

To start, I can tell that the authors have made an effort to address my and the other reviewer's comments. The paper also has improved in terms of readability, and figures 1, 2 and 3 are much better now.

But there remain a few issues that prevent publication, and therefore I again need to ask for a thorough revision. I still consider this an interesting dataset, and I know how much work went into the image analysis.

The first point is again the biomass estimation. While the figure the authors provided in response to my previous comment looks convincing (then again, it is a log scale), the way they got to their values is absolutely NOT correct, and since a paper like this will be read a lot and table S3 will be used by others, I absolutely insist that this is corrected. I still think it would be easiest to ditch all the four steps of ellipsoid biovolume calculation, wet weight and dry weight assumptions, and finally carbon conversions, and simply use the image area to carbon conversion regressions (sources for which are already halfway compiled in their own table S3).

In table S3, conversion factors from dry weight to carbon are listed. There are several errors in this table. First, citing is sloppy, making it quite difficult to track down the errors (apart from the fact that the full references are missing). For example, Marcolin et al. (2015) is cited a lot, but their values mostly originate from Lehette & Hernandez-Leon 2009.

More importantly, THESE VALUES ARE NOT THE FRACTION OF DRY WEIGHT THAT IS MADE UP OF CARBON!!!

Instead, these values are the parameter a (intercept) from the regression $DW = a * Image\ Area^b$. I suspect this mistake somehow happened because many values are near 0.4. However, one could have noted that a 99% carbon content in ostracods would mean they are basically charcoal.

Table 1 from Marcolin et al. 2015. Regression equations used to estimate carbon from size (image area). Red box marks the dry weight to carbon conversion factors. Green box marks the regression parameters from Lehette & Hernandez-Leon 2006.

Foraminiferida	$V \times 89$		Michaels <i>et al.</i> (1995)
Bryozoa	$43.38 \times A^{1.54}$	0.432	Uye (1982) and Lehette and Hernández-León (2006)
Echinodermata	$43.38 \times A^{1.54}$	0.432	
Copepoda	$45.25 \times A^{1.59}$	0.455	
Cladocera	$43.97 \times A^{1.52}$	0.481	
Polychaeta	$43.38 \times A^{1.54}$	0.299	
Ostracoda	$99.46 \times A^{1.28}$	0.369	
Mollusca	$43.38 \times A^{1.54}$	0.227	
Peracarida	$49.58 \times A^{1.48}$	0.407	Lehette and Hernández-León (2006) and Davis and Wiebe (1985)
Chaetognatha	$23.45 \times A^{1.19}$	0.283	
Decapoda and Euphausiacea	$49.58 \times A^{1.48}$	0.407	Lehette and Hernández-León (2006), Beers (1966)
Siphonophora	$43.17 \times A^{1.02}$	0.109	
Other Cnidaria	$43.17 \times A^{1.02}$	0.072	Lehette and Hernández-León (2006)
Thaliacea	$0.3396 \times A^{2.642}$		Gibson and Paffenhöfer (2002)
Appendicularia	$2.62 \times 10^{-8} \times L_{TR}^{2.83}$		Sato <i>et al.</i> (2001)

A: area (mm²); V (cm³): volume defined as $4/3 \times \pi \times (\text{feret length}/2)^3$; L_{TR}: trunk length (μm).

To recapitulate: in the first round of review (where the calculation methods were not described yet, so I could not find the error), I suggested to do this:

"Area_exc" = Pixel Area excluding holes



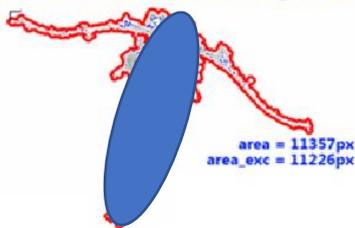
Use regression $DW(\mu g) = a * A^b$
 Where A is image area (mm²)
 And a and b are regression parameters from published literature (mainly L&H 2009)

Use published conversion factors for dry weight to carbon (mainly Kiorboe 2013)

Instead, the authors insisted that "Lehette and Hernandez-Leon (2009) provide estimated regression equations for different zooplankton taxa between scanned area and biomass (C content)((COMMENT: THIS IS INCORRECT, THEY PROVIDE DW FROM SCANNED AREA, NOT C CONTENT)). This is a direct relationship between size and C content which is actually the key parameter needed for estimates of migrant biomass and flux of C. We do not question such relationships by Lahette and Hernandez-Leon, but they are not suitable for our taxonomic composition and do not account for variable C-specific contents (which are independent of size). Instead, we used taxa-specific size and C content conversion factors previous published works which are suitable for our taxonomic groups and for the study region."

And they did this:

"Area_exc" = Pixel Area excluding holes



Calculate volume of rotating ellipsoid (blue)

Assume density of 1 (1mm³ = 1mgWW)

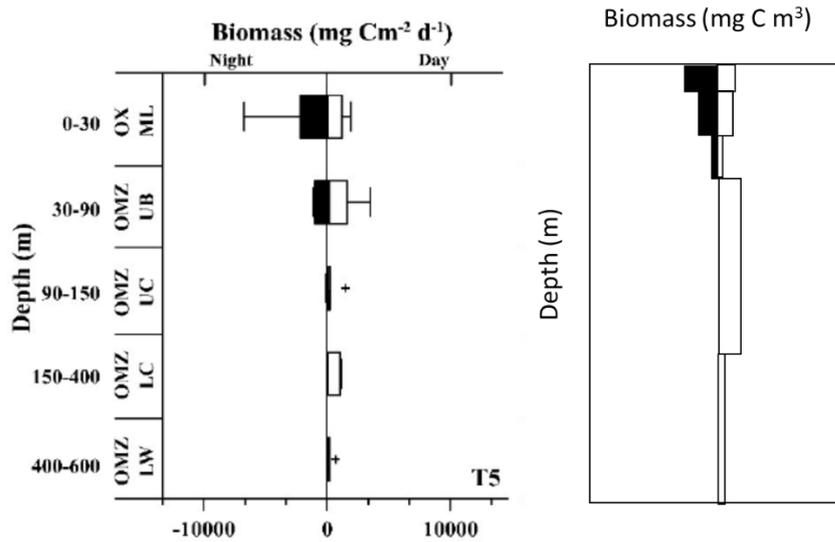
Assume water content of 90%
 1mg WW = 0.1mg DW

Use completely random numbers for carbon conversion mgDW->mgC

This needs to be corrected.

