weddell Sea, and Hopkins et al. (1993a, b) have shown that depth
94 was an important factor in determining spatial distance in an ordination of a series of samples
95 taken in the Scotia Sea MIZ in winter. A recent paper on diversity in the Arctic Ocean has
96 also shown depth to be a major structuring element for species distributions and communities
97 (Kosobokova et al., 2011 and references therein).

98 Extensive sampling of the mesopelagic rarely takes place these days, largely on
99 grounds of cost. Fortunately, *Discovery Investigations* routinely sampled the Southern Ocean
100 to 1000 m with the N70V net and many of these sets, taken ~80 years ago, are still available
101 for study. The aim of this investigation was to examine the basic properties (abundance,
102 species composition and diversity) of a series of plankton samples taken during 5 transects
103 completed by *Discovery Investigations* along 80°W in the Pacific sector just west of Drake
104 Passage, with a further aim of examining the spatial variability (horizontal and vertical) of the
105 plankton community structure. The transects, occupied between December 1933 and
106 November 1934, crossed three zones of the Antarctic Circumpolar Current, extending from
107 the AZ in the south, through the Polar Frontal Zone (PFZ) to the sub-Antarctic Zone (SAZ) in
108 the north (Fig. 1).

109 In this region of the Southern Ocean, temperatures have risen faster than elsewhere in
110 the Southern Hemisphere, with surface summer temperatures rising more than 1°C during the
111 last part of the 20th century (Meredith and King, 2005). Such changes have been predicted to
112 have detrimental effects on various marine species including Antarctic krill, a key species in
113 the Southern Ocean foodweb with a known dependence on the physical environment
114 (Atkinson et al., 2004), and marine benthos, which is also sensitive to temperature change
115 (Peck et al., 2004). In investigating zooplankton distributions from samples collected 80
116 years ago, we are establishing what patterns were prior to subsequent ocean changes and
117 potentially forming a baseline against which the future impacts of such physical change
118 might be measured.

119 **2. Method**

120 *2.1 Zooplankton sampling and processing*

121 A total of 41 stations were occupied during a series of 5 transects completed during
122 the *Discovery Investigations*. The transects extended from ~ 68°S to 55°S along 80° W and
123 were investigated during December 1933, March, September, October and November 1934
124 (Fig. 1). The position, number and horizontal spacing of stations varied between transects (5-
125 10 stations per transect, transect lengths of ~690-1400 km). From these stations, 248
126 plankton samples were taken with an N70V net used extensively during *Discovery*
127 *Investigations* to obtain stratified plankton samples. Of these, 9 samples were not located in
128 the Natural History Museum collections and were presumed missing. Others had suffered
129 some spillage during capture and initial processing, as indicated on the sample labels.
130 Although these latter samples were examined, species counts were not included in subsequent
131 data analysis reducing the total number of stations to 39 and samples to 215.

132 Full details of the N70V net construction are given in Kemp et al. (1929), but briefly
133 it was a 70 cm dia. ring-net with two grades of silk mesh; a forward section of 40 threads per
134 inch (TPI) and a rear section of 74 TPI equivalent to ~195µm and 440 µm respectively (Ward
135 et al. 2012), and a collar of ~6 mm knotted mesh. The net was deployed in an open state and
136 lowered to the bottom depth of the horizon to be fished before being hauled vertically
137 upwards at a rate of 1 m sec⁻¹. Net closure was accomplished by use of a messenger sent
138 down the wire at a predetermined time which triggered a closing mechanism as the net
139 reached the upper depth of the fished horizon. This released the net bridles allowing a rope
140 encircling the net and attached to the closing mechanism to throttle the net. Dependent on
141 water depth, up to six samples were obtained as follows: 50 m to surface, 100–50 m, 250–100
142 m, 500–250 m, 750–500 m and 1,000–750 m. A full suite of vertical net samples was carried
143 out in ~1.5-2hr.

144 The samples were taken almost 80 years ago and preserved in formalin. As might be
145 expected, overall condition was variable and not very good in the main. Crustaceans were
146 generally intact, although brittle, and internal tissue was often absent. Specimens of the
147 copepod *Calanoides acutus* had been removed from many samples and preserved separately.
148 A record of numbers removed was appended to the label in each jar, although the stage
149 structure of the absent fraction is unknown. Appendicularians were also often difficult to
150 identify because of tissue deterioration and thecosome pteropods likewise. Chaetognaths
151 were generally represented only by their jaws with the soft body tissue having dissolved
152 away. Numbers were therefore estimated by counting jaws and dividing by 2. Further
153 information on chaetognath species composition and other elements of the catch was obtained
154 from digitised records of photographs taken of *Discovery* logbook pages pertaining to each
155 haul (data provided by Andrew Mackey). Four species of chaetognaths were identified in
156 these logbooks, *Eukrohnia hamata* and 3 species of the genus *Sagitta*; *S. gazellae*, *S. maxima*
157 and *S. planctonis*. A comparison of *Discovery* logbook entries with jaw counts indicated that
158 overall, jaw counts underestimated abundance by ~10% (average jaw counts as a proportion
159 of *Discovery* counts = 0.91 (± 0.781)). However not all logbook entries were consistent in
160 recording chaetognath numbers and a number of blank entries were apparent although
161 chaetognaths were present in the corresponding samples. For the sake of completeness we
162 have therefore decided to use abundance data generated from jaw counts/2 .
163 Initially samples were looked at in their entirety and rare and/or large specimens removed and
164 counted. Samples were then split successively using a folsom splitter until the resultant
165 aliquots were estimated to contain between 100-200 individuals which were again identified
166 and counted. Finally, further splitting took place to produce aliquots from which counts of the
167 smaller and more numerous size fractions were made. Both aliquots from each final split
168 were counted. An average of 753 (± 297) individuals were counted per sample and, across all

169 samples, a total of 212 taxa /species /stages were identified (see Web appendix). Flow meters
170 were not used by *Discovery Investigations* and volume swept has been estimated from net
171 diameter and the depth of the water column that each net fished through. Thus, $\sim 19 \text{ m}^{-3}$ of
172 water was swept in a 50 m vertical tow, $\sim 58 \text{ m}^{-3}$ in 150 m tow, and $\sim 96 \text{ m}^{-3}$ in a 250 m tow.

173

174 2.2 Oceanography

175 Temperature and water for salinity analysis were obtained from each station with
176 Ekman reversing bottles fitted with protected and unprotected reversing thermometers.
177 Thermometers were scaled to 0.1°C and data presented in the station lists to two decimal
178 places. Salinity was analysed by titration against silver nitrate of known strength using
179 potassium chromate as an indicator (Kemp et al., 1929).

180 We used water mass properties at each of the *Discovery* stations to locate the
181 positions of the sub-Antarctic Front (SAF) and Polar Front (PF) on each transect, enabling
182 stations to be classified into three zones of the Antarctic Circumpolar Current (ACCZ): the
183 SAZ, PFZ and AZ. The frontal locations were determined from potential temperature-
184 salinity curves and confirmed with vertical sections of potential density along the transects
185 (Orsi et al, 1995; Read et al, 1995). Two transects (March and December) sampled south of
186 the southern ACC front but this front was not used to group stations in subsequent analyses.

187 2.3 Statistical analysis

188 PRIMER (v6) (Clarke and Gorley, 2006) was the principal statistical package used to
189 analyse the species-by-samples dataset. Routines were carried out on the species by stations
190 data matrix using $\log(x+1)$ transformed standardised data (ind. m^{-3} or ind. m^{-2} (0-1000 m)), to
191 determine Bray-Curtis similarities before undertaking hierarchical clustering of data into
192 sample groups (CLUSTER), and ordination (Multi dimensional scaling (MDS)). Analysis of
193 similarity (ANOSIM) was used to test for differences between resultant groups. ANOSIM

194 operates on the resemblance matrix and is approximately analogous to standard univariate
195 analysis of variance (ANOVA). Similarity percentages (SIMPER) was used to examine
196 taxonomic contributions towards group structure.

197 The PAST statistical package (Hammer et al. 2001), was used investigate species
198 diversity and specifically to undertake sample rarefaction (Mao tau) on the data converted
199 into a presence or absence matrix with respect to depth horizon. Rarefaction generates the
200 expected number of species in a small collection of n samples drawn at random from the
201 large pool of N samples (Gotelli and Colwell, 2001). The shape of the curve is steep at first
202 and then tends to plateau as only the rare species remain to be sampled. Rarefaction curves
203 generated from the present study were mostly beyond the steep phase but had yet to reach the
204 plateau phase. Therefore, we extrapolated the curves to determine (i) the expected number of
205 taxa within each depth interval and (ii) the level of sampling effort required to identify 90%
206 of this number of taxa. Both a 3-parameter power function and Michaelis-Menten function
207 were fitted to the rarefaction curves, with the former achieving the best levels of fit
208 ($R^2 > 0.99$). The power functions were then extended to three times the original sample size
209 (from ~36 to 108 samples) for each depth interval, following Colwell et al. (2004). The
210 rarefaction curves and extrapolations for each depth interval were plotted together for
211 comparative purposes.

212

213 **3. Results**

214 *3.1 Physical conditions along the transect*

215 Surface and subsurface positions of the fronts were located between the same station
216 pairs on all transects, with very few stations showing evidence of interleaving that is often
217 seen close to fronts. Our classification agrees with that of Foxton (1956) who partitioned

218 plankton standing crop into sub-Antarctic and Antarctic zones, although we additionally
219 define the PFZ (Table 1). At the southernmost stations of each of the transect passes,
220 temperature in the top 100 m of the water column was always less than 0 °C. At the
221 northernmost stations at the same depth, temperature ranged from ~2.7—7.75 °C. A
222 latitudinal increase of 1 °C per 155-200 km within the top 100 m was recorded for each
223 transect, with temperature increasing northwards. The range of the average temperature
224 calculated for the upper 100 m along each transect varied from 4.46 °C on the shortest
225 transect (November) to 8.49 °C on the longer one (March). As expected, this near-surface
226 temperature range was much larger (by a factor of 4.25 on average) than that recorded at
227 1000 m. Within all five transects, the difference between the minimum and maximum
228 temperature at 1000 m was only 2.15 °C.

229 *3.2 Mesozooplankton abundance by depth*

230 Initially we calculated abundance (ind. m⁻³) within each depth horizon and present
231 these data as medians across all months and ACCZs (Table 2, Fig. 2). Median abundance
232 within the top 100 m was significantly higher than in horizons deeper than 250 m (Kruskal–
233 Wallis H=106.44, $p=0.00$). Abundance declined down the water column such that below 750
234 m the median was ~4% of that in the near-surface layers. Within depth horizons, monthly
235 variation in range was greater (absolute and relative) in surface horizons than deeper (factor
236 of 3-4 in top 50 m, 2-3 in 100-50 m and <2 in 250-100 m). Below 250 m the range was <1
237 and broadly equal across months.

