

**Seasonal and diel variations in the vertical distribution,  
composition, abundance and biomass of zooplankton  
in a deep Chilean Patagonian fjord**

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## 21 Abstract

22 Comau Fjord is a stratified fjord in the northern part of Chilean Patagonia. In spite its deep waters  
23 being aragonite-undersaturated causing an energetic burden to coral calcification, dense  
24 populations of cold-water corals (CWC) thrive here as well. While the paradox has been  
25 attributed to a rich supply of zooplankton, quantitative data to support this hypothesis are so far  
26 lacking. In this study, we investigated the seasonal and diel changes of the zooplankton  
27 community over the entire water column. We used a Nansen net (100  $\mu\text{m}$  mesh) to take stratified  
28 vertical hauls between the surface and the bottom (0-50-100-200-300-400-450 m). Samples were  
29 scanned with a ZooScan, and abundance, biovolume and biomass were determined for 41 taxa  
30 identified on the web-based platform EcoTaxa 2.0. Zooplankton biomass was the highest in  
31 summer (209 g dry mass  $\text{m}^{-2}$ ) and the lowest in winter (61 g dry mass  $\text{m}^{-2}$ ). Abundance, however,  
32 peaked in spring, suggesting a close correspondence between reproduction and phytoplankton  
33 spring blooms (Chl *a* max. 50.86  $\text{mg m}^{-3}$ , 3 m depth). Overall, copepods were the most important  
34 group of the total zooplankton community, both in abundance (64-81%) and biovolume (20-70%)  
35 followed by mysids and chaetognaths (in terms of biovolume and biomass), and nauplii and  
36 Appendicularia (in terms of abundance). Throughout the year, diel changes in the vertical  
37 distribution of biomass were found with a daytime maximum in the 100-200 m depth layer and a  
38 nighttime maximum in surface waters (0-50 m), associated with the diel vertical migration of the  
39 calanoid copepod family Metridinidae. Diel differences in integrated zooplankton abundance,  
40 biovolume and biomass were probably due to a high zooplankton patchiness driven by biological  
41 processes (e.g., diel vertical migration or predation avoidance), and oceanographic processes  
42 (estuarine circulation, tidal mixing or water column stratification). Those factors are considered to  
43 be the main drivers of the zooplankton vertical distribution in Comau Fjord.

44 **Keywords.** zooplankton, seasonality, diel vertical migration, cold-water corals (CWC),  
45 ZooScan, EcoTaxa, Comau Fjord, Chilean Patagonia.

46

## 47 Introduction

48 Coastal marine ecosystems are among the most productive on earth (Mann & Lazier, 1991). They  
49 provide substantial economic and ecological services, such as high biological production, nutrient  
50 cycling or shoreline stability and erosion control (Escribano, Fernández & Aranís, 2003; Liu et  
51 al., 2010; Pan et al., 2013; Barbier, 2017). The Chilean fjord region extends over large and  
52 complex hydrographic and geomorphologic conditions, particularly rich in productivity and  
53 biodiversity (Försterra, Häussermann & Laudien, 2017; Häussermann, Försterra & Laudien,

2021), but also endangered by human exploitation, which has been increasing significantly – e.g. salmon aquaculture – over the last two decades (Iriarte, González & Nahuelhual, 2010; Buschmann, Niklitschek & Pereda, 2021; Navedo & Vargas-Chacoff, 2021).

The hydrography of the fjords and channels of Patagonia can be considered a transitional estuarine-marine system where a surface layer of silicate-rich terrestrial freshwater meets nitrate- and phosphate-rich marine waters. These fjords receive freshwater from rivers, surface runoff and groundwater flow due to the high rainfall and glacier melting (Pantoja, Iriarte & Daneri, 2011). The upper brackish layer within the top 10 m water depth is usually poor in nitrate and phosphate but rich in silicate and organic matter from terrestrial inputs (Sánchez, González & Iriarte, 2011). Below the halocline, a water mass with higher salinity (>31), named Modified Subantarctic Water (MSAAW), flowing landward from the adjacent oceanic area provides the fjords with macronutrients (nitrate and phosphate). Both water masses, surface-freshwater and MSAAW, generate a two-layer structure in the water column with sharp vertical and more gradual horizontal salinity gradients (Sievers & Silva, 2008; Pérez-Santos et al., 2014; Meerhoff et al., 2019). The summer stratification creates a barrier which may hinder the exchange of nutrients within the water column, altering the functioning of the pelagic food web and productivity patterns (Silva, Calvete & Sievers, 1997; González et al., 2011). During austral spring, the Comau Fjord receives an intense riverine input of fresh water, rich in silicic acid used by bloom-generating diatoms and, thus, leading to high primary production (González et al., 2010). In austral summer, the high concentration of phytoplankton promotes an increase in the abundance and biomass of zooplankton as secondary producers (Antezana, 1999; González et al., 2010). In fjord systems, seasonal patterns are modulated by other oceanographic processes, such as estuarine circulation (Palma & Silva, 2004), tidal regimes and lateral advection (Castro et al., 2011) or water column stratification (Sánchez, González & Iriarte, 2011) influencing the zooplankton biomass and community structure on shorter time scales.

Zooplankton plays an essential role in the functioning of marine ecosystems and in the oceanic carbon cycle. It includes a wide variety of organisms and displays extreme variability in terms of community composition and vertical, seasonal and geographical distribution (Palma & Kaiser, 1993). Many taxa are known to perform diel vertical migrations (DVM), most likely to evade predators (Stich & Lampert, 1981; Iwasa, 1982). According to the predator-evasion hypothesis, migrating zooplankton resides in deep waters during daytime hours where the probability of being perceived by visually orientated predators is lower than if they remained in better illuminated shallow waters, and at night, in the refuge of darkness, they migrate upwards to feed. However, DVM is not performed by all organisms in a zooplankton community or even not by all individuals of one species. For example, while late copepodites and adults of the copepod genus

89 *Metridia* migrate, a large fraction of the young developmental stages remains in surface waters,  
90 saving the energy of performing the DVM, suggesting a lower probability of being perceived and  
91 consumed by visual predators (Hays, 1995). Both, migrant and non-migrant species are important  
92 elements of the biological carbon pump via the production of sinking fecal pellets that transport  
93 carbon from surface waters to the seafloor (Urrère & Knauer, 1981; Fowler & Knauer, 1986;  
94 Emerson & Roff, 1987). Zooplankton also provides a trophic link between primary production  
95 and higher consumers such as fish, birds and mammals, but also invertebrate predators, such as  
96 corals (Nemoto, 1970; Gili et al., 2006; Höfer et al., 2018).