Second, the use of a log-axis on a stacked bar chart (figs 4 and 5) is not possible. Please use a linear axis. Actually I don't think the stacked bar chart are really necessary, the taxon-specific values are listed in the tables.

Third, visualizing the total distribution of biomass (Fig 6) should be kept, but this figure has some issues – first, the x-axis label is a rate. This also does not fit to the figure caption. I still think the most logical way to plot biomass distribution is as bar chart with variable widths on a linear depth axis. This way, you can plot biomass concentration (mg C m⁻³) as the height of the bars, and the bar area gives a good idea on the integral, which makes it nicer to compare depth strata with different thickness (and also relate to the hydrography in fig 2). I made a sketch to illustrate what I mean (it is just a sketch, so it lacks axis tick marks, error bars, and layer names). Left: one panel of Fig 6 now, right: suggestion.



Lastly, the results are still quite lengthy. The words “show” and “exhibit” are used both about fifteen times each. Try to streamline the writing a bit, using mostly active voice ("Copepod abundance was higher here than there" rather than "Copepods exhibited higher abundances here than there").

I do look forward to the final version of the paper.

1 Zooplankton diel vertical migration and downward C flux into the 2 Oxygen Minimum Zone in the highly productive upwelling region off 3 Northern Chile

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10 **Abstract.** The daily vertical movement of zooplankton, known as diel vertical migration (DVM), can enhance the vertical
11 flux of carbon (C) and so contribute to the functioning of the biological pump in the ocean. The magnitude and efficiency of
12 this active transport of C may depend on the size and taxonomic structure of the migrant zooplankton. However, the impact
13 that a variable community structure can have on zooplankton-mediated downward C flux has not been properly addressed.
14 This taxonomic effect may become critically important in highly productive eastern boundary upwelling systems (EBUS),
15 where zooplankton biomass becomes aggregated in the coastal zone and composed by a highly diverse community with
16 variable DVM behavior. In these systems, presence of a subsurface oxygen minimum zone (OMZ) can impose an additional
17 constraint to vertical migration and so influence the downward C export. Here, we address these issues based on a high-
18 resolution zooplankton sampling at three stations off northern Chile (20°S-30°S) during November-December 2015.
19 Automated analysis of zooplankton composition and taxa-structured biomass allowed us to estimate daily migrant biomass
20 by taxa and their amplitude of migration. We found that a higher biomass aggregates above the oxycline, associated with
21 more oxygenated surface waters and this was more evident upon a more intense OMZ. Some taxonomic groups, however,
22 were found closely associated with the OMZ. Most taxa were able to perform DVM in the upwelling zone withstanding
23 severe hypoxia. Also, strong migrants, such as Copepods Eucalanidae and Euphausiids, can exhibit a large migration
24 amplitude (~500 m), remaining either temporarily or permanently during the day or night condition within the core of the
25 OMZ and so contributing to the release of C below the thermocline. Our estimates of DVM-mediated C flux suggested that a
26 mean migrant biomass of $3.4 \text{ g C m}^{-2} \text{ d}^{-1}$ may contribute with about $0.6 \text{ g C m}^{-2} \text{ d}^{-1}$ to the OMZ system through respiration,
27 mortality, and production of fecal pellets at depth, accounting for ca. 25% of the net primary production, and so implying the
28 existence of a very efficient mechanism to incorporate freshly produced C into the OMZ. This downward C flux mediated by
29 zooplankton is however strongly dependent on the taxonomic structure due to variable migration amplitude and DVM
30 behavior.

31 1 Introduction

32 The Oxygen Minimum Zone (OMZ) in the Southeast Pacific, the fourth largest of the six permanent hypoxic regions in the
33 world oceans (Paulmier *et al.*, 2006), is a key component of the water column and a permanent feature intruding the coastal
34 zone of Ecuador, Peru and Chile (Fuenzalida *et al.*, 2009; Paulmier *et al.*, 2006). In the highly productive upwelling region
35 of Northern Chile the OMZ is closely linked to wind-driven upwelling in the coastal area and associated to the Equatorial
36 Subsurface Water (ESSW), which is transported southward along the continental shelf by the Peru-Chile Undercurrent
37 (PUC), as far south as 48°S (Fuenzalida *et al.*, 2009; Morales *et al.*, 1996a; Silva and Neshyba, 1979). Off Iquique (20°S) the
38 OMZ is characterized by being thick (500 m), very intense ($< 20 \mu\text{mol kg}^{-1}$) and with O_2 concentrations in the core of OMZ
39 among the lowest found in the global ocean reaching the current detection limit ($< 1 \mu\text{M}$) (Ulloa and Pantoja, 2009), although
40 it becomes thinner at about 30°S (Paulmier *et al.*, 2006).

41 During the last decades, the OMZ systems have attracted much scientific interest because of evidence showing that hypoxic
42 and anoxic conditions in coastal areas are expanding and becoming more intense (Ekau *et al.*, 2010; Stramma *et al.*, 2008).
43 At present, ongoing ocean deoxygenation is widely recognized as linked to global warming and it is rising much concern in
44 modern oceanography (Breitburg *et al.*, 2018).