238 *3.3 Month vs ACCZs*

239 There are some gaps in the data matrix when considering abundance with respect to
240 ACCZ by month, which makes it difficult to get a strictly comparable view of monthly
241 mesoplankton distributions by ACCZ (Table 1). Data from the SAZ are absent in November

242 and the PFZ is represented by only one sample in November and March. We have therefore
243 plotted data showing variation in abundance across months and depth zones (ie ACCZs
244 pooled) and also by variation across ACCZ and depth (ie months pooled). In the former (Fig
245 3a), the trend was for near-surface abundance to increase from September through till
246 December and then decline slightly by March. In depth horizons below 100 m, no systematic
247 change was observed across months, with the range of values tightly constrained compared to
248 the upper 100 m. Below 250 m, abundance was reduced in each successive depth horizon by
249 almost half in many cases. The ratio of the median abundance in the top 100 m to that in the
250 deepest horizon varied from ~8-31 times across months. Pooling stations by ACCZ,
251 irrespective of time of year (Fig. 3b), indicated that there was a greater range of values within
252 the near-surface AZ compared to elsewhere. Deeper horizons (>500 m) once again appeared
253 quite similar in overall abundance, irrespective of region. The high abundances seen in the
254 near-surface waters of the AZ were due to the presence of limacinid pteropods during
255 December. Median abundance and biomass by depth and ACCZ are presented in Table 2.
256 Sample biomass was estimated from settled volumes provided by Foxton (1956) who
257 included these samples along 80°W in an estimation of zooplankton standing stock in the
258 Southern Ocean. Foxton provided data as displacement volumes (cm^{-3}) which we have
259 converted to wet/dry mass assuming that 1 cm^{-3} equals 1 g wet weight and that dry weight
260 represents 10% of this (see Hopkins 1971). Estimated biomass (wet mass mg m^{-3}) largely
261 reflects patterns of abundance. Summed over the water column, median wet mass (gm^{-2} , 0-
262 1000 m) was uniform, ranging from ~16 g in the AZ and SAZ to ~20 g in the PFZ. Dry mass
263 using the 0.1 conversion advocated by Hopkins (1971) is therefore ~1.6-2.0 gm^{-2} (0-1000 m)
264 across all ACCZs.

265 *3.4 Spatial structure within the sample set*

266 Clustering and multi-dimensional scaling (MDS) of Bray-Curtis similarities (latter
267 not presented) showed that depth was by far the strongest factor in separating samples (Fig.
268 4). Three main clusters were apparent which could be classified as epipelagic (mainly
269 samples taken in the top 100 m), upper mesopelagic (mainly samples from 100-250 m and
270 250-500 m horizons) and lower mesopelagic (mainly depths >500 m). Analysis of similarity
271 (ANOSIM) to determine the extent to which the main clusters reflected depth was carried out
272 using sample depth zone as a factor. A value of $R = 0.66$ ($p = 0.001$), indicated that there was
273 indeed a strong case for viewing the water column as broadly divisible into 3 horizons
274 irrespective of ACCZ. We also tested whether month or time of sampling (day or night) had
275 any significant influence on the clustering and found that both had much lower, yet still
276 significant, values of R ($R = 0.23$, $p = 0.001$ for month; $R = 0.03$, $p = 0.046$ for day/night). The
277 larger magnitude of R (which is an absolute measure of differences between groups, as
278 compared with the p level, which is influenced by sample size) for depth shows that
279 differences occasioned by month are slight in comparison with depth.

280 *3.5 Community analysis*

281 Although depth clearly dominated the way in which the sample specific dataset
282 clustered, we also wished to determine the extent to which ACCZ might be a factor in
283 horizontal variability. For the 27 stations which had a full complement of samples down to
284 1000 m (see Table 1), species stages were aggregated within species resulting in 171
285 species/taxa and abundance (ind. m^{-2} , 0-1000 m). Data were $\log(x+1)$ transformed, before
286 clustering, MDS and subsequent routines were again carried out. We also wished to ascertain
287 whether 'plankton communities' could be defined in waters >500 m deep. By restricting our
288 analysis to the two deepest horizons fished by the N70V we increased the number of stations
289 available for analysis from 27 to 33. However our first iterations of the 0-1000 m data matrix
290 identified the strong influence on defining communities played by seasonal migrants (see

291 Atkinson 1988) and smaller, predominantly epipelagic species such as *Oithona similis*.
292 Therefore we first removed these to produce a species list that better reflected the deepwater
293 fauna (see legend Fig. 5). Following the removal of seasonal migrants and small near-surface
294 copepods, a total of 156 taxonomic categories remained. We examine these analyses in turn.

295 *0-1000 m--* A plot of the MDS ordination on the 0-1000 m data set is presented in Fig
296 5a. At the 73% similarity level station groups comprised an AZ group (9 stns), an AZ/ PFZ
297 group (11 stns) and a predominantly SAZ grouping (7 stns).

298 *500-1000 m--* By restricting our analysis to the two deepest horizons fished by the
299 N70V, we increased the number of stations available for analysis from 27 to 33. On this
300 occasion we observed two major groupings at ~72% similarity, the first of predominantly AZ
301 and PFZ stations and the second a diverse group of PFZ and SAZ stations (Fig. 5b).

302 A summary of ANOSIM performed on these data is presented in Table 3. Global R
303 for both ordinations was significant and among the pairwise comparisons of the ACCZs the
304 strongest difference was between AZ/SAZ. The value of R for the AZ/PFZ and PFZ/SAZ
305 pairwise comparisons more than halved (Table 3). This is consistent with the MDS plots
306 (Figs 5a-b) which suggest that stations along the transect form a continuum, with the spatial
307 extremes of AZ and SAZ stations differing most. We have therefore used the output from
308 ANOSIM and defined station groups with respect to ACCZ for subsequent analysis, rather
309 than trying to account for temporal and other differences in the data. The MDS plots
310 generally show stations thus identified as occupying different parts of the ordination.

311 We then ran the similarity percentages routine (SIMPER) to ascertain which
312 species/taxa were most responsible for within group structure. For the 0-1000 m ordination,
313 the small abundant species contributed most to within group similarity and between group
314 dissimilarity. In table 4 we have presented the 10 species/taxa contributing most to the above

315 for the 3 groupings identified by MDS (Fig. 5a), making a combined total of 20 species.
316 Between 50-57% of within group similarity was accounted for by the tabulated taxa and
317 ~40% of between group dissimilarity. Species/taxa so identified showed a range of
318 distributions and abundances and overall the PFZ stations had the greatest average abundance
319 of plankton (mean $\sim 8.5 \times 10^4$ ind. m^{-2} 0-1000 m, see table 5) although not significantly greater
320 than the other two groups. Among species that often contributed most to dissimilarity
321 between station groups were seasonal migrants including *Calanoides acutus*, *Calanus*
322 *simillimus* and *Subeucalanus longiceps* and species that were widely distributed throughout
323 the water column such as *Ctenocalanus* spp. and *Oithona similis*. The former species are
324 seasonal migrants that spend spring/summer in near-surface waters and reside in deeper
325 waters for the rest of the year. However, differences in the timing of life-cycles with respect
326 to latitude means they are not uniformly distributed with respect to depth over the length of
327 the transect.

328 In the 500-1000 m MDS plot (Fig. 5b) stations were once again broadly arrayed
329 across the ordination. The pairwise comparisons of ACCZs carried out within ANOSIM
330 showed a greater difference between the AZ and SAZ than of either with the PFZ (Table 3).
331 In this case the PFZ/SAZ groupings were not significantly different and we have therefore
332 pooled these before undertaking SIMPER. Again smaller copepod species proved to be the
333 greatest contributors to within group structure. All except two of the first 10 taxa defining
334 within group similarity and between group dissimilarity was the same in each of the two
335 ACCZ groupings (20 taxa in total), with only the rank order changing (Table 6). A number of
336 the deeper dwelling taxa, including the copepods *Metridia curticauda*, *Mormonilla* sp.,
337 *Paraeuchaeta biloba* and Scolecithriciid copepodites, as well as Siphonophora and
338 *Thysanöessa* spp. were particularly important in defining dissimilarity between groups.

339 *3.6 Seasonality in the mesopelagic*

340 One of the questions we wished to answer was whether seasonal changes in
341 abundance could be detected in the deeper depth horizons. We have presented 'spring'
342 (Sept/Oct/Nov) and 'summer' (Dec/Mar) abundances of the 20 most abundant taxa
343 (excluding seasonal migrants) in the 500-1000 m depth horizons across all ACCZs (Table 7).
344 Overall there was a suggestion that one or two taxa were more abundant in summer than
345 spring (e.g. *Oncaea* spp., *Metridia lucens* and *Paroithona* sp.) but only *Spinocalanus* spp. was
346 significantly different and the majority of taxa showed very little difference between our
347 seasonal groupings. However, given the length of the transect, there are likely to be
348 'seasonal' differences observed when comparing stations at one end with another. Within the
349 Scotia Sea, Ward et al. (2004, 2006), have found differences in population age of biomass
350 dominants *Calanoides acutus* and *Rhincalanus gigas*, to be as much as 3 months and possibly
351 more, over transects of similar lengths, sampled quasi-synoptically. The possibility that
352 grouping all stations together might be aliasing seasonality was investigated by looking for
353 'seasonal differences' within ACCZs.

354 We grouped 500-1000 m samples according to season, as before, but undertook
355 comparisons within the AZ, and the PFZ/ SAZ ACCZs combined. The majority of ANOVA
356 comparisons showed no difference between seasons. The most significant differences (5/40)
357 were found in the AZ for Chaetognatha, *Paraeuchaeta antarctica* and Ostracoda where
358 spring averages were all greater than summer ($p < 0.05$). Among PFZ and SAZ stations,
359 Chaetognaths (Spr < Su) and *Spinocalanus* spp. (Spr < Su) were significantly different. None of
360 the taxa which showed significant seasonal differences in one ACCZ showed them in the
361 other.

362 3.7 Diversity

363 The results of the rarefaction analysis are illustrated in Figure 6. The curves represent
364 taxa numbers by depth, pooled across all ACCZs based on between 35-37 samples in each

365 depth horizon. Diversity increased systematically with depth, although the two deepest
366 horizons were virtually indistinguishable. None of the curves reached a plateau indicating
367 that diversity was underestimated. The slope of the curves was broadly identical for the 3
368 horizons within the top 250 m but steeper below 500 m, indicating a greater level of
369 undersampling of the total number of taxa at deeper depths. We have extrapolated the curves
370 for each horizon to 108 samples (approximately tripling the original sampling size within the
371 limits suggested by Colwell et al., 2004) to estimate the number of taxa expected at that
372 sampling intensity. The number of taxa observed in our analysed sample sets was between
373 70-80% of the estimated number based on 108 samples. To obtain 90% of the total taxa
374 predicted at 3 times the present sample size would require sample size to be doubled to
375 between 70 and 80 samples.

376 We have also plotted the median number of species/taxa found within each depth
377 horizon with respect to ACCZ (Fig. 7). The overriding pattern is of an increase with depth
378 across all ACCZs down to 500 m as well as from south (AZ) to north (SAZ). At depths >500
379 m, species numbers in all but the SAZ dropped somewhat. Further, we performed a stepwise
380 regression on the number of taxa recorded in each sample using depth, ACCZ and month as
381 predictors. This analysis clearly showed that depth alone was the most important factor
382 influencing diversity (r^2 adj = 0.60), with the fit being improved by ACCZ and month
383 ($r^2=0.71$, Mallows CP = 4.0). Overall plankton diversity was highest in the SAZ.