97 Cold-water corals (CWC) rely on zooplankton as their principal food source to maintain their  
98 physiological processes, such as respiratory metabolism and growth (Carlier et al., 2009; Mayr et  
99 al., 2011; Naumann et al., 2011). Therefore, their diet depends on the zooplankton seasonal, diel  
100 and vertical distribution. In Comau Fjord, azooxanthellate scleractinian CWC are wide-spread  
101 even in deep aragonite-undersaturated waters (Häussermann & Försterra, 2007; Fillinger &  
102 Richter, 2013a; Jantzen et al., 2013a). In the latter, the dissolution of exposed parts of the  
103 skeleton, enhanced bioerosion, and reduced CWC growth and survival have been observed  
104 (McCulloch et al., 2012; Maier et al., 2016). The calcification of the CWC skeleton is  
105 energetically costly, and thus food requirements in these adverse environments are higher  
106 compared to aragonite-saturated waters (Maier et al., 2016). According to Fillinger & Richter,  
107 (2013b), in Comau fjord, the CWC *Desmophyllum dianthus* thrives but coral densities decrease  
108 below 280 m despite available substrate, suggesting that lower oxygen and pH concentrations,  
109 combined with a shortage of zooplankton could be limiting coral growth. However, up to now  
110 little is known about the zooplankton of Comau Fjord in the northern Patagonian region and its  
111 role in sustaining the CWC communities living in the fjord. Most studies were carried out in the  
112 central-southern part of Patagonia (from Penas Gulf to Cape Horn, S 46.50° – S 55.55°), while  
113 studies performed in the northern area (from Puerto Montt to San Rafael Lagoon, S 41.20° – S  
114 46.40°) mainly focused on selected microzooplankton taxa in the upper water column, on bulk  
115 measurements of zooplankton biomass, or on the carbon flow through the pelagic food web (e.g.  
116 Palma, 2008 and references therein; Villenas, Soto & Palma, 2009; González et al., 2010, 2011;  
117 Palma et al., 2011; Sánchez, González & Iriarte, 2011). Other studies addressed the physical  
118 oceanographic processes and their effect on zooplankton distribution (e.g. Marín & Delgado,  
119 2009; Castro et al., 2011) and their relationship with zoo- and ichthyoplankton growth and  
120 feeding (Landaeta et al., 2015a,b). Recent studies have investigated zooplankton migration  
121 patterns by acoustic backscatter and vertical velocity profiles (Valle-Levinson et al., 2014; Díaz-  
122 Astudillo, Cáceres & Landaeta, 2017; Pérez-Santos et al., 2018). The information on zooplankton

123 diversity and migration patterns is, however, still very fragmentary and the linkage between the  
124 abundance of CWC and zooplankton supply in Comau Fjord remains unknown.

125 In this study, we aim to describe the diel, vertical, and seasonal distribution of mesozooplankton  
126 groups of Comau Fjord, with a focus on the dominating taxa, particularly those that migrate and  
127 are more likely to aggregate. Samples were collected at day and night hours with vertical net  
128 hauls through the whole water column in spring, summer, autumn and winter. They were  
129 processed with a high-resolution image analysis system (ZooScan, Gorsky et al. 2010). The  
130 zooplankton taxa were identified using EcoTaxa 2.0 (Picheral, Colin & Irsson, 2017), allowing to  
131 assess the influence of seasonal environmental changes on zooplankton dynamics, and the food  
132 naturally available to CWC.

133

## 134 **Materials & Methods**

135 Field work was carried out in Comau Fjord, Northern Patagonia, Chile (Fig. 1). Zooplankton was  
136 sampled four times at a fixed station (42°14.95S, 72°28.83W) in central Comau Fjord: in austral  
137 spring (28<sup>th</sup> September 2016; three days before spring tide-new moon), summer (17<sup>th</sup> January  
138 2017; two days after spring tide-full moon), autumn (7<sup>th</sup> May 2017; three days before spring tide-  
139 full moon), and winter (29<sup>th</sup> July 2017; one day before neap tide), during both day (noon) and  
140 night (midnight). Samples were collected with a 70 cm-diameter Nansen closing net (mesh size:  
141 100 µm) equipped with a non-filtering cod end. Vertical hauls were carried out at 0.45 m s<sup>-1</sup> to  
142 sample the depth strata 0-50-100-200-300-400-450 m. Immediately after the collection, the  
143 samples were sieved through a 50 µm mesh and preserved in 4% borax-buffered formaldehyde  
144 for laboratory analyses. After every zooplankton haul, a CTD multi-probe (SBE 19plusV2  
145 Profiler - with RS 232 Interface, Sea-Bird Electronics Inc.) was deployed from the surface to the  
146 bottom, measuring conductivity, temperature, oxygen, pH and chlorophyll *a* (Chl *a*)-fluorescence.

147 In the laboratory, fixed zooplankton samples were washed with fresh water and prepared for  
148 analysis with a ZooScan digital imaging system (Grosjean et al., 2004; Gorsky et al., 2010).  
149 ZooScan (CNRS patent, [www.hydroptic.com](http://www.hydroptic.com)) provides a quick and reliable method for the  
150 analysis of preserved plankton samples, storing digitized images for later examination,  
151 reprocessing and dissemination. Concentrated samples were subsampled with a Folsom plankton  
152 splitter to avoid images being cluttered with more than approximately 1000-1500 individuals. Up  
153 to six binary splitting steps were carried out (corresponding to a minimum 1/64<sup>th</sup> fraction of the  
154 original sample). Routinely, the two final splits were scanned with ZooScan yielding images of  
155 2400-dpi resolution (14200×22700 pixels). The hinged base of the ZooScan allowed the recovery