45 The presence of oxygen-depleted water becomes a critical physiological constraint for pelagic and benthic organisms
46 inhabiting the upwelling zone, impacting their biomass and productivity, the species diversity, distribution and behaviour
47 (Wishner *et al.*, 2018; Ekau *et al.*, 2010; Grantham *et al.*, 2004). For instance, diel vertical migration (DVM), a common
48 feature of the various size groups of zooplankton and also one of the most important movements of biomass in the ocean, can
49 also be affected by changes in intensity and distribution of the OMZ (Wishner *et al.*, 2018, 2013; Escribano *et al.*, 2009;
50 Fernández-Álamo and Färber-Lorda, 2006; Hidalgo *et al.*, 2005; Morales *et al.*, 1996; Judkins, 1980). The OMZ can act as
51 an ecological barrier for vertical distribution of many organisms, constraining most zooplankton to a narrow (50 m) upper
52 layer, as shown in the coastal upwelling zone off Chile according to the works of Escribano (2006) and Donoso and
53 Escribano (2014). Zooplankton also become limited to the upper 150 or 300 m in the eastern tropical north Pacific (Wishner
54 *et al.*, 2013). However, the OMZ can also offer refuge for species adapted to live there, creating microhabitats of differing
55 oxygen concentration that are characterized by layers of high zooplankton biomass and abundance, with distinct species
56 zonation (Antezana, 2009; Wishner *et al.*, 2008; Fernández-Álamo and Färber-Lorda, 2006), which, in turn, may have
57 important consequences for carbon (C) cycling and its vertical flux. For example, it is known that zooplankton in the coastal
58 upwelling region off northern Chile may play a significant biogeochemical role by promoting carbon flux into the subsurface
59 OMZ (Escribano *et al.*, 2009). Therefore a significant proportion of the vertical material flux from the euphotic zone to the
60 deep sea ($> 200 \text{ m}$) and within the food chain could be determined by DVM of zooplankton (Longhurst and Williams, 1992;
61 Steinberg and Landry, 2017).

62 As important contributors to the functioning of the biological pump, diel zooplankton migrants can actively increase the
63 magnitude of C export by transporting surface-ingested material in their guts to deep waters where it can be metabolized

64 (Steinberg and Landry, 2017). Zooplankton moults or mortality at depth can also contribute to the transportation of
65 assimilated organic biomass into the deep waters (Ducklow *et al.*, 2001). The biological pump process is also thought to be
66 related to the size structure of dominant zooplankton. This because some groups with large body sizes may exhibit a greater
67 range of vertical migration and sometimes higher levels of biomass, and so influencing the biogeochemical fluxes (Dai *et al.*,
68 2016; Ducklow *et al.*, 2001). However, the effect of variable size structure on DVM performance and its consequence for
69 active C transport has not indeed been assessed. Size-structure is certainly related to zooplankton composition which has
70 hardly been properly addressed when examining the role of DVM on C flux. For instance, in areas with hypoxic subsurface
71 layers some species are more active migrants and thus more efficient C-transporters, because they have developed
72 adaptations to low oxygen conditions and can even use the OMZ as their habitat, either temporarily or permanently
73 (Escribano *et al.*, 2009; Gonzalez and Quiñones, 2002; Seibel, 2011).

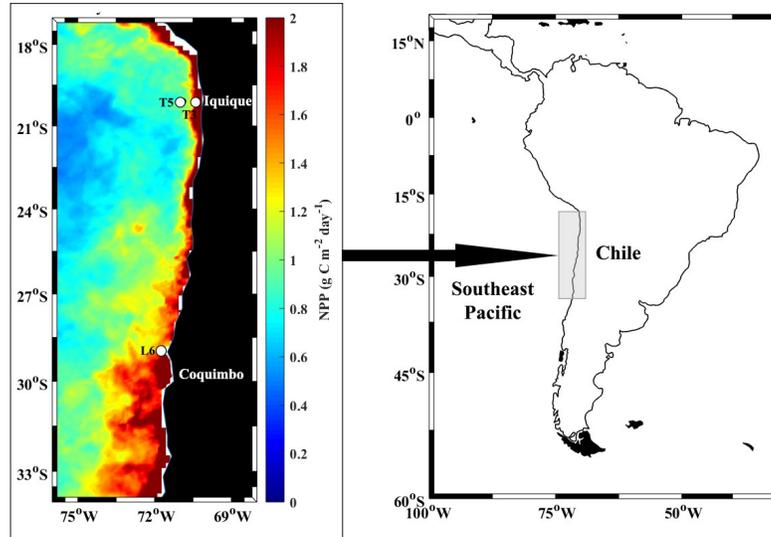
74 C-export to depth may also depend on the amount of biomass being produced in the photic zone. Primary production
75 promotes zooplankton feeding and growth and therefore determining C availability for both passive and active transport to
76 depth. In this context, highly productive upwelling zones can be assumed as systems where the C-flux mediated by
77 zooplankton DVM can be enhanced, although certainly depending on the size and taxonomic structure of zooplankton. In
78 these regions, a shallow OMZ might exert a further impact on the C-flux by affecting DVM or zooplankton metabolism at
79 depth. In the present study, based on high resolution sampling and automated analysis of mesozooplankton, we assessed
80 zooplankton vertical migration and downward C to the OMZ in the highly productive upwelling region of Northern Chile.
81 We aimed at understanding the role that taxonomic and size-structure can play in the magnitude and variability of the DVM
82 behaviour interacting with a shallow OMZ, and the implications this interaction can have on the magnitude of the downward
83 C flux in a highly productive coastal upwelling zone

84 **2 Methods**

85 **2.1 Study Area**

86 The study area was located in the Southeast Pacific Ocean and covered the coastal zone of the northern upwelling region of
87 Chile (21-29°S) (**Fig. 1**), which is a region known to be subjected to wind-driven upwelling throughout the year and
88 containing an intense and shallow OMZ (Ulloa *et al.*, 2012). The sampling design comprised three stations: Two stations (St.
89 T3 and St. T5) across a zonal section off Iquique (20°S) and a coastal station (St. L6) off Coquimbo (29°S). The study was
90 carried out during the LowpHox cruise conducted in November-December 2015 onboard the R/V *Cabo de Hornos*. At each
91 station, temperature, salinity and dissolved oxygen (DO) were recorded from 1000 m (St.T5 and St.T3) and only 356 m (St.
92 L6) using an oceanographic rosette with a CTD SeaBird 911 (SBE 911 plus) equipped with a SeaBird SBE-43 oxygen
93 sensor and a Sea tech Fluorometer. Discrete water samples were also obtained for chemical measurements of oxygen.