384 Across all depths, the mean number of taxa in the 0-1000 m water column with
385 respect to ACCZ was significantly greater in SAZ stations than elsewhere (Table 8). To
386 illustrate the general trends in the vertical and horizontal distributions of the major taxa, we
387 have plotted the median abundances (ind. m^{-3}) of families and groups with respect to depth
388 and ACCZ (Fig. 8). Within copepod families, trends were apparent both with depth and
389 across ACCZs. The Calanidae (*Calanoides*, *Calanus*, *Neocalanus*), Clausocalanidae

390 (*Clausocalanus*, *Ctenocalanus*, *Microcalanus*) and Oithonidae (*Oithona*, *Paroithona*) are all
391 copepod families that were essentially more abundant within the surface 100 m. In the case of
392 the Calanidae, this was largely due to the presence of younger stages during the summer
393 months. To a greater or lesser extent, calaniids are interzonal migrants and spend a good
394 proportion of the late summer and winter at depth, coming back into the surface layers in
395 spring to reproduce (Andrews, 1966; Atkinson, 1991; Voronina, 1972). The distributional
396 'tail' reflects overwintering/overwintered stages distributed down to 1000 m. Within the
397 Clausocalanidae, *Ctenocalanus* spp. was largely responsible for the increased abundance in
398 the upper 100 m and *Microcalanus pygmaeus* below this. Within the Oithonidae, *Oithona*
399 *similis* was largely responsible for the extremely high abundances in the surface 100 m with
400 the less abundant *O. frigida* and *Paroithona* lying deeper. The Eucalaniids were most
401 abundant in the 100-250 m zones with the two most abundant and widespread interzonal
402 species (*Rhincalanus gigas* and *Subeucalanus longiceps*) spread down the water column with
403 a tendency to be more abundant at northern stations. Ostracoda were widely distributed
404 within the top 500 m but also occurred down to 1000 m. A number of families tended to
405 peak within the 50-100 m and 100-250 m depth horizons e.g. Scolecitrichidae, Euchaetidae,
406 Metridinidae, Aetideidae and Augaptilidae, whereas the Heterorhabdidae, Spinocalanidae,
407 Phaennidae and Lucicutidae were generally more abundant below 500 m. A summary of the
408 distribution of principal copepod families shown in the figure with respect to ACCZ is given
409 in Table 9. Here the number of species identified across all ACCZs is given along with the
410 number found within each defined ACCZ. A total of 102 copepod species were identified in
411 the course of analysis although a number of copepod families such as the Oncaidae, and
412 Spinocalanidae and other non copepod taxa such as the Ostracoda and Siphonophora, were
413 not resolved to species level and will certainly contain more species that are likely to have

414 distinct regional distributions. The overall pattern of copepod distribution indicated that
415 ~10% more copepod species were found within the SAZ compared to the AZ.

416

417 **4. Discussion**

418 *4.1 Sampling methodology*

419 Given that the samples were collected 80 years ago at a time when navigational and
420 oceanographic equipment were less technically advanced than today, we first discuss how
421 this might have impacted upon sample collection and our subsequent interpretation. In a pre-
422 satellite era, ships' navigation and positioning away from land was dependent upon celestial
423 sightings and, between times, dead reckoning. Practical accuracy would therefore have been
424 to the nearest nautical mile, far short of the level of accuracy achieved by the Global
425 Positioning System (GPS) today. However, this does not directly influence our interpretation
426 of data as we are dealing with an oceanographic context determined from the temperature and
427 salinity data rather than from geographical coordinates. Seawater properties were determined
428 from water samples taken with Ekman reversing bottles. The sampling depth of these bottles
429 was estimated from a metering sheave on the hydrographic gantry and checked against depth
430 determined from protected and unprotected thermometers located on some of the bottles. In
431 the *Discovery* station lists (Anonymous, 1942) nominal metered depth can be checked against
432 actual depth for some of the bottles. The two estimates are generally to within a few meters of
433 one another even at depths of 3-4 km indicating that we can be confident of the hydrographic
434 sampling. Temperature is given to 2 decimal places (less than the 4 decimal places generally
435 used today) although the thermometers used were described as being scaled to 0.1°C (Kemp
436 et al., 1929). Given that we used the data to determine whether a front lay between relatively
437 widely spaced stations, this level of accuracy is sufficient for our interpretation.

438 It is more difficult to assess whether the nominal net depth is as indicated, as no
439 independent estimate of depth was apparently undertaken. However, even though wire angles
440 must have deviated from the vertical at times, the accuracy of the deeper water bottle
441 sampling suggests that depth determination during vertical net sampling may also have been
442 reasonably accurate. Wire deviations are expected to be greater in bad weather. Data on wind
443 speed at each of the 41 stations indicates that, at the vast majority (~88%), the Beaufort scale
444 was ≤ 4 at the commencement of operations further suggesting that this may not have unduly
445 influenced net depths.

446 Both bottle sampling and netting were accomplished in a relatively short space of
447 time. Generally, two stations were worked each day commencing at around 9am and 8pm.
448 Netting operations of all types took ~ 4 hours to complete, with an N70V net series to 1000 m
449 generally taking ~1.5-2 hours. A series of Ekman water bottles could be hauled to the surface
450 at ~5.5 min per 1000 m of wire out although would have been deployed at a somewhat slower
451 rate. Thus time on station was not excessive and the zooplankton and hydrographical
452 sampling would be spatially related. It is unlikely that, at any one station, hydrographic
453 sampling was undertaken in one water mass and zooplankton sampling in another.

454 The nets were deployed open to the lower point of the sampled horizon and then
455 hauled vertically upwards before being closed which raises the question of catch
456 contamination, particularly of deeper horizons with surface contaminants. Previous studies
457 have shown that contamination can take place when nets are deployed in this way and
458 plankton are captured as the net jerks upwards when the ship rolls (Grice and Hulsemann
459 1968, Harding 1972). However, *Discovery Investigations* deployed this net over an
460 accumulator spring which would have tended to have minimised this problem (Atkinson
461 1988). Additionally, many of the particularly abundant species that occur throughout the
462 water column such as *Oithona similis* and *Ctenocalanus* spp. and which are more likely to be

463 potential contaminants are known to be seasonal migrants and/or widely distributed within
464 the water column (Atkinson 1988). Although contamination cannot be quantitatively
465 examined, the orders of magnitude lower abundances seen in all of the deeper horizons (Figs
466 2, 3) suggest it is minimal. The isolated occurrences of deeper species in near-surface nets
467 can in all probability be attributed to insufficient washing down of nets between hauls.

468 *4.2 Abundance and biomass*

469 The pattern of mesozooplankton abundance decreasing with depth was similar across
470 all months. Abundance was highly variable in the surface 100 m although greatly exceeded
471 that in the deeper water column. Although somewhat lower overall in the SAZ, both
472 abundance and biomass were not significantly different between ACCZs when viewed over
473 the whole sampled water column. Higher zooplankton abundance near the PF has previously
474 been observed, particularly where phytoplankton biomass was also high (Foxton, 1956,
475 Pakhomov et al., 2000, Dubischar et al., 2002). In our study area, elevated chlorophyll levels
476 are today generally restricted to the near continental shelf region (Fig 1), well south of the
477 position of the PF. It is also likely that, regionally, zooplankton standing stock is lower here
478 than in many other parts of the ACC because of the low primary production seen in the
479 region of 80°W (Fig. 1). Abundance and biomass in the depth horizons below 250 m
480 generally exhibited less variation whether pooled by ACCZ or month.

481 Biomass estimates along 80°W derived from Foxton's (1956) data were somewhat
482 lower (~1.6 to 2.0 g dw m⁻²) than estimates determined by Hopkins (1971) for the Pacific
483 sector, which were 2.67 and 2.58 g dw m⁻² (0-1000 m) for the Antarctic and sub-Antarctic
484 regions respectively. However Foxton excluded gelatinous zooplankton from his
485 determinations and the two estimates may be closer than first appears. Different net
486 performance could also be a factor (see Table 3 Atkinson et al., 2001). Samples for Hopkins'
487 study were taken with a net mesh size of 202 µm. Recent research has shown systematic

488 differences between N70V catches and a 200 μ m bongo net, with the latter catching ~3 times
489 more (by abundance) than the former which translates to ~1.6 times greater biomass (Ward et
490 al., 2012).

491 *4.3 Community analysis*

492 Both MDS ordinations suggested a continuum of stations and species distributions,
493 rather than groups divided by distinct boundaries. This probably reflects the greater number
494 of taxa pooled over a greater depth and the ironing out of seasonal differences when the data
495 were pooled across months. In the deeper ordination (Fig. 5b) community structure was less
496 clear cut, with PFZ stations appearing close in ordination space to AZ and SAZ stations. This
497 may in part be due to manipulating the data by removing seasonal migrants and restricting the
498 analysis to defined depth horizons but also to the lower and more uniform abundances found
499 at depth as well as the rarity of many species. Numerous studies have previously identified
500 plankton communities within the ACC (Errhif et al., 1997; Hunt and Hosie 2005; Pakhomov
501 et al., 2000; Ward et al., 2003). These have generally been based on samples taken from the
502 epipelagic and are very often defined by changes in species abundance rather than by a
503 fundamentally different species composition. In these studies, the PF has generally
504 represented a significant community boundary but, for mesopelagic species, it is less clear cut
505 (Atkinson and Sinclair, 2000). In the deeper horizons where seasonality is less pronounced
506 than in the near-surface, abundances are more uniform and many of the copepod species in
507 particular, whilst widespread, are uncommon. Additionally, previous studies have been
508 largely synoptic rather than using data that have been pooled from September through to
509 March.

510 Mackintosh (1934), Chiba et al. (2001) and Mackey et al. (2012) have all underlined
511 the importance of temperature in determining the distributions of many species of Southern

512 Ocean plankton. Atkinson and Sinclair (2000) have also argued that there is little evidence
513 that the PF forms a biogeographic barrier to the distribution of many species owing to their
514 wide-spread distribution at depth. Our analysis also reflects the wider distribution of many
515 species (Tables 4 and 6). The relatively low values of the R statistic for the AZ/ PFZ and the
516 PF/SAZ pairwise comparisons versus its high values for the AZ/SAZ comparison (Table 3),
517 suggests that the PFZ represents a transition zone. The PFZ has been characterized by the
518 presence of a mixture of sub-Antarctic, sub-tropical and Antarctic species and as such
519 represents a biogeographic ecotone-type community (Pakhomov et al., 2000). There is also
520 no doubt that many species do have distributions that are relatively unaffected by the
521 presence of the PF. Many of the horizontal gradients at ACCZ boundaries are weak relative
522 to vertical gradients and are too weak to limit species distributions which tend to form a
523 continuum, characterised by core regions and regions of expatriation (Angel 1997).
524 Nevertheless, many other species exhibit step changes in abundance in passing from one side
525 of a front to the other and watermass preferences are marked (Boltovskoy et al., 1999). In the
526 wider South Atlantic, Boltovskoy et al. (1999) plotted species distributional boundaries
527 against latitude and found that by far the highest number occurred in the transition zone
528 between the subtropical and sub-Antarctic around 30-40°S. This warm-cold water transition
529 was stronger than the second highest found in the vicinity of the PF.

530 *4.4 Seasonality*

531 Despite pronounced seasonality in the near-surface Antarctic pelagial (Clarke 1988;
532 Smetacek et al., 1990) we found no indirect evidence that this was also apparent in the
533 mesopelagic. Abundance and biomass varied little below 250 m (Table 2) and testing by
534 ACCZ only 5 of 40 taxa/species seasonal abundance comparisons were significantly
535 different, suggesting that seasonal differences were not widespread within the 500-1000 m
536 horizons. In many oceans, seasonal flux of organic material to the deep-sea has been detected

537 at many thousands of metres depth (Billett et al., 1983; Asper et al., 1992). In the North
538 Atlantic, Koppelman and Weikert (1999) found evidence that, below 1000 m, many
539 species/taxa had summer:spring ratios of >1, suggesting a reproductive response to the spring
540 bloom, particularly in the upper bathypelagic zone (1050-2250 m). This increase was
541 pronounced among calanoid copepods and in particular the Metridinidae which increased by
542 a factor of 13.5 between spring and summer. In regions where blooms are largely absent, it
543 was suggested that such responses would not be detected (Koppelman and Weikert, 1999).
544 They cite summer biomass profiles from the Madeira Abyssal Plain (Roe, 1988), where
545 spring blooms do not occur, as being similar to pre-bloom spring profiles from the
546 BIOTRANS site in the temperate northeast Atlantic where their study was carried out.