156 of the complete undamaged subsample, which was later stored in 70% ethanol for archiving. Most  
157 overlapping individuals on the scanning surface were manually separated to ensure an even  
158 distribution before scanning. Image analysis was performed with the software ZooProcess  
159 (Gorsky et al., 2010), a plug-in for the image processing and analysis software ImageJ (Schneider,  
160 Rasband & Eliceiri, 2012). The processing involved (1) the automatic subtraction of background  
161 noise, (2) the automatic thresholding and detection of objects, and (3) the automated storage of  
162 detected objects in separate images (“vignettes”). Below 300  $\mu\text{m}$ , organisms were often too  
163 blurred to be identified. Thus, the ZooScan detection limit was set at the standard of 300  $\mu\text{m}$  so  
164 that detected zooplankton sizes ranged from 0.3 to 59 mm. The automatic processing of the scans  
165 was successful in 75-80% of the cases where vignettes with one individual were obtained.  
166 However, despite the manual separation, some individuals overlapped, resulting in vignettes with  
167 two or more objects. These objects on the pictures were manually separated using the “separation  
168 with mask” tool of the ZooProcess software. Separated vignettes were stored, while the original  
169 vignette, containing multiple objects was eliminated from the database to avoid duplicate counts.  
170 In some cases, the separation of individuals was not possible as cutting the vignette would mean  
171 losing information about the morphology of the organisms (i.e. cutting overlapping urosomes  
172 from two different copepods or small copepods embedded in cnidarians). Overall, the  
173 contribution of vignettes with multiple objects that could not be separated was always <10% of  
174 the total amount of vignettes.

175 Vignettes were subjected to the semi-automated taxonomic classification in EcoTaxa 2.0  
176 (Picheral, Colin & Irsson, 2017). This web-based machine learning application uses training sets  
177 of expert-identified taxa and random forest classification to automatically identify and sort the  
178 objects. Although EcoTaxa contains more than 160 million objects on its server, no ZooScan  
179 training set was available for Patagonian waters. Therefore, manual identification of individuals  
180 on a subset of the images was first necessary to train an initial model, which was later used by the  
181 system to classify the scanned organisms. The initial learning set with Patagonian organisms  
182 improved progressively its prediction by sorting more objects into the given categories. This  
183 produced the final learning set for the classification of the entire image data set. At the end, all  
184 classified objects were individually validated to assure a correct classification. The organisms  
185 were classified to the lowest possible taxonomic level; for most copepods this was the family  
186 level. However, small calanoid copepods (<1.5 mm) were not distinguishable on family level and  
187 were thus comprised as one category: “Calanoids (<1.5 mm)” including five groups (copepodites  
188 (all calanoid taxa, <1mm), Clausocalanidae, Microcalanidae, small Calanidae (*Neocalanus* spp.)  
189 and Paracalanidae). Developmental stages were included in the corresponding taxon as long as  
190 they were clearly identifiable. Only for calanoid copepods, the classification of some

developmental stages was not clear and therefore, they were included into the category “copepodites”, which included developmental stages of different calanoid copepod taxa. The category Cnidaria was constituted by organisms from the class Hydrozoa (mostly medusa and Siphonophorae). Another category contained all images that were out of focus (“bad focus”) and likely comprised individuals from all copepod taxa, in total 6,766 vignettes. From the total of 83,516 vignettes, 23,227 could not be assigned to zooplankton taxa, but were labelled as “detritus”, “feces”, “fiber”, “leg” “bubble” and “other” and were not considered in our analyses. ZooProcess provides information about the length and width of each object, allowing the calculation of its volume as a proxy for its biomass (Gorsky et al., 2010). The program automatically fits an ellipse around the object, from which the major and minor axis and volume (V) is computed:

$$V (mm^3) = \frac{4}{3} \times \pi \times \frac{major\ axis (mm)}{2} \times \left( \frac{minor\ axis (mm)}{2} \right)^2$$

Biovolume (BV) was then calculated as the sum of the volumes of all objects ( $\Sigma V$ ) divided by the fraction of the sample (e.g.,  $F=1/64$ ) and by the volume filtered by the Nansen net ( $V_N$ ):

$$BV (mm^3/m^3) = \frac{\frac{\Sigma V (mm^3)}{F}}{V_N (m^3)}$$

$V_N$  was calculated as:

$$V_N (m^3) = \left[ \pi \times \left[ \frac{net\ diameter (m)}{2} \right]^2 \right] \times depth\ interval (m) \times filtration\ efficiency$$

where filtration efficiency was assumed as the theoretical 100% efficiency (value = 1). Flowmeter readings were not used because the speed of the net haul was below the measuring range of the mechanical flowmeter.

For the estimation of biomass, a regression between the dry mass (DM) of a specimen and its body area [ $DM (\mu g) = (a A^b)$ ] was used (Hernández-León & Montero, 2006; Lehet & Hernández-León, 2009), where  $A$  is the area ( $mm^2$ ) of each scanned individual. The regression required different conversion factors depending on the organism, as for instance, gelatinous zooplankton with high water content may not be compared to crustaceans or echinoderms (Table 1). Such coefficients have been successfully published in previous studies for mid-latitude shelf areas (Marcolin, Gaeta & Lopes, 2015) or the Chilean upwelling region (Tutasi & Escribano, 2020).

219 The biomass (B) of each taxon was then calculated as the sum of the individual dry masses of the  
220 respective taxon ( $\Sigma DM$ ) divided by the fraction of the sample (e.g.,  $F=1/64$ ) and by the volume  
221 filtered by the Nansen net ( $V_N$ ):

$$222 \quad B \text{ (mg dry mass/m}^3\text{)} = \frac{\frac{\Sigma DM(mg)}{F}}{V_N(m^3)}$$

223 Biovolume and biomass were calculated to obtain the sum of the values of all individuals for a  
224 given taxon. In multiple vignettes, the automatic calculation of biovolume and biomass was not  
225 possible because of overlapping specimens from different taxa. Then, the organisms were counted  
226 manually and biovolume and biomass were estimated by multiplying the mean volume or DM of  
227 the given taxon from all automatic calculations by the extra number of multiple vignettes. Groups  
228 with large size variability (e.g., chaetognaths, cnidarians or Euchaetidae) were divided into two  
229 categories, small (0.003-4.242 mm<sup>3</sup>) and large (4.243-90.083 mm<sup>3</sup>) in order to get a better  
230 biovolume/biomass assessment.

231 The integrated values of abundance, biovolume and biomass, were calculated down to 400 m  
232 water depth for all seasons, taking out the last 50 m from summer and winter, to make it  
233 comparable to the spring and autumn seasons, where samples were collected down to 400 m.