94
95



98
99 **Fig. 1.** Study area at the northern upwelling region of Chile, indicating sampling stations during the LowpHox cruise
100 conducted in November-December 2015 onboard the R/V *Cabo de Hornos*. Satellite estimated Net Primary Production
101 (NPP), averaged for November-December 2015 is shown. NPP was obtained from
102 <http://science.oregonstate.edu.ocean.productivity>.

103 2.2 Zooplankton sampling

104 Zooplankton samples were collected during daytime and night-time conditions in two consecutive days at the three stations
105 off northern Chile (T5-T3-L6) (**Fig. 1**), also as indicated in **Table S1 (Supplemental Material)**. Zooplankton hauls were
106 performed from 600 to 0 m depth with a Multi Plankton Sampler Hydro-Bios Multinet system with a 0.25 m² opening area
107 and equipped with 200- μ m mesh-size nets. The Multinet towing speed was 1 m s⁻¹ and the flowmeter was in the mouth of
108 the Multinet to estimate the volume of filtered water. At each station, 40 discrete samples were obtained from 600 to 0 m
109 depth. Each sample corresponded to a different depth stratum (30-0, 90-30, 150-90, 400-150 and 600-400m depth). These
110 strata were defined in according to distribution of oxygen concentration and localization of the OMZ. Therefore, they were
111 assumed as representing the near surface oxygenated layer, the upper oxycline, the upper OMZ, the OMZ core and the lower
112 boundary of the OMZ indicated in **Table S2 (Supplemental Material)**. Once onboard the collected zooplankton samples
113 were preserved immediately in 5% buffered formalin-seawater solution.

114

115 2.3 Taxonomic and size measurements

116 Taxonomic identification and enumeration of taxa were carried out by analysis of digitized images obtained with the
117 Hydroptic ZooScan digital imaging system (Gorsky *et al.*, 2010). Each sample was wet-sieved through a 1000 μm mesh into
118 two size fractions. Then, each size-fraction was fractionated again separately with a Motoda splitter until the zooplankton
119 concentration was sufficiently diluted to avoid contact between organisms in the ZooScan scanning frame. Fractioning into
120 small and large organisms and consequent separate image acquisition of the two size classes prevented underestimates of
121 large-rare objects which may need less fractioning (Gorsky *et al.*, 2010). 179 subsamples were scanned and digitized at a
122 resolution of 2400 dpi after manual separation of objects on the scanning tray. After processing the samples with ZooProcess
123 software, each of the objects was automatically sorted with the help of a learning set, and then the sorting was visually
124 validated by an expert (for details, see Chang *et al.*, 2012; Gorsky *et al.*, 2010). Organisms making up the ZooScan datasets
125 were enumerated, measured, biomass-estimated and classified into 27 taxonomic groups, such as Copepods, Chaetognaths,
126 Euphausiid, Gelatinous and other zooplankton. The abundance (ind. m^{-3}) and stratum-integrated abundance (ind. m^{-2}) of
127 total zooplankton or of each taxonomic group was calculated following Eq. (1):

128

$$129 \text{ Abundance}/\text{m}^3 = \frac{N * \text{Subpart}}{\text{Vol} * \text{Townb}}, \quad (1)$$

130

131 where, N is the number of individuals with same prediction (e.g., in last column written “copepod”), Subpart is the splitting
132 ratio, Vol is net volume and Townb is the number of net tows in a sample. Stratum-integrated abundance (ind. m^{-2}) was
133 obtained after multiplying by width (m) of a given stratum.

134

135 2.4 Biomass estimates and Carbon fluxes

136 The ZooScan Integrated System also provided zooplankton body size for each organism. The ellipsoidal biovolume (EBv)
137 was computed instead of equivalent spherical diameter (ESD). Biovolume was used because most of the zooplankton shapes
138 were elongated, and also because ESD overestimates biovolume since spheres have a higher volume: cross-sectional-area
139 ratio than other shapes (García-Comas *et al.*, 2014; Herman and Harvey, 2006; Sprules *et al.*, 1998). Ellipsoidal biovolume
140 EBv (mm^3) was calculated following Eq. (2):

141

$$142 \text{ EBv} = \frac{4}{3} * \pi \left(\frac{\text{Major}}{2} \right) * \left(\frac{\text{Minor}}{2} \right)^2, \quad (2)$$

143

144 where, the EBv corresponds to the biovolume of an ellipsoid, Major is the longest axis of the object and Minor is the minor
145 axis of a perfect ellipse of the same area of the measured object, both axes are in mm and were provided by ZooScan. We
146 used EBv (mm^3) for estimating biomass (mg C) for all taxonomic groups. Conversions from the EBv to carbon were made in

147 three steps: First EBv was converted into wet weight (WW), assuming 1 mm⁻³ was equivalent to 1 mg WW (Wiebe *et al.*,
148 1975). Then WW was converted to dry weight (DW) assuming that the water content of zooplankton in average was 90%
149 (DM = 0.1 × WW), as suggested by Yamaguchi *et al.* (2014) and Sato *et al.*(2015). Finally, individual biomass was
150 converted to carbon using a factor carbon for each taxonomic group from previous published data **Table S3 (Supplemental**
151 **Material)**.