547 In this part of the Southern Ocean, levels and periodicity of primary production may
548 also be important. Our contemporary composite of ocean colour (Fig. 1) shows that highest
549 levels of chlorophyll are generally found south of the transect locations and may be
550 associated with the retreating ice-edge. To the north of this, chlorophyll is uniformly low
551 ($<0.5 \text{ mg l}^{-1}$). Production patterns have probably not changed dramatically over the years
552 because the transect lies upstream of the main sources of iron input into the ACC in this
553 sector, namely shelf deposits from the Antarctic Peninsula and topographically induced
554 upwelling (Park et al., 2010). Data on the presence of phytoplankton in the 80°W samples are
555 given by Foxton (1956) (Table 11c). December and March samples from the AZ contained
556 phytoplankton in the near-surface layers which implies that 'summer' stations in the AZ in
557 1933/34 were only just experiencing a bloom, suggesting that deep-water fauna would in any
558 case have had no time to respond to this burst of production. A reproductive response later in
559 the year (early winter onwards) may explain why spring abundance of some species was
560 greater than during summer. Interannual variability also may be partly responsible insofar as

561 the summer grouping (Dec. 1933 and Mar. 1934) precede rather than succeed the spring
562 grouping (Sept, Oct, Nov 1934) and so effectively different cohorts were being assessed.

563 4.5 Diversity

564 Within this study a number of taxonomic categories were unresolved into species. Our
565 ‘lumping’ of species manifested itself variously at the level of phylum, e.g. Chaetognatha,
566 Ctenophora; class or order e.g. Ostracoda, Siphonophora, Appendicularia; family or genus
567 e.g. Lucicutiid copepodites, *Oncaea* spp. etc. Of the taxa identified, copepods were the most
568 highly resolved to species level. Reasons for this variously included the state of preservation
569 of some elements of the samples, expedience in wishing to analyse a large number of samples
570 in a timely manner and levels of taxonomic expertise. Nonetheless some clear patterns
571 emerged. Rarefaction analysis highlighted the relationship of increasing diversity with depth
572 when species presence/absence data from all ACCZs were pooled with respect to depth (Fig
573 6). This was also to an extent mirrored in the clustering of station data with respect to depth
574 (Fig 4) which effectively partitioned the water column across all ACCZs into 3 distinct
575 horizons. The rarefaction curves from the two deepest horizons corresponding to the lower
576 mesopelagic (*sensu* Fig. 4) indicate a greater diversity and have a steeper slope than the
577 shallower depth-horizons indicating a different rate of species accumulation. It should be
578 borne in mind that each deep sample represents 5 times the amount of water swept per sample
579 in the upper two horizons in the top 100 m. However, the slope of the species accumulation
580 curve was different and even with an extrapolated sample number (equivalent to a greater
581 volume of water swept) diversity never approaches that of the deeper horizons. Modern net
582 sampling of the deep water column is usually undertaken with larger nets than the N70, using
583 oblique rather than vertical hauls and generally sampling a greater volume of water. This
584 would undoubtedly influence the rate of accumulation of species per haul but the relative
585 pattern between the different horizons is unlikely to change dramatically.

586 It is difficult to summarise patterns of diversity across such a heterogenous group as
587 plankton. Some groups are taxonomically better investigated than others, using both
588 traditional morphology and modern molecular techniques such as genetic bar coding (Bucklin
589 et al. 2011) but, at a wider scale, the polar and sub-polar biomes are less rich in species across
590 many groups than their tropical and subtropical counterparts (Angel et al., 2007; Boltovskoy
591 et al., 1999; Woodd-Walker et al., 2002). In near-surface samples, Woodd-Walker et al.
592 (2002) have demonstrated the large-scale spatial variation in taxonomic richness of copepod
593 genera across the whole Atlantic Ocean. Higher diversity was apparent in the tropics and
594 sample evenness and diversity also reduced dramatically around 40°S and towards the poles.
595 This has been attributed to the former possessing a relatively stable environment where
596 seasonal changes are minimal, allowing for a largely retentive system in which primary and
597 secondary producers are closely coupled and diversity is characteristically high (Conover,
598 1979; Longhurst and Pauly, 1987). Polewards, seasonality increases and production and
599 consumption become increasingly uncoupled and diversity is lower. Increasing diversity
600 with depth may therefore reflect the relative stability of the lower part of the mesopelagic in
601 contrast to the seasonal breakdown of the thermocline and the winter overturning of the upper
602 water column (Woodd-Walker et al., 2002). At a smaller spatial scale our analysis suggested
603 that slightly more copepod species were present in the SAZ than in either the AZ or PFZ.
604 This was in large measure through higher species numbers within the families Aetideidae and
605 Augaptilidae being found in the SAZ (Table 9). However, based on the wider literature,
606 Razouls et al. (2005-2012) record 47 and 39 members of the Aetideidae and 27 and 20
607 members of the Augaptilidae in the AZ and SAZ respectively. Our samples only recovered a
608 relatively small proportion of these totals which reflects sampling effort, regional bias and
609 overall rarity of many of the deeper dwelling species (see E-Table). Large-scale physics in
610 the form of mesoscale eddies (Nowlin and Klinck 1986) may also influence diversity by

611 introducing species polewards or in some cases equatorwards across the PF. Based on the
612 wider literature extending to the entire Southern Ocean, Razouls et al. (2005-2013) estimate
613 that the number of copepod species occurring in all Antarctic waters totals 295 compared to
614 275 reported for the sub-Antarctic. They have also suggested that ~80% of copepod species
615 observed in the sub-Antarctic are immigrants and originate from sub-tropical and temperate
616 zones. It is possible that these form 'pseudo-populations' (Razouls et al., 2005-2013) and
617 many of these may already be at the limits of their distributional ranges. Other taxonomic
618 groups such as Amphipoda, Ostracoda and Salpidae, appear to have a greater number of
619 species in the SAZ whereas others such as euphausiids, chaetognaths and some radiolarians
620 appear more evenly distributed (Boltovskoy et al., 1999).

621 *4.6 Long-term change*

622 Profound physical changes have occurred in the Southern Ocean in the intervening 80
623 or so years since *Discovery Investigations*. At the WAP, close to the 80°W transect, climate
624 change has been rapid. Atmospheric temperatures have risen by ~3°C since 1951 (Vaughan et
625 al. 2003) and extensive glacial retreat has occurred (Cook et al., 2005). The surrounding
626 ocean temperature has also increased by ~1°C in summer (Meredith and King 2005), and sea-
627 ice duration and cover has declined in this region (Cavalieri and Parkinson 2008). The
628 potential and actual impacts of such changes are being recognised in various parts of the
629 marine ecosystem, (eg Atkinson et al., 2004; Ducklow et al., 2007; Clarke et al., 2007;
630 Forcada et al 2006; Schofield et al., 2010) and are generally complex.

631 The transect along 80°W lies north and west of the WAP and is largely north of the
632 seasonal sea-ice zone. Ocean warming throughout the region has not been uniform with depth
633 and, along with decreased salinity, is greater in near-surface waters (Meredith and King,
634 2005; Böning et al., 2008). Because of warming and freshening, Böning et al. (2008) estimate
635 that density surfaces between the PF and SAF at 800-1000 m have been displaced southwards

636 50–80 km in the last 40 years and perhaps 2-3 times this distance at the surface. This may
637 imply a southwards displacement of frontal zones but this is likely to be difficult to detect
638 against the natural background variation in the position of the ACC fronts (Moore et al.,
639 1999; Venables et al., 2012). Plankton distributions frequently show strong temperature
640 dependence in the Southern Ocean (Mackintosh, 1934; Chiba et al., 2001; Mackey et al.,
641 2012). Simplistically, as a consequence of warming, we might expect a southwards
642 penetration of some ‘warmer’ water species and a contraction in the range of cold water
643 species as predicted by Mackey et al. (2012) for macrozooplankton. This seems plausible as
644 Ward et al. (2004, 2006) have shown that, in extensive surveys of the Scotia Sea,
645 mesoplankton, particularly copepods, were much less abundant in ice influenced waters. A
646 polewards movement of the ice-edge may therefore progressively result in increased
647 zooplankton diversity and abundance further south. However, many species have wide
648 distributional ranges, occurring in all ACCZs sampled and over wide depth ranges and
649 without the benefit of time-series data change is not going to be easy to detect (Ward et al.,
650 2008, 2012; Mackey et al., 2012). Nonetheless, *Discovery* samples were taken over a large
651 part of the Southern Ocean using the same sampling gear and provide a valuable resource
652 with which to undertake comparisons on a regional basis and with contemporary collections.

653

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659 [collections/invertebrate-collections/historical-marine-collections/nhm-](http://www.nhm.ac.uk/research-curation/collections/our-collections/invertebrate-collections/historical-marine-collections/nhm-collections/discovery/index.html)
660 [collections/discovery/index.html](http://www.nhm.ac.uk/research-curation/collections/our-collections/invertebrate-collections/historical-marine-collections/nhm-collections/discovery/index.html)

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887 Fig. 1. Composite satellite image showing mean chlorophyll *a* during the period September –
888 March 2004–2012, east of Drake Passage. White areas denote missing data due to
889 land/cloud/sea ice cover. Superimposed are *Discovery* station positions coded by month.
890 Shaded grey rectangles represent the limits within which the sub-Antarctic Front (SAF) and
891 Polar Front (PF) were found during these transects. The chlorophyll *a* data used in Fig. 1 are
892 MODIS-Aqua 9 km resolution, level 3 data provided by the NASA Goddard Earth Sciences
893 Data and Information Services Center (Acker and Leptoukh, 2007).

894
895 Fig. 2. Logged Mesozooplankton abundance (ind m⁻³) with respect to depth. Data shown are
896 median plus interquartile range (box) and 10th and 90th percentiles (whiskers). Dots represent
897 outliers. Data have been pooled within depth horizons across ACC Zones.

898

899 Fig. 3. a) Mesozooplankton abundance (ind. m⁻³) versus depth by month. Open symbols
900 represent individual data points, filled circles, median values by depth. Key for depth: 1 = 50-
901 0 m, 2 = 100-50 m, 3 = 250-100 m, 4 = 500-250 m, 5 = 750-500 m, 6=1000-750 m.

902 b) Mesozooplankton abundance (ind.m⁻³) versus ACC zone by depth. Open symbols
903 represent individual data points, filled circles, median values by ACC zone. Key for depth, as
904 for Figure 3. ACC zone abbreviations as Table 1.

905 Fig. 4. Results of nearest neighbour clustering on the Bray-Curtis similarity matrix containing
906 all sample data. Samples have been coded according to depth horizon. We have sliced the
907 cluster dendrogram at 70% similarity. At $\geq 70\%$ similarity, the dendrogram has been
908 collapsed and stations are represented by symbols at the similarity level at which they first
909 became statistically indistinguishable. Red lines at $< 70\%$ similarity extend from the point at
910 which the grouping became statistically indistinguishable from one another. Black lines
911 extending to the x axis are statistically dissimilar to other stations.

912
913 Fig. 5a) MDS ordination carried out on the 0-1000 m data matrix. Lines encircling stations
914 represent 73% similarity.

915 5b) MDS ordination carried out on the 500-1000 m data matrix with seasonal migrants
916 omitted (see below). Lines encircling stations represent 72% similarity.