234 The relationships among physicochemical variables and each taxon abundance were analyzed  
235 using a redundancy analysis (RDA). RDA is a constrained ordination procedure which allows the  
236 assessment of how much of the variation of one set of response variables (i.e. zooplankton  
237 abundances) is explained with another set of variables (i.e. physicochemical variables). The RDA  
238 is a multiresponse analysis which summarizes the linear relationships among dependent and  
239 independent variables into a matrix followed by a principal component analysis (PCA). Mean  
240 values of temperature, salinity, oxygen, and Chl *a* (log transformed) for the entire water column  
241 were used as explanatory variables. RDA was performed in R (R Core Team, 2021) by using the  
242 *rda* function of the *vegan* package (Oksanen et al., 2019). The problems caused by non-normal  
243 distributions in testing the significance of RDA results were solved by a permutation test (10,000  
244 iterations) (Borcard, Gillet & Legendre, 2011) using the *anova.cca* function from the *vegan*  
245 package. All abundance data were logarithmically transformed before analysis.

246 The centroid depths (CD) of the zooplankton groups for each sampling event were calculated as:  
247  $CD = \sum (p_k \times z_k) / \sum p_k$ , where  $p_k$  is the number of organisms in the stratum  $k$ , and  $z_k$  is the mean  
248 depth of the stratum  $k$ . Due to the lack of replication, these values were compared using a  
249 contingency table by means of a Chi-square test, in order to test the significance of night-day  
250 changes in CD. The Chi-square test compares the critical values to assess their significance



251 according to their degrees of freedom ( $df = (r - 1)(c - 1)$ ), where  $r$  is the number of rows and  $c$   
252 the number of columns in the contingency table. After showing significant day-night differences  
253 in CD (Chi-square  $Pr < 2.2 \times 10^{-16}$ ,  $p < 0.001$ ), a dissimilarity analysis (function *simper* from *vegan*  
254 package) was performed to assess those zooplankton groups that cumulatively contribute at least  
255 to 70% of the night-day differences observed.

256

## 257 Results

258 The physicochemical parameters measured throughout the water column of Comau Fjord showed  
259 a stronger seasonal variability in surface waters (0-50 m) than in deep waters (50-450 m) (Fig. 2).  
260 The temperature profile indicated summer stratification down to 17 m water depth, followed by  
261 surface cooling, breakdown of the thermocline in autumn, and reverse temperature gradients in  
262 winter and spring. Accordingly, the surface temperature values were the lowest in winter and  
263 spring (8.6-11 °C), and the highest in summer (16.7 °C), getting cooler again in autumn (12 °C).  
264 In deeper waters, temperatures were more stable with an average value of  $11.4 \pm 0.2$  °C (mean  $\pm$   
265 ~~sd~~SD; Fig. 2A). Salinity was between 10-30 in the upper 20 m and  $32.9 \pm 0.4$  below 20 m (Fig.  
266 2B). The pH ranged between 8.5 and 7.7 in the upper 50 m and was  $7.9 \pm 0.1$  in deeper waters in  
267 all seasons, except for autumn, where we interpret the sudden drop of pH values as an instrument  
268 malfunction (Fig. 2C). Oxygen concentration showed the largest variations in the upper 50 m  
269 during the spring season ( $137.2\text{-}410.5 \mu\text{mol kg}^{-1}$ ), while below 100 m depth it was on average  $180$   
270  $\pm 9.3 \mu\text{mol kg}^{-1}$  (Fig. 2D). The chlorophyll *a* (Chl *a*) concentration peaked in early spring ( $50.86$   
271  $\text{mg Chl } a \text{ m}^{-3}$  at 3 m depth), followed by a decrease towards the end of the summer and low  
272 values through autumn and mid-winter ( $2.5\text{-}5 \text{ mg Chl } a \text{ m}^{-3}$  at 5-10 m depth) (Fig. 2E). Below 25  
273 m, the concentration of Chl *a* was  $<1.8 \pm 0.5 \text{ mg m}^{-3}$  throughout the year (Fig. 2E).

274 The zooplankton community exhibited large seasonal and diel differences. Abundance, integrated  
275 over the upper 400 m of the water column (individuals  $\text{m}^{-2}$ ) showed the highest values in spring  
276 and the lowest during autumn for both; day and night (Fig. 3A). The integrated biovolume ( $\text{cm}^3$   
277  $\text{m}^{-2}$ ) and biomass ( $\text{g dry mass m}^{-2}$ ) showed a different pattern with the highest values in summer  
278 and the lowest in autumn and winter (Fig. 3B, C). Diel differences in integrated abundances  
279 generally showed higher values during the day than at night, except for autumn (Fig. 3A). Diel  
280 differences in integrated biovolume and biomass were surprisingly large, with generally higher  
281 values during the night, except for spring (Fig. 3B, C), indicating that during daytime, particularly  
282 in spring, zooplankton was more numerous but smaller in size. Overall, the centroid depth  
283 dissimilarity analysis showed that in spring, fewer taxa contributed to the significant night-day

Commenté [d1]: You should put the result of the test here, not in the materials and methods

284 differences both, in abundance and biovolume, while in summer and autumn the number of  
285 contributing zooplankton groups increased (Fig. 4).

Commenté [d2]: Because it deals with vertical distribution (centroid depth), then it should be in the following paragraph....

286 The zooplankton vertical distribution showed that abundance exhibited the highest values during  
287 day and night in the 0-50 m layer at all seasons, with the exception of autumn during day time. ~~In~~  
288 ~~autumn, the daytime peak was absent with, where abundances showed overall~~ low values  
289 throughout the water column (Fig. 5A). The lowest abundances were found in >300 m depth in  
290 spring, summer and winter, and in 50-100 m depth in autumn during day and night time (Fig. 5A).  
291 The highest biovolume and biomass daytime values were observed in 0-50 m and 100-200 m  
292 water depth in spring and summer, and in 100-300 m in autumn. However, in winter, daytime  
293 biovolume and biomass were similarly low throughout the entire water column (Fig. 5B, C). At  
294 night, the highest biovolume and biomass values were found at the surface (0-50 m) in all  
295 seasons. The lowest biovolume and biomass values were found in >300 m depth in spring and  
296 summer, in 50-100 m and 200-300 m depth in autumn and in >50 m depth in winter (Fig. 5B, C).