152 To calculate the migrant biomass, we integrated biomass in the upper 90 m layer from our two sampled strata 0-30 m and
153 30-60 m. This 0-90 m stratum was considered the approximate above-oxycline layer after examining the vertical profiles of
154 oxygen. Biomass at night was thus subtracted from the corresponding day biomass in this layer to assess daily changes
155 involving migrants as in Putzeys *et al.*(2011). Thus, the negative values of the day-minus-night biomass corresponded to
156 migrant biomass that reached the epipelagic layer at night including organisms inhabiting above and below the oxycline.

157 The proportion of migrant biomass with respect to observed biomass in the upper 90 m of a given taxonomic group was
158 defined at the rate of migration on a daily basis. This rate of migration could thus be used as an index of DMV behaviour for
159 a taxonomic group.

160 To estimate the active C flux at each sampling station, we used a daily respiration fraction of 0.12. This proportion of
161 biomass being respired was estimated by Hernández-León and Ikeda (2005) for total zooplankton biomass at a temperature
162 ranging between 13°C and 18 °C for mid-latitude areas. We used this value considering a similar temperature range between
163 surface water and the core of the OMZ (**Fig. 2**). The daily contribution of egestion rate to C flux was assumed to be 0.09 of
164 migrant biomass, as suggested by Escribano *et al.* (2009). This estimate was derived from a combined biomass of large-sized
165 copepods and euphausiids which are the major groups contributing to migrant biomass in this upwelling area. This value also
166 considers that most egested faecal pellets may be produced rapidly in the upper layer upon feeding (Escribano *et al.*, 2009).
167 The contribution of mortality to C flux was assumed to be 0.08 d⁻¹ of migrant biomass, as a conservative estimate suggested
168 by Ohman and Wood (1996). We thus estimated total C flux following Eq. (3):

169

$$170 \text{ C Flux} = \left(\frac{RZ + MZ}{2} \right) + EZ \quad (3)$$

171

172 where Rz, Mz, and Ez, are Respiration, Mortality and Egestion estimates of fractions of total migrant biomass (MB) for a 12-
173 h period. We thus estimated total flux as,

174

$$175 \text{ C Flux} = (0.12*(MB) + 0.08*(MB)) / 2 + 0.09*(MB) \quad (4)$$

176 **2.5 Patterns of vertical distribution of migrating zooplankton**

177 For the analysis of vertical distribution of organisms, the density estimates of the organisms were standardized to number of
178 individuals per m³ (for each stratum) or per m² (for integrated values). In order to quantify the presence and extent of DVM

179 of various taxa at each station, we calculated both night: day (N:D) ratios in each stratum and weighted mean depth (WMD)
180 for zooplankton abundance, as a measure of the center of gravity of a population's vertical distribution for each taxon and
181 haul, in according to Andersen *et al.* (2004) following Eq. (5):

182

$$183 \quad WMD = \frac{\sum(ni*zi*di)}{\sum(ni*zi)}, \quad (5)$$

184

185 where, d is the mean depth of the strata (m), z the width (m) of the strata and ni the abundance (ind. m⁻³) of a given i
186 taxonomic group.

187 We calculated the amplitude of vertical migration (Δ DVM) as the difference between the WMD of the organisms during the
188 day and the night and therefore this Δ DVM was considered as the criterion to assess the DVM behaviour for each taxonomic
189 group. Positive values indicated normal DVM (pattern of nocturnal ascent by individuals that reside at depth by day) and
190 negative values indicated reverse DVM (pattern of nocturnal descent by individuals that reside near the surface by day). The
191 individuals that occupied the same depth stratum by day and by night, whether near the surface or at depth were considered
192 as non migrant in according to Ohman (1990).

193 2.6 Statistical analysis

194 For statistical analysis, as a criterion for determining if the DVM was significant, we tested for differences in the WMD
195 mean between day and night using a two-tailed t test. We considered the occurrence of DVM when the difference in the
196 WMD mean between day and night was significant ($p < 0.05$). In order to evaluate the similarity/dissimilarity in the
197 abundance and biomass among stations, strata, and day-night conditions, the multivariate grouping techniques were applied
198 ("cluster analysis"), ANOSIM (Two-Way Crossed Analysis) tests and multidimensional scaling (MDS) with the data
199 transformed in PRIMER v 6.1.16 (2013), prior to the application of the Bray-Curtis similarity index (Bray and Curtis, 1957).
200 In general, WMD for taxonomic groups did not exhibit a pronounced bimodal vertical distribution.