917 Open circles = AZ stations, filled triangles PFZ stations and open squares SAZ stations.

918 Taxa omitted from 500-1000m data matrix were *Calanoides acutus*, *Calanus simillimus*,
919 *Neocalanus tonsus*, *Rhincalanus gigas* plus nauplii, *Subeucalanus longiceps*, *Ctenocalanus*
920 spp. *Clausocalanus laticeps*, *Clausocalanus brevipes*, *Ctenocalanus/Clausocalanus*

921 copepodites, *Scolecithricella minor*, *Oithona similis*, *Limacina helicina*, *Limacina retroversa*
922 and Appendicularia spp.

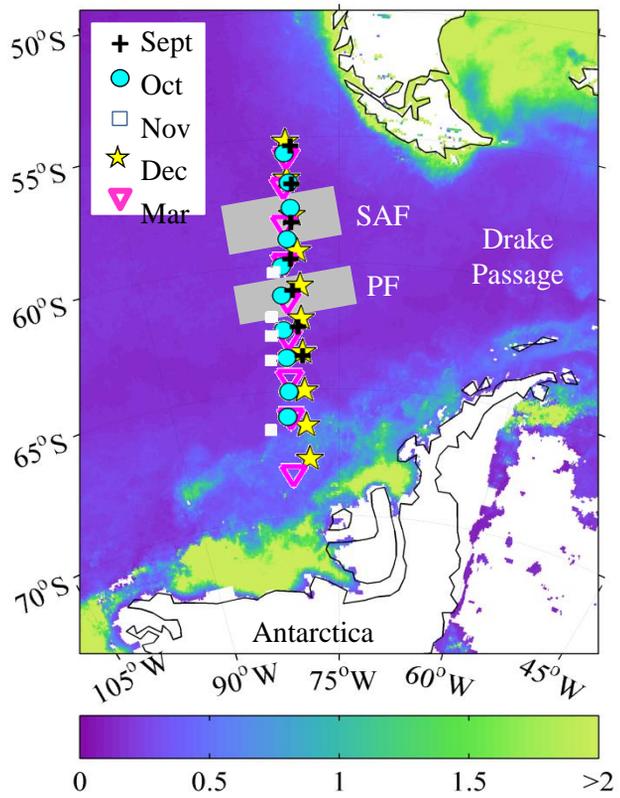
923 Fig. 6. Rarefaction curves based on a presence/absence species by depth matrix. Between 35-
924 37 samples were used from each depth horizon to construct the curves using the “Moa tau”
925 method (bold lines). A 3 parameter power function fitted to each rarefaction curve was then
926 extrapolated to 3 times the sample size (108 samples) following Colwell et al., (2004; fine
927 lines).

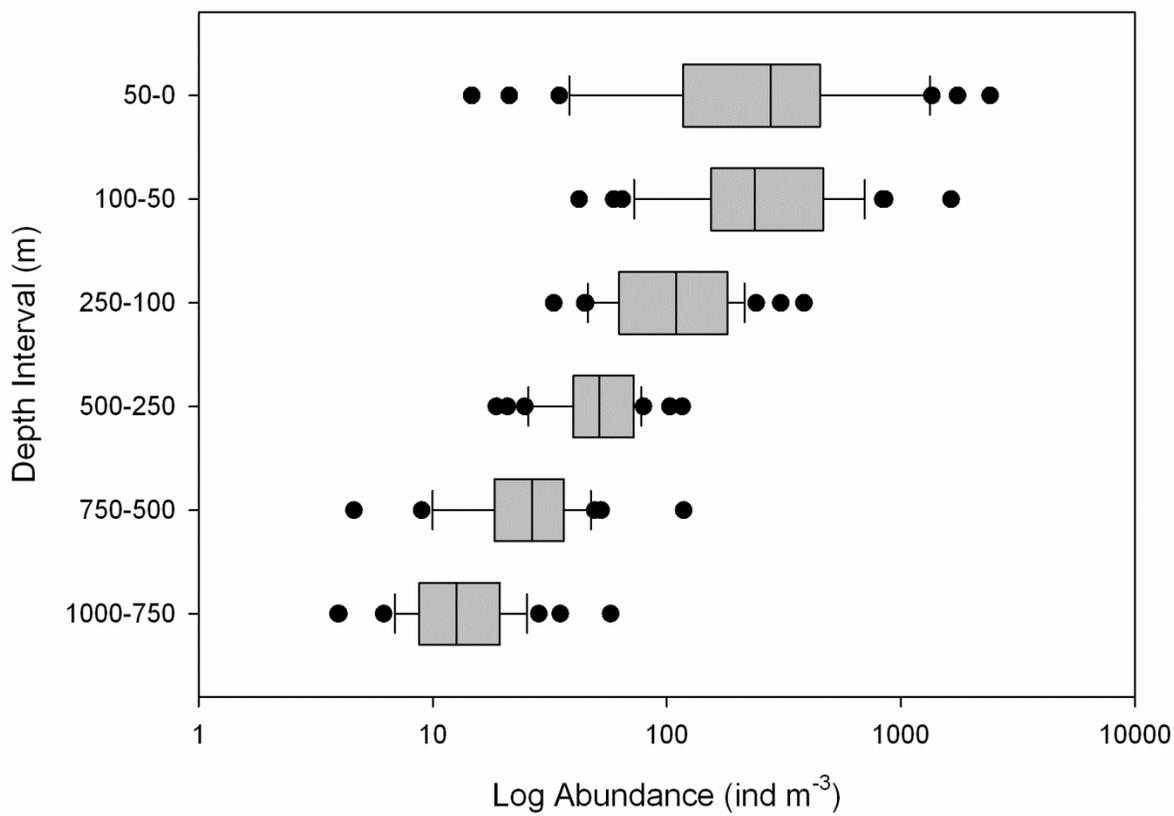
928 Fig. 7. Boxplot of median number of species/taxa in each depth horizon with respect to ACC
929 zone. Data shown are median plus interquartile range (box) and max and min (whiskers) * =
930 outlier. Depth coding as for Fig. 3.

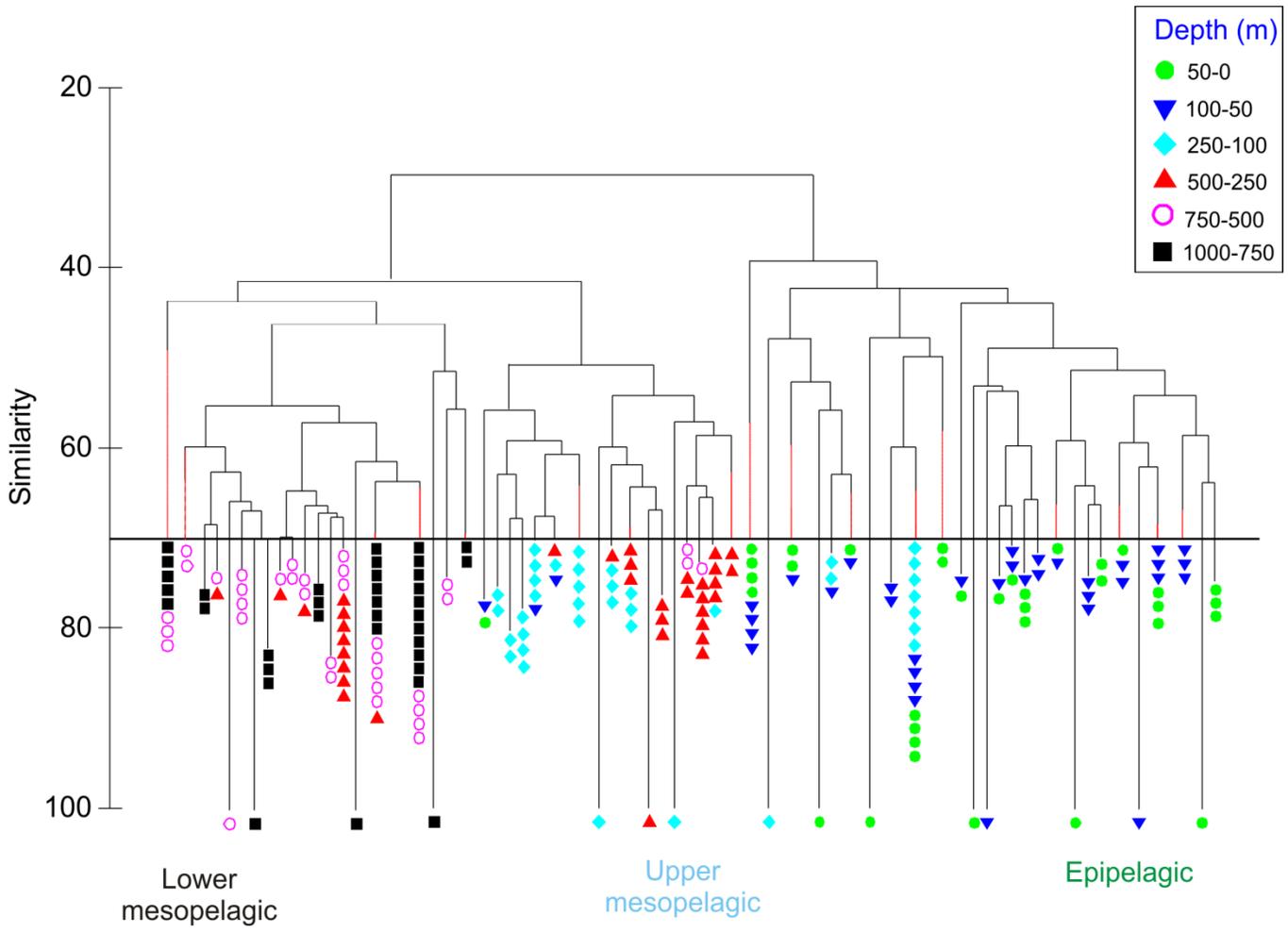
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932 Fig. 8. Median abundance (ind. m⁻³) of major copepod families and other taxa by ACC zone
933 and depth. Data shown are median plus interquartile range (box) and max and min
934 (whiskers), * = outlier. Depth coding as for Fig. 3.