297 The ~~taxa-taxonomic~~ composition greatly differed ~~between when considering~~ biovolume and  
298 abundance (Fig. 6, 7A; biovolume—not biomass—was used to better represent the gelatinous  
299 taxa). Copepoda were the largest group, constituting 20-70% of the total biovolume, and 64-81%  
300 of the total abundance. Within the copepod community, individuals smaller than 1.5 mm included  
301 (a) copepodites and adults of small calanoid genera, such as Clausocalanidae, Microcalanidae,  
302 Neocalanidae and Paracalanidae; (b) cyclopoids of the genera *Oithona* and *Oncaea*; (c)  
303 harpacticoids; and (d) nauplii. These small copepods accounted for 58-86% of the total abundance  
304 within the copepod community. Overall, 14 out of 41 taxa contributed 45-98% of the total  
305 biovolume and 45-86% of the total zooplankton abundance (Fig. 6). The other 27 taxa were  
306 constituted by copepods (Acartiidae, Aetideidae, Calanoida (non-identifiable), Candaciidae,  
307 Centropagidae, Copepoda (non-identifiable), Eucalanidae, Heterorhabdidae, Lucicutiidae,  
308 Oncaeidae, Pontellidae, Rhincalanidae), Actinopterygii (eggs and larvae), Appendicularia,  
309 Ascidiacea (larvae), Brachiopoda (larvae), Bryozoa (larvae), Cirripedia (nauplii and cypris),  
310 Cladocera, Decapoda (zoa), Echinodermata (larvae), eggs, Isopoda, Nemertea (larvae),  
311 Platyhelminthes (larvae), Polychaeta (larvae) and non-identifiable organisms.

Commenté [d3]: Dominant? Largest could be confusing, large being indicative of size...

Commenté [d4]: Larvae? Because you may have also megalopae...

312 The contribution of groups other than copepods to the total zooplankton community differed  
313 regarding biovolume and abundance. For biovolume, Cnidaria (2-23%) and Mysida (1-21%)  
314 constituted a large part of the zooplankton community, followed by Chaetognatha (3-15%).  
315 Metanauplii and calyptopis stages of Euphausiacea were found mostly in spring and summer,  
316 accounting for 0-7% of the total biovolume. A single adult specimen found during winter in 0-50  
317 m water depth at night raised the total biovolume of this taxon to 65%. Regarding abundance,

318 ~~Copepoda~~ nauplii (2-9%, with ~~its~~ their maximum in spring and minimum in autumn) and  
319 Appendicularia (0.4-6%) were the second and third most abundant groups after Copepoda,  
320 respectively. The fourth most abundant group differed among seasons: Echinodermata larvae (5%  
321 in spring), Mollusca larvae (2-5% in summer, 3-9% in autumn), Ostracoda (4-5% in autumn) and  
322 Bryozoa larvae (2-10% in winter). Across all samples and seasons, taxa that represented more  
323 than 5% of total biovolume were Cnidaria (13%), Calanidae (12.9%), Mysida (12.7%),  
324 Metridinidae (12.6%), Chaetognatha (9.6%) and Euchaetidae (8.7%). The most abundant groups  
325 with more than 5% of the total abundance were Harpacticoida (14.0%), Cyclopoida (9.1%) and  
326 Metridinidae (8.4%) (Fig. 6, 7A).

327 The integrated abundance of small and large copepods and chaetognaths showed the highest  
328 abundance in spring and generally low values in autumn and winter (Table 2). Highest  
329 abundances of copepod nauplii and cnidarians were found in early spring with a minimum in  
330 autumn and raising up again in late-winter. Mysida presented a stable abundance throughout the  
331 year with a minimum in summer. These groups clearly presented different vertical distributions  
332 (Fig. 7). Metridinidae, a family of large copepods with high biovolume and abundance, resided  
333 generally above 200 m, exhibiting a peak between 100-200 m during the day and a bimodal  
334 distribution (peaks at 0-50 m and 100-200 m) during the night (Fig. 7B). The largest individuals  
335 ( $0.62 \pm 0.3 \text{ mm}^3$ ; average size  $\pm$  sd) were found in 100-200 m waters during the day and night, but  
336 at nighttime, a fraction of those large individuals was found in shallow waters (Fig. 8A). In  
337 spring, a significant proportion of the Metridinidae was also found during the day in shallow  
338 waters where smaller individuals ( $0.14 \pm 0.08 \text{ mm}^3$ ) were observed (Fig 8A). The Calanidae were  
339 found mainly in intermediate waters (100-300 m) during daytime and nighttime where the largest  
340 individuals ( $1.37 \pm 0.5 \text{ mm}^3$ ) resided (Fig 8B). Small specimens of Calanidae ( $0.30 \pm 0.003 \text{ mm}^3$ )  
341 were found in shallow waters also during daytime in spring (Fig 8B). At night, in spring, summer  
342 and winter, individuals were present in shallow waters (<50 m depth). Despite the low number of  
343 Calanidae found at shallow waters, they contributed considerably to the biovolume in these layers  
344 (Fig. 7C). Euchaetidae showed higher abundance and biovolume in the deeper part of the water  
345 column during day time, ascending to shallower waters at night (Fig. 7I). Small copepods of the  
346 taxa Harpacticoida and Cyclopoida were very abundant, but, as expected, they only represented a  
347 small fraction of the total biovolume. Harpacticoida were mainly abundant below 50 m during  
348 both, day and night time, and their abundance increased with depth (Fig. 7D). Cyclopoida were  
349 mainly present in the upper 200 m during day and night in all seasons (Fig. 7H). Cnidarians were  
350 overall not very abundant, but constituted an important fraction of the biovolume, especially in  
351 200-300 m depth during day in all seasons, and 50-100 m depth during day time in summer and  
352 autumn. At night, they were mainly present in deeper layers 200-450 m (Fig. 7E). Mysida were

353 mainly present in >100 m depth during the day, while a small fraction was found at night between  
354 0-50 m in spring and between 50-100 m in autumn and winter (Fig. 7F). Chaetognatha were  
355 present mainly in intermediate layers between 100-300 m during day and night in all seasons (Fig.  
356 7G).

357 Overall, the four predictor variables (fluorescence (Chl *a*), oxygen, temperature, salinity)  
358 explained 44.3% and 33.69% of the total variation in abundance and biovolume, respectively  
359 (Fig. S1). The first axis of the RDA explained 34.15% (abundance) and 25.4% (biovolume) of the  
360 total variation while the second axis only accounted for 10.15% (abundance) and 8.29%  
361 (biovolume). Relationships between the parameters and the first RDA axis were highest for Chl *a*,  
362 while temperature was related to the second RDA axis. The ~~effect of the~~ physicochemical  
363 variables ~~on abundance and biovolume had~~ higher influence on abundance and biovolume  
364 between daytime and nighttime and within the most productive seasons (spring and summer) ~~on~~  
365 in shallow waters (0-50 m) than in deeper ones. The abundance and biovolume of the major  
366 zooplankton groups were positively influenced by Chl *a* concentration, except for mysids and  
367 ostracods, while temperature did not seem to have a strong influence on the majority of the  
368 groups, except for the large zooplankton (amphipods, Euphausiacea and euphausiids) (Fig. S1).