201 3 Results

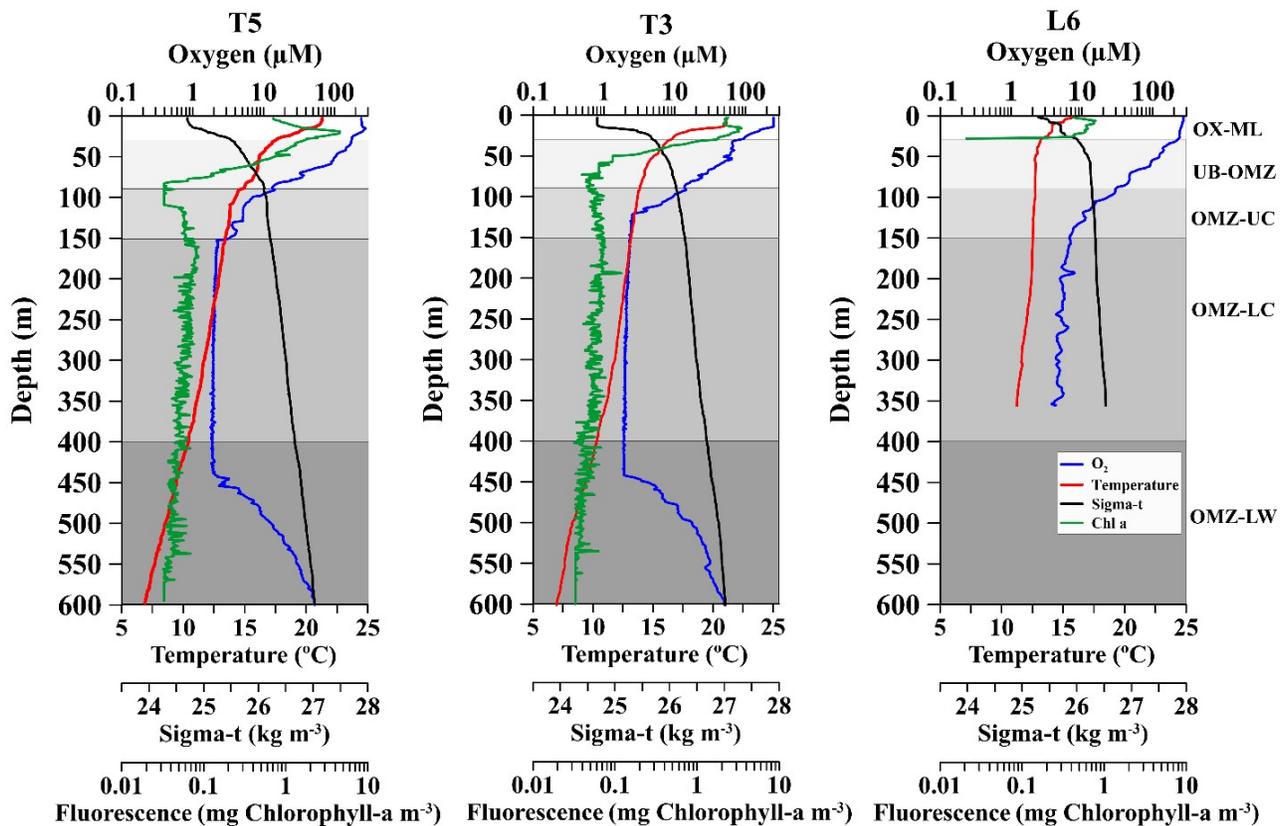
202 3.1 Hydrographic conditions

203 Vertical profiles of temperature, DO, and sigma-t show general oceanographic conditions in all stations (Fig. 2), such as the
204 vertical distribution of oxygen and location of the OMZ. From the vertical profiles of oxygen and coinciding with the
205 sampled layers of the Multinet different strata were defined: A well oxygenated stratum with oxygen approaching air
206 saturation ($>250 \mu\text{mol O}_2 \text{ kg}^{-1}$) defined as oxic mixed layer (OX-ML), an upper O₂ gradient (oxycline), defined by the level
207 where O₂ reaches 4% of the surface O₂ (Paulmier *et al.*, 2006), at whose base is located the upper boundary of the OMZ (45
208 $\mu\text{mol O}_2$ isoline, OMZ-UB) (Escribano *et al.*, 2009; Hidalgo *et al.*, 2005; Morales *et al.*, 1999); the OMZ core defined by an



209 upper boundary (OMZ-UC) with the lowest concentration of O_2 ($<20 \mu\text{mol } O_2 \text{ kg}^{-1}$) and a lower boundary (OMZ-LC) (1
 210 to $<20 \mu\text{mol } O_2 \text{ kg}^{-1}$), and finally a lower O_2 gradient (OMZ-LW). 

211 Across the zonal section off Iquique the offshore station (St. T5) and onshore station (St. T3) showed two contrasting
 212 hydrographic regimes regarding the OMZ. Station T5 had a less pronounced and thicker OMZ than station T3. At both
 213 stations the 5 strata were well defined in the water column (Fig. 2). The OX-ML ($>250 \mu\text{mol } O_2 \text{ kg}^{-1}$) was present at 18 m
 214 (St. T5) and 15 m (St. T3). The oxycline exhibited a gradual decrease from oxic ($\sim 250 \mu\text{mol } O_2 \text{ kg}^{-1}$) to suboxic ($<20 \mu\text{mol}$
 215 $O_2 \mu\text{mol } O_2 \text{ kg}^{-1}$) conditions associated with a strong stratification in the upper 80 m depth. The $45 \mu\text{mol } O_2$ isoline (OMZ -
 216 UB) was present at the base of the oxycline at 70 m (St. T5) and 59 m (St. T3). The OMZ core ($<20 \mu\text{mol } \mu\text{mol } O_2 \text{ kg}^{-1}$) was
 217 located below the thermocline and below the 26.5 kg m^{-3} isopycnal following description of Paulmier *et al.* (2006). In the
 218 oceanic station (St. T5) the OMZ core was located between 80 to 514 m, while in the coastal station (St. T3) it was between
 219 80 to 507 m with 423 m thickness. The O_2 concentration in the core was ca. $1 \mu\text{mol } O_2 \text{ kg}^{-1}$. The OMZ-LW at both stations
 220 was delimited above the core and below the depth where the O_2 slope changed significantly (slope break $>20 \mu\text{mol / m}$) (Fig.
 221 2).
 222



223
 224

225 **Fig. 2.** Vertical profiles of dissolved oxygen (O₂), temperature, Chlorophyll-a and water density (sigma-t), at 3 stations off
226 northern Chile (see **Fig. 1**) during the LowpHox Cruise in the austral spring 2015. Shaded areas represent different layers
227 defined according to oxygen concentration. OX-M=Oxic mixed layer, OMZ-UB= Upper boundary of the oxygen minimum
228 zone, OMZ-UC= Upper core of the oxygen minimum zone, OMZ-LC= Lower core of the OMZ and OMZ-LW=lower
229 oxygen gradient.