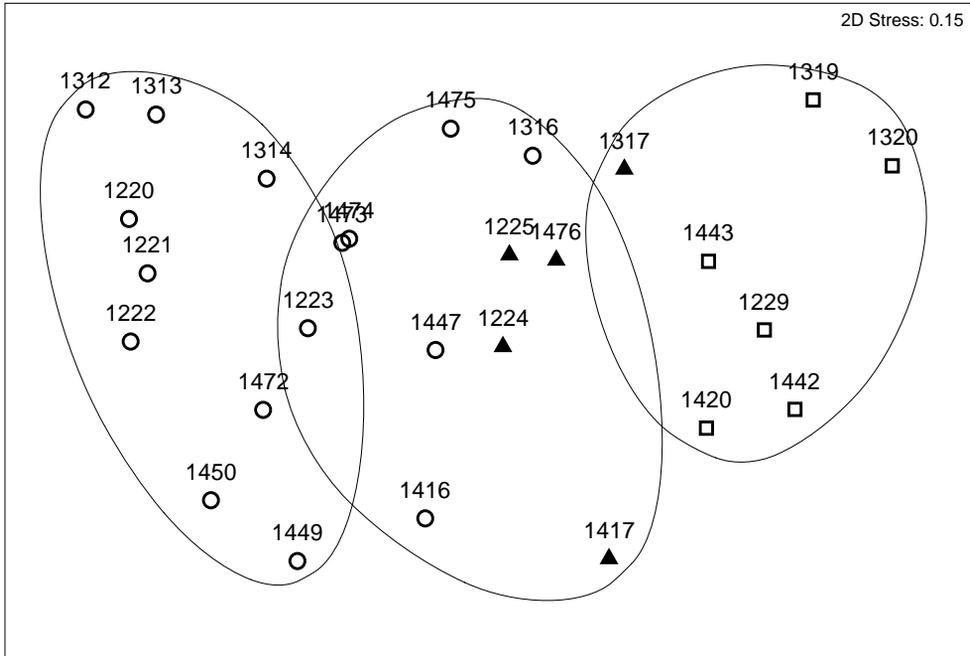
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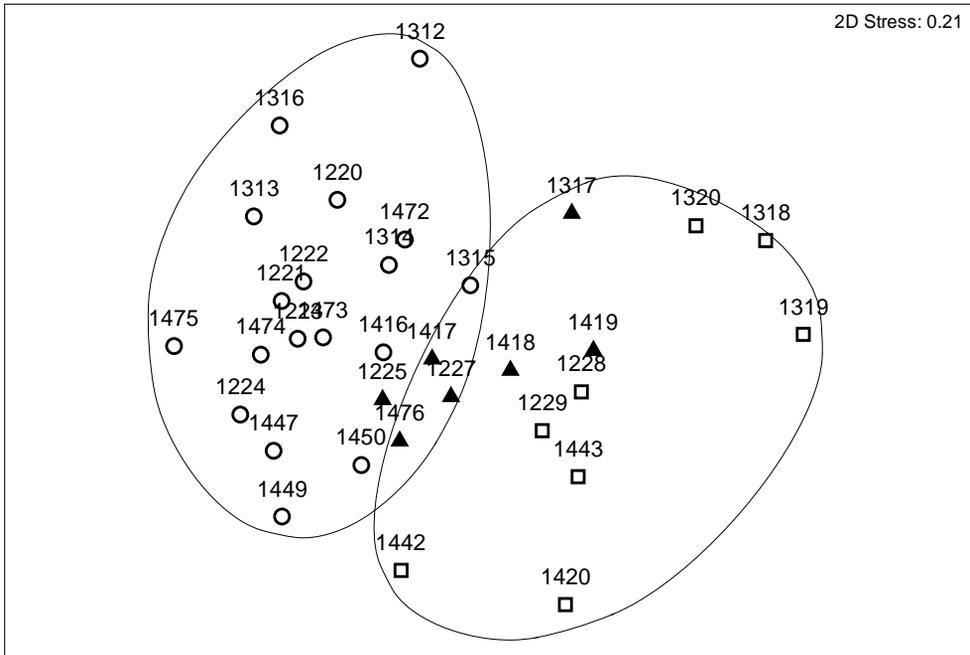


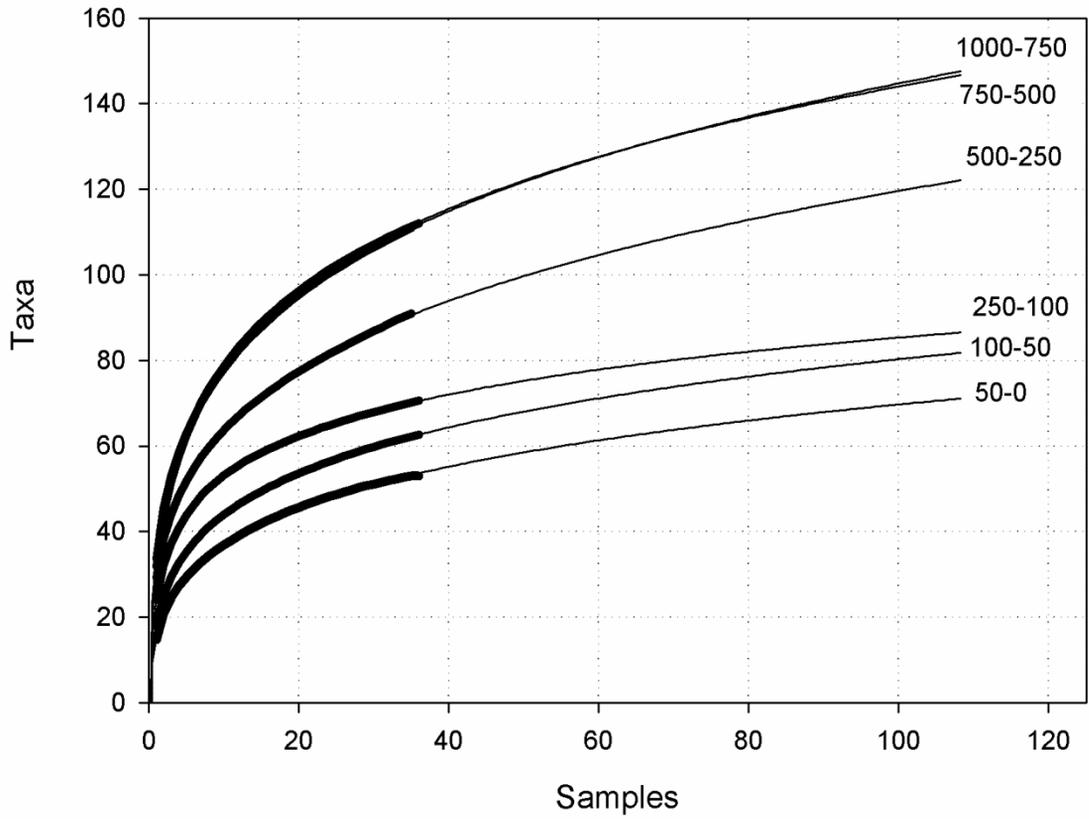


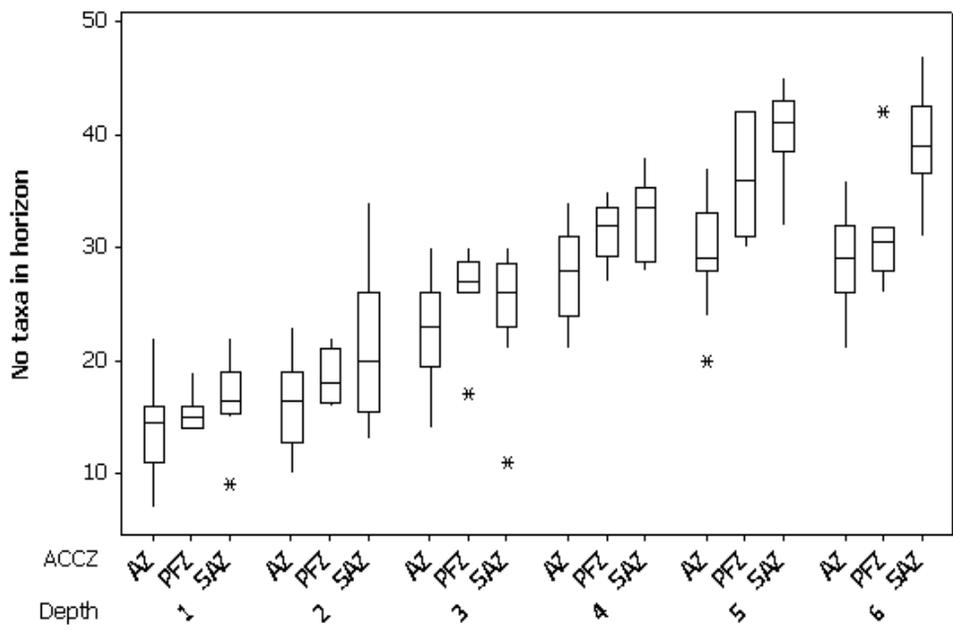
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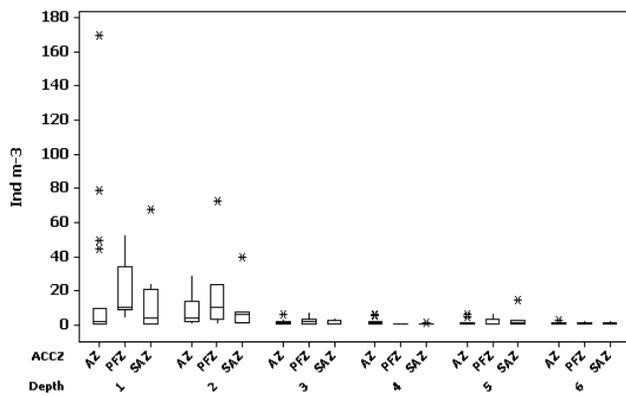
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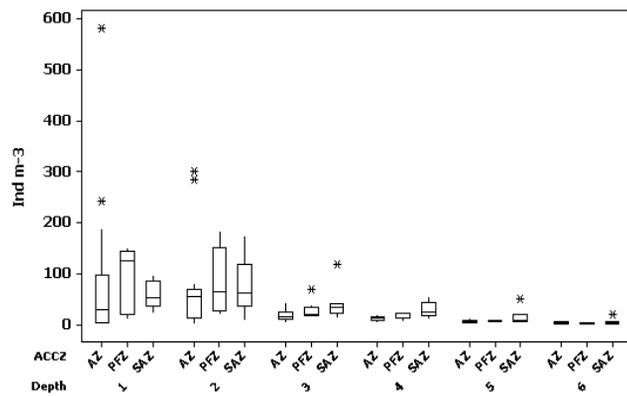