369

## 370 Discussion

### 371 *Physicochemical properties and mesozooplankton seasonal dynamics*

372 The physicochemical observations in our study are consistent with earlier descriptions from  
373 Comau Fjord, a temperate fjord connected to the Pacific Ocean by the Chacao Channel and  
374 Ancud Gulf. Comau Fjord has an estuarine circulation and is characterized by a strong  
375 pycnocline, where surface waters (0-50 m) present higher variability than the deeper and quasi-  
376 homogeneous water layer (Fig. 2). Clear differences were observed in temperature, pH, oxygen  
377 and Chl *a* in relation to the season, likely caused by the strong seasonal variability in solar  
378 radiation (maximum between spring and summer), precipitation and river discharge (maximum in  
379 late autumn and winter) (González et al., 2010).

380 In Comau Fjord, a thermal inversion of the surface water (0-50 m) in winter is visible (Fig. 2),  
381 probably due to heat loss in the surface layer caused by winds and the discharge of cold  
382 freshwater from rivers and glaciers (Silva, Calvete & Sievers, 1997). Later in the year, the thermal  
383 density stratification stabilizes the water column, triggering spring phytoplankton blooms (Iriarte  
384 et al., 2007), which usually follow rain events and thus the input of nutrients (unpublished data).

Commenté [d5]: Please indicate if this is yours or of other authors...

385 This leads to a strong increase in Chl *a* (Fig. 2E). As shown by previous studies in the area (Palma  
386 & Silva, 2004; Vargas et al., 2008; González et al., 2010), this peak in Chl *a* was most likely due  
387 to blooming chain-forming diatoms. The high phytoplankton biomass is expected to be grazed  
388 predominantly by copepods, increasing their biomass and establishing the classical diatom-to-  
389 zooplankton food web (Palma & Silva, 2004; Vargas et al., 2008; González et al., 2010).  
390 Accordingly, we found the highest zooplankton abundance during the most productive season, the  
391 austral spring. At this time, copepod nauplii and young stages of calanoid copepods, which are  
392 indicative of intense zooplankton reproduction, accounted for a large proportion of the  
393 zooplankton community. In summer, biovolume and biomass reached their maxima (Fig. 3),  
394 which together with a lower abundance, suggests the presence of larger individuals or different  
395 groups including species with larger individuals. Subsequently, in autumn and winter,  
396 zooplankton abundance, biovolume and biomass decreased. At this time, primary production  
397 should be low, as reflected by low Chl *a* values (Fig. 2E), and likely zooplankton growth was  
398 limited by food availability (Escribano et al., 2007). In winter, the plankton in the fjord typically  
399 shifts towards a microbial loop based community grazed by heterotrophic nanoflagellates, which  
400 become the main mesozooplankton prey (Vargas et al., 2008; González et al., 2010) however, this  
401 does not support a high secondary production.

402 The northern part of the Chilean fjord region, the area between Puerto Montt and Guafo Mouth,  
403 represents the most productive area of Patagonia in terms of primary production and zooplankton  
404 biomass (Palma, 2008). In contrast, the phytoplankton production in the southern area is low due  
405 to the stronger influence of glaciers, resulting in cold, fresh and turbid waters (Palma & Rosales,  
406 1997; Palma & Silva, 2004; Iriarte et al., 2007; Palma, 2008), and consequently low zooplankton  
407 survival and growth (Giesecke et al., 2019). Previous studies described ranges for zooplankton  
408 biovolume, expressed as the plankton wet volume, of 65 to more than 1386 ml zooplankton 1000  
409 m<sup>-3</sup> outside Comau Fjord, in the Inner Sea off Chiloé island (Palma, 2008). This is in line with the  
410 present results (250-1500 ml zooplankton 1000 m<sup>-3</sup>), showing an especially high biovolume  
411 during ~~the summer season in Comau Fjord~~. Palma & Rosales (1997) also found the highest values  
412 of zooplankton **biomass** in the northern part (interior of Reloncaví Fjord and Ancud Gulf) with  
413 values that ranged between 56- ~~and~~ 1626 ml zooplankton 1000 m<sup>-3</sup>, but a low zooplankton  
414 biomass in the inner of Comau Fjord. The observed variations may potentially be due to (a)  
415 interannual differences with a much lower Chl *a* concentration during the same season of their  
416 year of study (Ramírez, 1995); and/or (b) methodological and analytical differences, e.g. different  
417 sampling gears and proxies for biomass estimation (i.e. measurement of zooplankton wet volume  
418 vs. image analysis in this study). ~~The Moreover, the~~ vertical hauls used in this study may most  
419 likely underestimate the abundance of very motile organisms able to avoid the net, such as **adult**

Commenté [d6]: Biovolume (because of the unit)?

420 euphausiids, megalopae of *Munida gregaria* (~~megalo~~pae) or large fish larvae, which are also  
421 important in terms of density and biomass across west Patagonia.

Commenté [d7]: If you have a reference for this (in particular for *Munida*, why this one)...

422 Pronounced seasonality of environmental variables often results in high biological production and  
423 are associated to seasonal changes in the holoplankton community (Mauchline, 1998; Balbontín  
424 & Bustos, 2005; Aracena et al., 2011), and meroplankton abundance (Ladah et al., 2005;  
425 Landaeta & Castro, 2006). In upwelling areas, where seasonal changes of environmental  
426 conditions are also pronounced, the highest abundances of *Calanus chilensis* were observed in  
427 spring and low abundances through autumn and winter (Hidalgo & Escribano, 2007). The  
428 population dynamics of copepods in this study followed the phytoplankton seasonal cycle, with  
429 the highest abundance in spring, associated with the maximum concentration in phytoplankton,  
430 decreasing over time with minimum abundances in autumn and winter (Table 2). Considering that  
431 copepods are the main food source for cnidarians and chaetognaths (Palma & Kaiser, 1993), the  
432 higher abundance of carnivorous zooplankton occurring in spring, can be attributed to the large  
433 copepod abundance at this time of the year. The overall decline in copepod abundance from  
434 spring to winter may be explained by the decrease in phytoplankton occurrence and the increasing  
435 predation pressure exerted by carnivorous zooplankton.