230 The structure of the OMZ at the coastal station (St.L6) off Coquimbo (29° S) (**Fig.1**), exhibited similar characteristics to St.
231 T3 (21° S), but in this area the OMZ was deeper and thinner. The OX-ML was shallower. The OMZ -UB (45 μmol O₂) in the
232 base of oxycline was down to 80 m. The low O₂ concentrations in the core were less intense than at 21°S (4 to 20 μmol O₂kg⁻¹)
233 and it was located below 100 m (**Fig. 2**). The OMZ-LW could not be assessed because of lack of CTD data below 350 m.
234 Additional oceanographic variables showed a surface warming (>20 °C) and strongly stratified conditions at the three
235 stations with a sharp thermocline in the upper 100 m, coinciding with the oxycline, whereas Chlorophyll-a maximum (>5 mg
236 Chlorophyll-a m⁻³ was located in the upper 20 m (**Fig. 2**).

237 **3.2 Zooplankton composition and abundance**

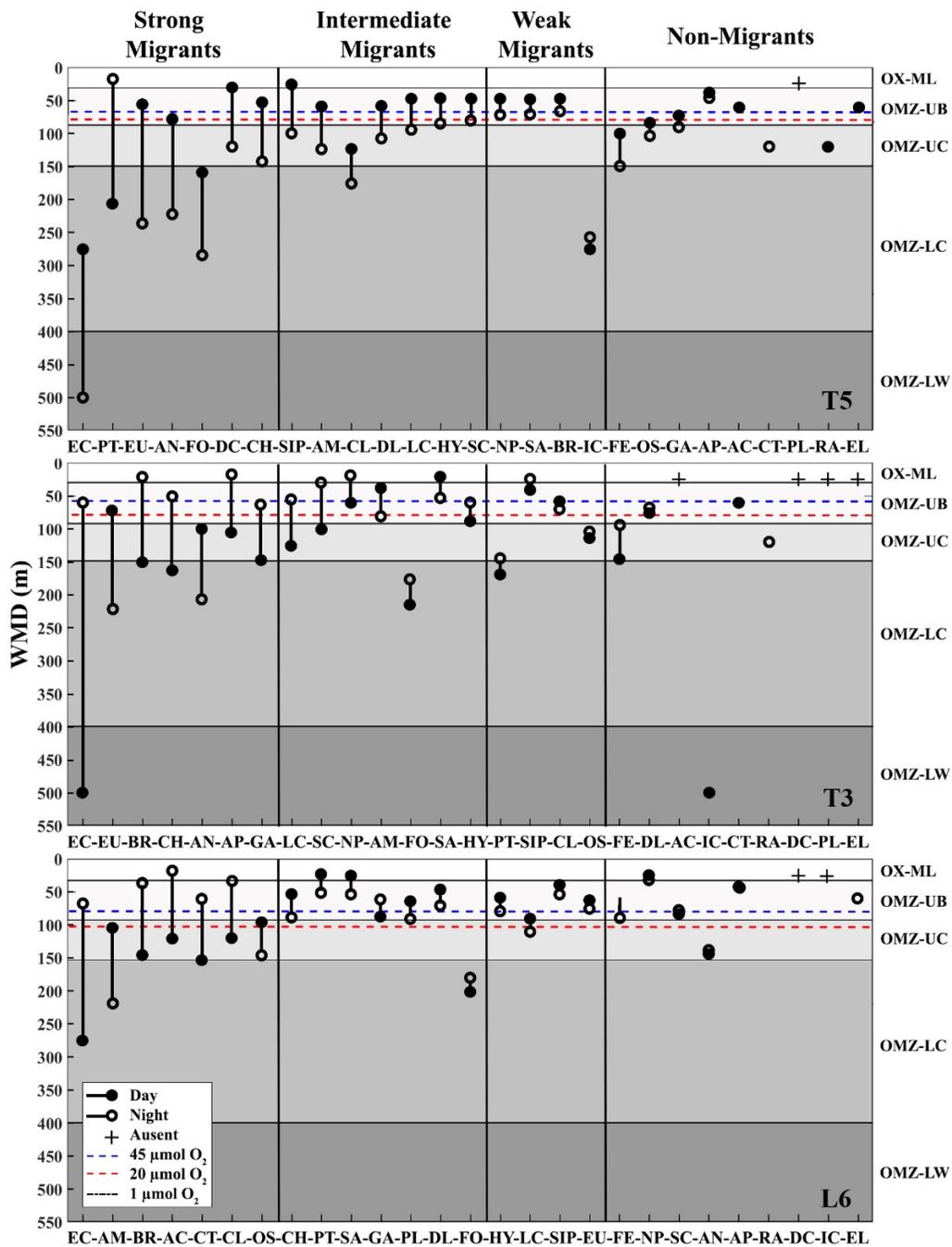
238 A total of 27 zooplankton taxa were identified by the ZooScan and ZooProcess (**Table S4 Supplemental Material**). The
239 number of taxa varied among stations and strata. Across the zonal section off Iquique the number of taxonomic groups
240 fluctuated between 23 (St. T3) and 26 (St. T5), whereas 25 taxa were present off 29° S (St. L6). The most dominant taxa
241 both daytime and night conditions, were: Copepods 87% (in St. T5), 79% (in St. T3) and 69% (in St. L6). This group was
242 constituted by small Copepods, large Copepods, the Copepods Eucalanidae and the Copepods *Acartia* spp; fish eggs
243 constituting 2% (in St. T5), 5% (in St. T3) and 6% (in St. L6); Nauplii being <1% (in St. T5), <1% (in St. T3) and 7% (in St.
244 L6); Appendicularian 5% (in St. T5), 4% (in St. T3) and 3% (in St. L6) (**Table S5 Supplemental Material**). The remaining
245 19 pooled groups only constituted <6% (in St. T5), 11% (in St. T3) and 15% (in St. L6). The total integrated abundances of
246 zooplankton (0-600 m) by sampling station are shown in **Table S4** (Supplemental Material). As based on a Two-Way
247 Crossed Analysis ANOSIM test, this water-column integrated abundance did not show significant differences between day
248 and night samples (p>0.05). Significant differences among stations (Two-Way Crossed Analysis ANOSIM p<0.05) were
249 found, so that the stations were treated independently. Off Iquique the abundance showed the lowest values at the onshore
250 station (St. T3 with 18% lower), which was characterized by the strongest and most extensive OMZ in the study area. These
251 values increased at the offshore station (St. T5 with 31% greater), where the OMZ was less pronounced and thicker. Unlike
252 stations T3 and T5, the onshore station off 29°S (St. L6) had a weaker and less extensive OMZ showing the highest
253 zooplankton abundance (51% greater).