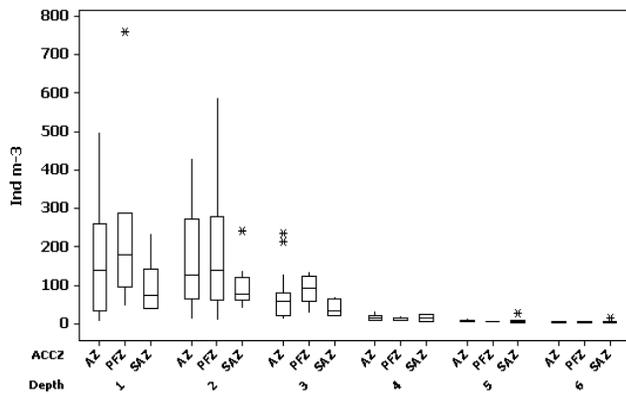
Calanidae



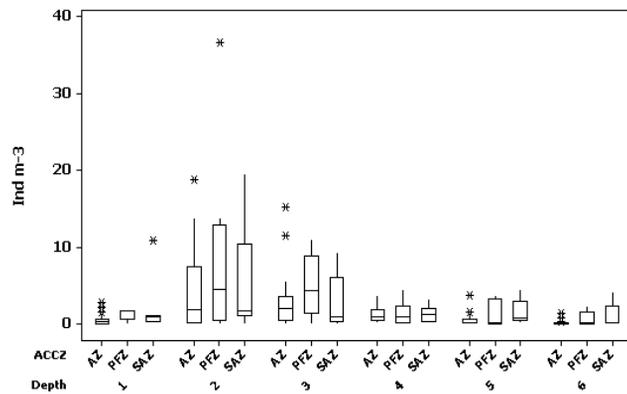
Clausocalanidae



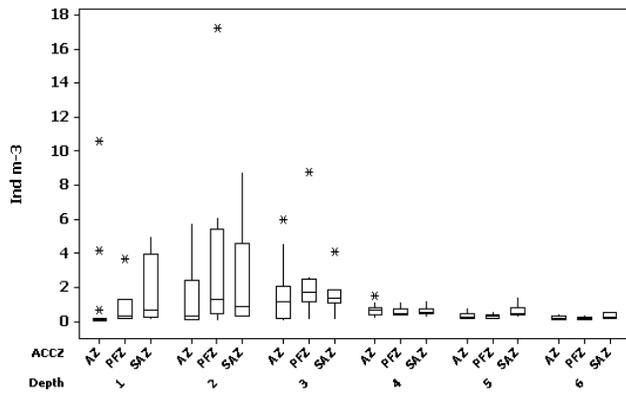
Oithonidae



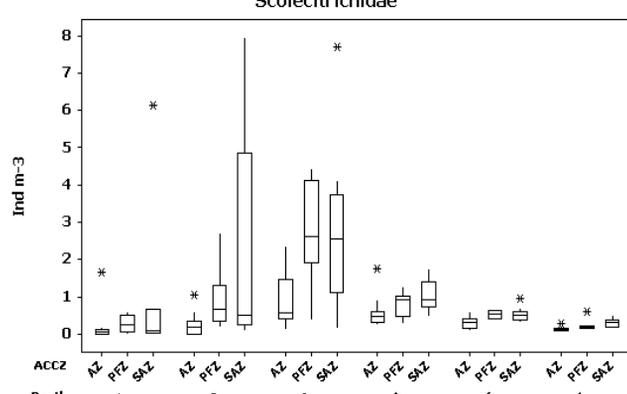
Eucalanidae



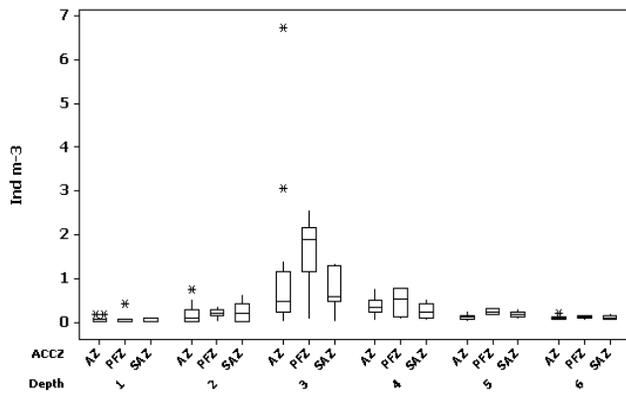
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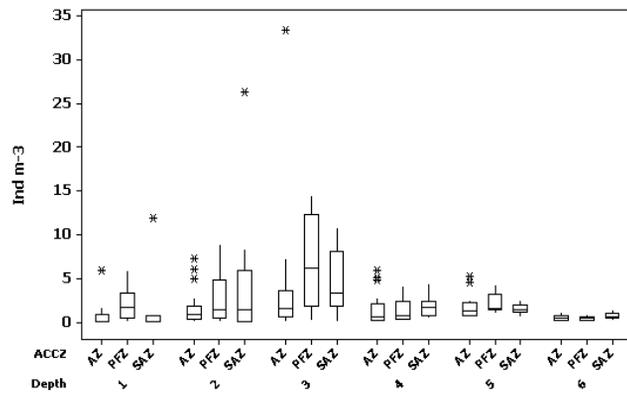
Scolecitrichidae



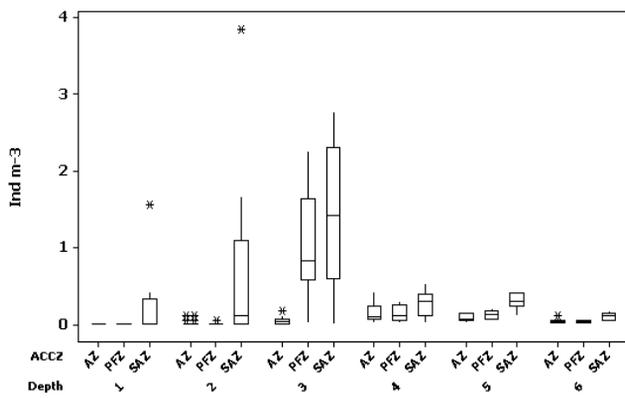
Euchaetidae



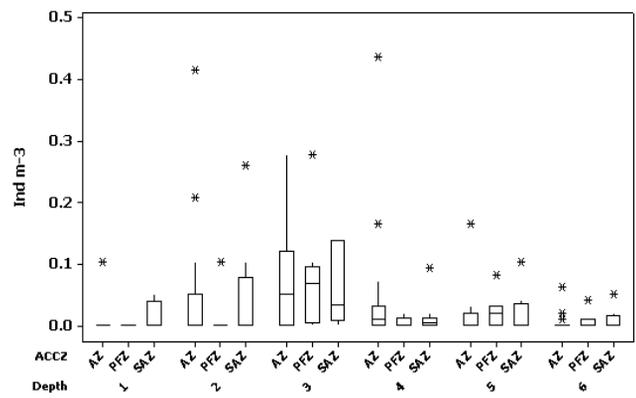
Metridinidae



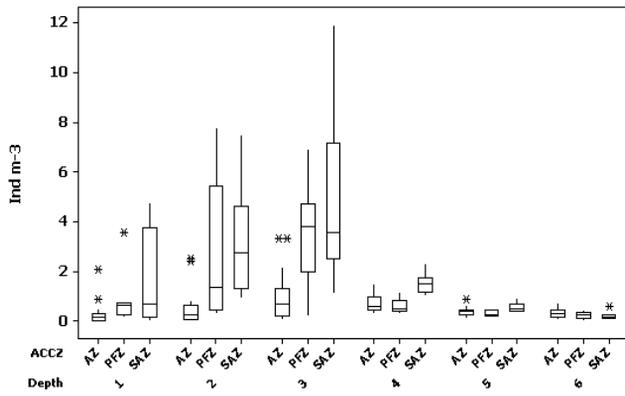
Aetidiidae



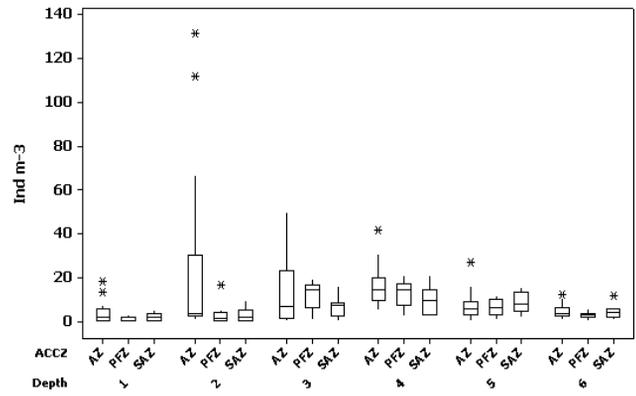
Augaptilidae



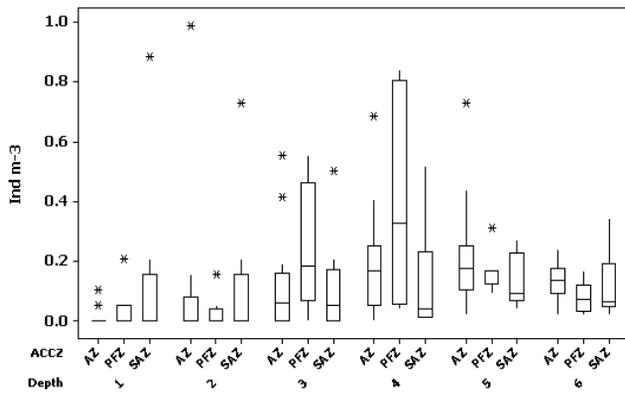
Ostracoda



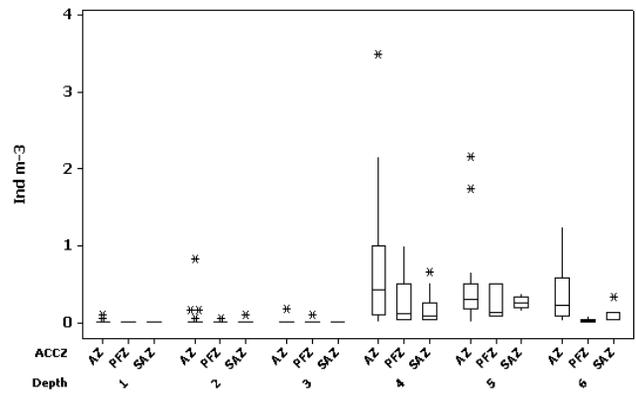
Oncaeidae



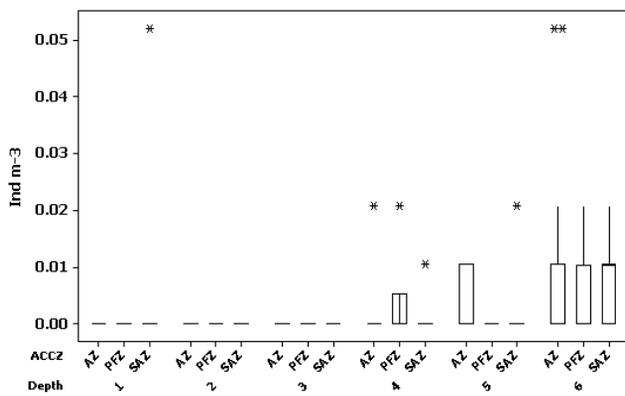
Heterorhabdidae



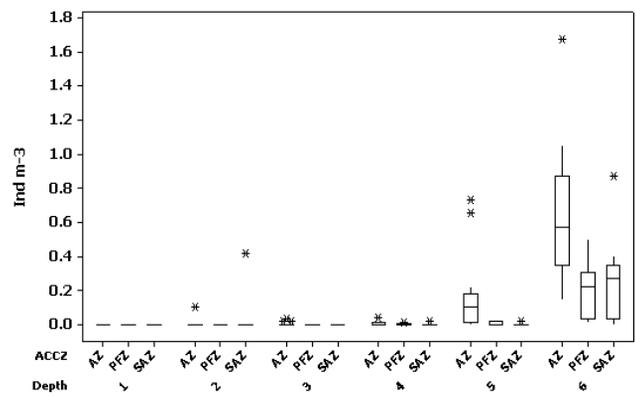
Spinocalanidae



Phaennidae



Lucicutidae



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937 Table 1. *Discovery* stations sampled along 80°W by month with respect to ACC Zones.

938 Stations emboldened were originally classified as Antarctic stations by Foxton (1956) and the

939 remainder as sub-Antarctic. Antarctic Circumpolar Current Zones (ACCZ) defined according
940 to potential temperature, salinity and potential density sections and potential temperature-

941 salinity curves (see text). Stations with an asterisk are those where one or more samples are

942 missing or the contents had been spilt.

943

	ACCZ		
	Antarctic Zone (AZ)	Polar Frontal Zone (PFZ)	sub-Antarctic Zone (SAZ)
December (1933)	1220, 1221, 1222, 1223, 1224	1225, 1226*, 1227*	1228*, 1229
March (1934)	1312, 1313, 1314, 1315*, 1316	1317	1318*, 1319, 1320
September (1934)	1415*, 1416,	1417, 1418*, 1419*	1420, 1421*
October (1934)	1447, 1449, 1450	1446*, 1444*	1441*, 1442, 1443,
November (1934)	1472, 1473, 1474, 1475	1476	

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949 Table 2: Median plankton abundance (ind. m^{-3} (Q1-Q3)) and biomass (wet mass $mg\ m^{-3}$ (Q1-
 950 Q3)). Biomass derived from settled volume (cm^3) assuming $1\ cm^3 = 1\ g$ wet mass. See
 951 original data in Foxton (1956) Table 11c. . The formula $\pi \times r^2 \times h$ has been used to estimate
 952 abundance and volume per m^2 and thence m^3 . ACC Zone definitions and abbreviations as
 953 Table 1, (n) = no. of samples within each ACC Zone.