Commenté [d8]: This sounds a bit redundant with paragraph 385-401. Both can be merged...

436 In fjord systems, oceanographic processes such as estuarine circulation, tidal mixing or water  
437 column stratification may influence the composition and abundance of zooplankton communities  
438 (Palma & Silva, 2004; Sánchez, González & Iriarte, 2011). Overall, copepods were the main  
439 contributors to the total biomass and biovolume of the zooplankton community, especially during  
440 summer (69-78%). This is in agreement with previous studies showing that in Chilean fjords,  
441 planktonic crustaceans, such as copepods and euphausiids, have the highest abundances and  
442 biomasses, followed by chaetognaths and gelatinous plankton (Defren-Janson, Schnack-Schiel &  
443 Richter, 1999; Palma & Silva, 2004). Copepods are the most abundant and diverse component of  
444 marine zooplankton worldwide (Mauchline, 1998), and the abundance of small copepods (<1.5  
445 mm) generally surpasses the abundance of larger ones (Fransz, 1988; Gallienne & Robins, 1998;  
446 Gallienne, Robins & Woodd-Walker, 2001). Similarly, small copepods accounted for 58-86% of  
447 the total copepod community in the present study. Another important contributor to zooplankton  
448 communities in Chilean fjords is the euphausiid *Euphausia vallentini*. The present study revealed  
449 young stages of euphausiids during spring and summer but only one adult specimen in winter,  
450 indicating that euphausiids are present in Comau fjord, but were not caught efficiently in our  
451 samples. This is likely related to the small volume filtered by our net, the patchy distribution of *E.*  
452 *vallentini*, as well as the ability of these micronekton organisms to avoid small or slow  
453 approaching nets (Brinton, 1962). Like euphausiids, mysids can form dense swarms, making them  
454 a potential food resource for a wide range of organisms, from predatory fishes to benthic CWC.

Commenté [d9]: Redundant with lines 418-421

455 Despite their important contribution to the total zooplankton biovolume in deep waters (Fig. 7F),  
456 poor attention has been given to their presence in Chilean Patagonia. To our knowledge, there are  
457 only two studies describing mysids in this area: Guglielmo & Ianora (1997) found that the most  
458 abundant species for the Strait of Magellan is the deep-dwelling *Boreomysis rostrata*; Díaz-  
459 Astudillo, Cáceres & Landaeta (2017) found higher abundances of mysids during night and inside  
460 the Reloncaví Fjord and Gulf of Ancud. Thus, this study constitutes the first record of mysids in  
461 Comau Fjord.

#### 462 ***Zooplankton diel vertical distribution and migration***

463 Diel Vertical Migration (DVM) is usually associated with differences in light intensity within the  
464 photic zone, taking place periodically in 24 h cycles (Brierley, 2014). During daytime,  
465 zooplankton organisms migrate to deeper, darker waters to avoid visual predators, such as fishes,  
466 while they come to the surface for feeding at night (Hays, Webb & Frears, 1998). In Comau  
467 Fjord, DVM seemed to be related to the size of the zooplankton. Here, a large proportion of  
468 zooplankton organisms that inhabited the intermediate waters (100-300 m) during daytime  
469 ascended to shallow waters at night while small organisms (individuals <1.5 mm), composed  
470 mainly of cyclopoids and harpacticoids, did not perform a clear DVM. Throughout the year,  
471 Cyclopoida centroid depth fluctuated between 40-85 m water depth and Harpacticoida between  
472 149-254 m water depth. These two copepod groups, however, differed with regard to their vertical  
473 distribution. The highest abundances of cyclopoid copepods were mainly found in the upper 100  
474 m, whereas harpacticoids were found between 200-400 m (Fig. 7D).

475 Large individuals from several calanoid copepod taxa (Metridinidae, Calanidae, Euchaetidae),  
476 mysids, chaetognaths and cnidarians did perform DVM over 300 m. This agrees with the findings  
477 by Hays (1995) that DVM is pronounced in large and pigmented species due to their  
478 susceptibility of being perceived by visually orientated predators. Similarly, studies in northern  
479 Patagonia have shown that conspicuous zooplankton organisms tend to avoid well illuminated  
480 waters (Villenas, Soto & Palma, 2009). According to Hays, Kennedy & Frost (2001), large  
481 individuals of *Metridia* usually reside in deep waters, and only a fraction of these ascends to  
482 shallow waters at night, whereas smaller individuals stay at the surface continuously. In Comau  
483 Fjord, Metridinidae showed the highest values of biovolume and abundance at intermediate  
484 depths (100-200 m) during daytime, but in the surface layer (0-50 m) at night (Fig. 7B), which  
485 suggested the migration of the largest individuals towards the surface (Fig 8). In spring, a  
486 significant proportion of the Metridinidae population was found in shallow waters during the day.  
487 This was probably due to the smaller size of the individuals and the higher amount of food  
488 available here. Following the same pattern as Metridinidae, larger individuals of Calanidae were

**Commenté [d10]:** You first have to explain that you observed DVM, based on your results... Then you can elaborate on the size.

**Commenté [d11]:** Please avoid as there is no statistics for this particular point.

**Commenté [d12]:** Yes, and you have the corresponding observation Fig. 8A...



489 found in deeper waters during daytime, whereas small specimens were found in shallow waters  
490 also during daytime in spring, where they may not be hunted by visual predators due to their  
491 small size. It is possible that those small organisms were individuals of earlier life stages (i.e.  
492 small) or species with a smaller size. Euchaetidae also performed DVM where most of the  
493 organisms lived continuously in the deepest waters (200-450 m) during the day, but a small  
494 proportion of the population migrated to the surface during night (Fig. 7I).

495 Mysids were mainly detected in deep waters (>200 m water depth) accounting for up to 70% of  
496 the total zooplankton biovolume. According to Sánchez, González & Iriarte (2011), the  
497 distribution of mesozooplankton in the upper 30 m is strongly influenced by the salinity  
498 stratification, which may act as a barrier for the DVM of some oceanic larger zooplankton  
499 organisms. The variability in surface temperature, salinity, pH, and oxygen might act as a barrier  
500 to the mysids' upward migration in Comau Fjord, resulting in the low mysid abundance in the  
501 surface layer.