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	Depth (m)	AZ (111)	PFZ (50)	SAZ (54)
Abundance	50-0	306 (43-700)	294(194-464)	187 (127-299)
	100-50	294 (177-565)	211 (107-552)	180 (157-239)
	250-100	92 (55-153)	182 (122-195)	93 (66-147)
	500-250	44 (39-72)	54 (35-67)	56 (40-83)
	750-500	20 (13-36)	26 (20-29)	30 (23-46)
	1000-750	13 (8-21)	10 (8-18)	14 (7-24)
Biomass	50-0	21 (5-47)	44 (29-51)	36 (13-60)
	100-50	52 (10-88)	47 (23-153)	36 (23-59)
	250-100	21 (10-40)	42 (16-49)	26 (14-41)
	500-250	19 (12-22)	14 (11-17)	12 (16-23)
	750-500	12 (8-19)	14 (9-17)	14 (11-19)
	1000-750	7 (5-10)	10 (8-13)	9 (7-10)

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963 Table 3: Analysis of Similarities (ANOSIM), testing how well station grouping reflects
 964 ACCZ definitions (see text). ACC Zone definition and abbreviations as Table 1. * =
 965 significant difference. No. of permutations = 999.

966

Data matrix	R statistic	Significance Level (<i>p</i>)	Watermass Comparison
<hr/>			
0-1000 m			
Differences between ACCZs	0.32	0.008	AZ/PFZ*
	0.87	0.001	AZ/SAZ*
	0.39	0.010	PFZ/SAZ*
500-1000 m (less seasonal migrants)			
Differences between ACCZs	0.25	0.016	AZ/PFZ*
	0.74	0.008	AZ/SAZ*
	0.12	0.130	PFZ/SAZ
<hr/>			

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976 Table 4: Abundance (mean ind. m⁻², 0-1000 m (±SD)) of the first ten species/taxa in the analysis shown to be contributing
 977 most to within group similarity and between group dissimilarity across all 3 groups of stations (no. of stations). Taxa ranked
 978 in terms of overall abundance across all groups. ACC Zone definitions and abbreviations as Table 1.
 979

Species/taxa	AZ(16)	PFZ(5)	SAZ(6)
<i>Oithona similis</i>	22399 (11274)	32167 (15881)	11616 (6456)
<i>Ctenocalanus</i> spp.	8240 (5557)	17167 (15859)	12565 (7039)
Pteropoda	15094 (43509)	359 (753)	62 (128)
<i>Oncaea</i> spp.	10827 (6623)	7742 (6091)	6415 (3691)
<i>Oithona frigida</i>	6300 (2412)	5720 (2774)	4128 (1779)
<i>Microcalanus pygmaeus</i>	3970 (1601)	4587 (820)	7562 (3594)
Calanoid nauplii	3315 (3511)	3480 (3891)	1650 (1075)
<i>Metridia lucens</i>	709 (616)	2598 (2258)	1701 (1165)
<i>Paroithona</i> sp.	31 (39)	858 (815)	4492 (4889)
<i>Rhincalanus gigas</i>	1024 (656)	1784 (499)	864 (952)
<i>Calanus similimus</i>	740 (2413)	1667(1165)	1664 (1736)
<i>Calanoides acutus</i>	1342 (1051)	610 (1143)	9 (10)
<i>Ctenocalanus/ Clausocalanus</i> copepodites	447 (761)	486(469)	2257 (1949)
Ostracoda	489 (201)	998 (335)	1389 (545)
Chaetognatha spp.	619 (378)	855 (753)	847 (451)
Appendicularians	881 (1803)	236 (266)	533 (1088)
Copepod spp. (Unidentified copepodites)	742 (828)	332 (256)	774 (399)
<i>Rhincalanus gigas</i> nauplii	783 (1383)	347 (455)	31 (76)
<i>Clausocalanus laticeps</i>	34 (73)	266 (263)	859 (1277)
<i>Subeucalanus longiceps</i>	19 (18)	147 (112)	908 (445)

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Table 5: Mean abundance ind.m⁻² (\pm SD) within the depth horizons indicated with respect to ACC Zone grouping (see Table 3). ACC Zone definition and abbreviations as Table 1. (n) number of stations within respective ACC Zones.

Depth	ACC Zone Grouping		
	AZ (16)	PFZ (5)	SAZ (6)
0-1000 m	8.18 x10 ⁴ (5.17 x10 ⁴)	8.50 x10 ⁴ (4.40 x10 ⁴)	6.47 x10 ⁴ (3.08 x10 ⁴)
	AZ (18)		SAZ /PFZ(15)
500-1000 m	0.80 x10 ⁴ (0.44 x10 ⁴)		0.86 x10 ⁴ (0.44 x10 ⁴)

Table 6: Abundance (mean ind. m⁻², 500-1000 m (±SD)) of the first ten species/taxa in the analysis shown to be contributing most to within group similarity and between group dissimilarity across all 3 groups of stations (no. of stations). Taxa ranked in terms of overall abundance across all groups. ACC Zone definitions and abbreviations as Table 1.

Species/taxa	AZ (18)	PFZ & SAZ (15)
<i>Oncaea</i> spp.	2989 (2268)	2874 (1501)
<i>Oithona frigida</i>	1983 (963)	1452 (656)
<i>Microcalanus pygmaeus</i>	1238 (1225)	2150 (1831)
<i>Metridia lucens</i>	243 (353)	372 (259)
<i>Metridia curticauda</i>	223 (78)	152 (66)
Calanoid nauplii	194 (151)	170 (84)
Ostracoda spp.	178 (80)	165 (62)
<i>Spinocalanus</i> spp.	225 (215)	72 (43)
Chaetognatha spp.	115 (66)	171 (118)
<i>Lucicutia ovalis</i>	200 (96)	54 (62)
<i>Paroithona</i> sp.	5 (15)	241 (551)
Scolecithriciid copepodites	56 (30)	83 (50)
<i>Pleuromamma robusta</i>	43 (72)	85 (45)
<i>Mormonilla</i> sp.	20 (28)	83 (74)
Siphonophora	12 (13)	51 (53)
<i>Clione antarctica</i>	15 (21)	30 (37)
<i>Paraeuchaeta biloba</i>	8 (14)	22 (13)
Euphausiid nauplii	14 (25)	14 (27)
<i>Thysanoëssa</i> spp.	6 (21)	11 (9)
<i>Euaetideus australis</i>	0 (0)	12 (15)

Table 7: Mean (\pm SD) and median (Q1-Q3) abundance (ind.m⁻², per 250 m haul in the 500-1000m depth horizon) of the 20 most abundant species/taxa within the 500-1000 m horizons. Data represent spring (Sept/Oct /Nov) and summer (Dec/March). Only *Spinocalanus* spp. showed a significantly different seasonal abundance (Anova $F_{1,69}=5.74$, $p = 0.019$).

Species/taxa	Spring mean (ind. m ⁻² , per 250 m haul in the 500-1000 m depth range)	Summer mean (ind. m ⁻² , per 250 m haul in the 500-1000 m depth range)	Spring median (ind. m ⁻² , per 250 m haul in the 500-1000 m depth range)	Summer median (ind. m ⁻² , per 250 m haul in the 500-1000 m depth range)
<i>Oncaea</i> spp.	1199 (879)	1674 (1381)	915 (585-1681)	1122 (686-2359)
<i>Oithona frigida</i>	835 (835)	850 (454)	717 (364-1065)	727 (587-1086)
<i>Microcalanus pygmaeus</i>	710 (612)	898 (1317)	546 (177-1081)	447 (125-1122)
<i>Metridia lucens</i>	122 (207)	165 (252)	30(8-131)	94 (27-194)
<i>Metridia curticauda</i>	103 (61)	90 (72)	87 (64-126)	83 (23-152)
Calanoid nauplii	89 (87)	86 (77)	83 (42-117)	83 (16-120)
Ostracoda	91(46)	79 (53)	87 (61-110)	73 (35-118)
<i>Spinocalanus</i> spp.	47*(61)	100 (115)	23 (10-62)	73 (31-125)
Chaetognatha	68 (43)	70 (70)	64 (31-89)	42 (25-104)
<i>Lucicutia ovalis</i>	67 (96)	62 (70)	26 (0-102)	34 (3-112)
<i>Paroithona</i> sp.	12 (21)	91 (323)	0 (0-21)	0 (0-31)
<i>Heterorhabdus</i> spp.	36 (25)	39 (32)	32 (18-45)	34 (16-51)
Scolecithriciid copepodites	34 (29)	35 (33)	26 (16-52)	21(10-56)
<i>Pleuromamma robusta</i>	35 (60)	26 (41)	16 (0-45)	5 (0-38)
<i>Mormonilla</i> sp.	28 (39)	23 (35)	10 (0-42)	10 (0-31)
<i>Gaetanus tenuispinus</i>	16 (18)	18 (24)	9 (3-19)	10 (3-27)
Siphonophora	9 (13)	19 (34)	3 (0-12)	8 (0-19)
<i>Pareuchaeta antarctica</i>	14 (13)	9 (14)	10 (5-21)	5 (1-10)
Scaphocalaniid copepodites	9 (13)	13 (15)	5 (2-10)	8 (3-19)
<i>Scolecithricella dentipes</i>	13 (11)	8 (8)	9 (2-23)	8 (3-13)

Table 8: Mean (\pm SD) number of species/taxa within the 0-1000 m water column with respect to ACC Zone. Of the comparisons carried out among ACC Zones using all taxa and copepods only that between AZ and SAZ was significant; ANOVA $F_{1, 20 \text{ df}} = 14.41$, $p = 0.001$ for all taxa and $F_{1, 20 \text{ df}} = 16.58$, $p = 0.001$ for copepods alone. ACC Zone definition and abbreviations as Table 1.

Watermass (no. stns)	No. taxonomic categories	No. copepod categories
AZ (17)	60.4 \pm 6.1	46.9 \pm 3.4
PFZ (4)	61.5 \pm 6.2	47.3 \pm 4.7
SAZ (6)	73.7 \pm 9.6	57.0 \pm 8.5

Table 9. Number of species of copepods identified within each ACC Zone with respect to Order and Family. Family (n) = no. of species identified within family across all groupings. ACC Zone definition and abbreviations as Table 1. ACCZ (n) = no. samples analysed.

Order	Family	AZ (117)	PFZ 44)	SAZ (54)
Calanoida				
	Calanidae (4)	4	3	4
	Eucalanidae(5)	2	4	4
	Spinocalanidae(2)	2	1	1
	Clausocalanidae (4)	4	4	4
	Tharybidae (1)	1	1	1
	Stephidae (1)	1	0	0
	Aetideidae (16)	8	8	15
	Euchaetidae (7)	5	4	6
	Phaennidae (5)	3	4	3
	Scolecitrichidae (13)	11	8	8
	Arietellidae (1)	1	0	1
	Augaptilidae (11)	4	5	9
	Heterorhabdidae (6)	4	3	6
	Lucicutiidae (6)	6	2	3
	Metridinidae (6)	5	6	5
	Phyllopodidae (1)	1	1	1
	Candaciidae (1)	1	1	1
	Bathypontidae (1)	1	1	1
Mormonilloida				
	Mormonillidae (1)	1	1	1
Cyclopoida				
	Oithoniidae (4)	4	3	4
Harpacticoida				
	Ectinosomatidae (1)	1	1	1
	Aegisthidae (1)	0	0	1
Poecilostomatoida				
	Oncaeidae (2)	2	2	2
Siphonostomatoida				
	Ratanidae (1)	0	0	1
	Totals	72	63	83

1

2 Highlights

3 1. Plankton samples from 1930's *Discovery Investigations* in the Southern Ocean analysed

4 2. Depth was the strongest factor separating samples

5 3. Mean zooplankton abundance in the epipelagic was ~ 25 times greater than at 1000 m

6 4. Seasonal signals across all data became less distinct with depth.

7 5. Rarefaction analysis revealed that depth was a major influence on diversity.

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