502 Chaetognaths were distributed throughout the water column with the highest biovolumes between  
503 100-300 m, during both day and night (Fig. 7G). This is in accordance with the distribution found  
504 by Guglielmo & Ianora (1995) for the Strait of Magellan. Cnidarian highest biovolumes were  
505 found in 50-100 m and in deeper layers (>200 m) during day and night, respectively. South of  
506 Comau Fjord, between Boca del Guafo and the Pulluche Channel, the vertical distribution of  
507 cnidarians (i.e. Hydromedusae and Siphonophorae) indicated highest abundances in the upper 100  
508 m and only in deeper waters in Moraleda and Darwin channels (Palma, Apablaza & Soto, 2007).

509 In the 50-100 m water layer a minimum in zooplankton abundance and biomass was found in all  
510 seasons (Fig. 5). This “zooplankton gap” could be related to a high concentration of predators in  
511 this water depth. In northern Chilean fjords, the high biomasses and abundances of gelatinous  
512 organisms are correlated to a decrease in chitinous biomass of other organisms (Palma & Silva,  
513 2004; Villenas, Soto & Palma, 2009). It is known that chaetognaths and cnidarians can grow at  
514 fast rates, forming dense aggregations that seasonally dominate the zooplankton biomass  
515 (Casanova, 1999; Brodeur, Sugisaki & Hunt Jr, 2002) by feeding voraciously on copepods and  
516 larvae from other organisms (Lie et al., 1983; Palma & Rosales, 1997). In our study, biovolume of  
517 carnivorous organisms ranged between 18 and 83% of the total zooplankton in this water layer,  
518 reaching particularly high values in summer and autumn (50-83%) (Fig. 9). These previous  
519 evidences suggest that carnivorous organisms may have reduced zooplankton abundance in this  
520 depth stratum and predation may play a role in structuring the vertical zooplankton community in  
521 Comau Fjord.

Commenté [d13]: Would any observation in your data show this? RDA?



522 Mesozooplankton plays an important role in the food web as a food source for many pelagic and  
523 benthic organisms (González et al., 2013), including CWC (Gili et al., 2006; Carlier et al., 2009;  
524 Mayr et al., 2011). Due to the difficulty of studying CWC *in situ*, little is known about their  
525 natural diet and its availability. In Comau Fjord, CWC thrive also in deeper, naturally acidified  
526 waters (Häussermann & Försterra, 2007; Försterra, Häussermann & Laudien, 2017). Although the  
527 environment is unfavorable, this might be due to the high ingestion rate (e.g. of the CWC  
528 *Desmophyllum dianthus*) which showed a positive impact on their calcification rates, regardless  
529 of the seawater pH (Martínez-Dios et al., 2020). Zooplankton abundance and biomass are highly  
530 influenced by seasonality. Therefore, CWC must be adapted to differences in food availability,  
531 i.e. high zooplankton abundances during spring and summer and low concentrations during  
532 autumn and winter. Indeed high growth rates of *D. dianthus* were found in summer (Jantzen et al.,  
533 2013b) which may be associated with higher zooplankton availability. In winter, when  
534 zooplankton biomass is low, CWC may slow down their metabolism to cope with the lower food  
535 availability (Naumann et al., 2011). Only recently, a study confirmed that *D. dianthus* preyed on  
536 medium and large sized calanoid copepods and euphausiids (Höfer et al., 2018). Based on the  
537 present DVM data, shallow-dwelling CWC in Comau Fjord might feed on small copepods (e.g.  
538 cyclopoid and calanoid) during the day, and on larger organisms during night hours when  
539 zooplankton migrates upwards. Deeper-dwelling corals, by contrast, may mainly encounter larger  
540 prey, such as mysids or large calanoid copepods (Calanidae, Euchaetidae) and thus may gain  
541 enough energy to upregulate their internal pH in an acidified environment.

542 In the present study, zooplankton abundance, biovolume and biomass, integrated over the entire  
543 water column, differed between day and night. There are few zooplankton groups that contributed  
544 to the day-night differences in their vertical distribution. For instance, Rhincalanidae influenced  
545 the differences between day and night in spring and autumn as the centroid depth changed for  
546 biovolume but not for abundance (Fig. 4), indicating that probably ca. 10% of the difference are  
547 caused by the migration of large individuals. Mysida contributed to ca. 20% of the differences  
548 between day and night both for abundance and biovolume in spring, while in summer the  
549 percentage of biovolume was much less than for abundance, suggesting that a larger number of  
550 smaller individuals were migrating and causing the differences between day and night centroid  
551 depth (Fig. 4). These differences were probably due to the high patchiness the zooplankton  
552 exhibited during day and night, especially by larger zooplankton individuals, and those able to  
553 form swarms (i.e. mysids and euphausiids). Biological processes, such as diel vertical migration,  
554 predator avoidance, location of food patches and mating are mechanisms by which the underlying  
555 zooplankton behavior present high spatial heterogeneity (Folt & Burns, 1999). Therefore, in  
556 Comau Fjord, those biological drivers together with oceanographic processes, such as estuarine

557 circulation, tidal mixing or water column stratification are considered to be the main drivers of  
558 zooplankton patchiness.

559

## 560 **Conclusions**

561 The seasonal changes of zooplankton across the entire water column showed that abundance  
562 peaked in spring, likely due to spring phytoplankton blooms. In summer, biovolume and biomass  
563 were the highest and decreased thereafter over time, reaching the lowest values in late autumn and  
564 mid-winter. Probably, the low concentration of Chl-*a* during the cold seasons could not support  
565 secondary production. The vertical distribution of zooplankton biovolume and biomass differed  
566 between day and night, with a daytime maximum in the 100-200 m water depth and a nighttime  
567 maximum in surface waters (0-50 m) associated with the diel vertical migration of the calanoid  
568 copepod Metridinidae. Overall, copepods were the dominant group of the total zooplankton  
569 community with an important contribution of small organisms (individuals <1.5 mm), followed  
570 by mysids, chaetognaths and cnidarians (biovolume and biomass), and nauplii and Appendicularia  
571 (abundance). The integrated abundance, biovolume and biomass also showed significant  
572 differences between daytime and nighttime values. These differences were probably due to the  
573 high zooplankton patchiness driven by both biological and oceanographic processes. Diel vertical  
574 migration, predation avoidance, location of food patches as well as estuarine circulation, tidal  
575 mixing or water column stratification are considered to be the main drivers of the zooplankton  
576 distribution in Comau Fjord.

577

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584

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