254

255 3.3 Diel vertical migration (DVM) and vertical distribution

256 3.3.1 Main migrant groups of zooplankton

257 The diel vertical migration of 27 zooplankton taxa in the 0–600 m water column is shown in **Fig. 3**. These taxa were
258 classified into four groups according to their amplitude of migration (Δ DVM) (**Table 1**): **1) Strong migrants**, represented
259 by taxa that exhibited a strong DVM and showed a broad range of Δ DVM from 225 to 99 m (in St.T5), 440 to 84 m (in St.
260 T3) and 208 to 87 m (in St. L6). This group constituted 70% of taxa with higher Δ DVM. The composition of taxa in this
261 group was variable at each station (**Table 1**), but in general was well represented by Eucalanidae Copepods (EC),
262 Euphausiids (EU), *Acartia* Copepods (AC), Ctenophores (CT), Decapods (DC), Annelidae (AN), Bryozoan (BR), Pteropods
263 (PT) and Chaetognaths (CH). These taxa were mostly concentrated in the oxic surface stratum (**OX-ML**), and the **OMZ**
264 **core** showing a strong interaction with both the **OMZ-UC** and the **OMZ-LC**, and so changing from normoxia to hypoxia
265 condition and vice versa between 0 to 550 m (**Fig. 3**), **2) Intermediate migrants**, represented by taxa that exhibited a
266 moderate DVM and showed a range of Δ DVM from 73 to 34 m (in St. T5), 70 to 27 m (in St. T3) and 49 to 22 m (in St. L6).
267 This group constituted 23% of taxa with moderate Δ DVM. The composition of taxa in this group was also variable at each
268 station (**Table 1**), but it was mostly represented by small (SC) and large Copepods (LC), Amphipods (AM), Cirripedia
269 Larvae (CL), Gastropods (GA), Siphonophores (SIP) and Appendicularian (AP). These taxa were mostly concentrated in the
270 oxic surface strata (**OX-ML**) and in the **OMZ-UC** showing some interaction with the **OMZ core** and vertically changing
271 from normoxia to hypoxia condition, and vice versa between 0 to 200 m, **3) Weak migrants**, represented by taxa that
272 exhibited a weak DVM and showed a range of Δ DVM of 24 to 18 m (in St.T5), 23 to 12 m (in St.T3) and 21 to 11 m (in St.
273 L6). This group constituted 5% of taxa of low range of Δ DVM. The composition of taxa in this group was also variable at
274 each station (**Table 1**), but in general it was represented by Hydrozoa (HY), Salps (SA), Platyhelminthes (PT), Decapoda
275 Larvae (DL), Ostracods (OS), Nauplii (NL) and Ichthyoplankton (IC). These taxa were concentrated mainly in the oxic
276 surface strata (**OX-ML**) and in the **OMZ-UP**, but also in the **OMZ-UC** at the onshore stations (StationT3 and Station L6),
277 showing much less interaction with the **OMZ core**, while spatially moving from normoxia to hypoxia condition and vice
278 versa between 0 to 100 m, and **4) Non-migrants**, represented by taxa which did not exhibit a significant DVM and had a
279 range of Δ DVM from 16 to 0 m (in St. T5), 7 to 0 m (in St. T3) and 6 to 0 m (in St. L6). This group constituted 1% of taxa
280 with not significant Δ DVM. The composition of taxa in this group was also variable at each station (**Table 1**), but in general
281 it was represented by fish eggs (FE), Radiolarian (RA) and Echinoderm larvae (EL).



282

283

284 **Fig. 3.** Weighed mean depth distribution (WMD) of the zooplankton community interacting with the OMZ off Iquique
 285 (Stations T5 and T3) and off Coquimbo (Station L6) at the northern upwelling area of Chile during the austral spring 2015.

286 Shaded gray areas represent different layers defined by their oxygen levels (defined in Methods). The taxonomic groups

287 were classified by automated analysis (ZooScan): EC= Eucalanidae Copepods, AM= amphipods, BR=Bryzoan, AC= *Acartia*
 288 Copepods, CT= Ctenophores, CL=Cirripedia larvae, OS=Ostracods, CH=Chaetognaths, PT= Pteropods, SA=salps, GA=
 289 Gastropods, PL= Platyhelminthes, DL= Decapoda larvae, FO=Foraminifera HY=Hydrozoa, LC=Large Copepods,
 290 SIP=Siphonophores, EU=Euphausiids, FE=fish eggs, NP=Nauplii, SC=Small copepods, AN= Annelidae,
 291 AP=Appendicularian, RA=Radiolarian, DC=Decapods, IC=ichthyoplankton, EL=Echinoderm larvae.

292

293 **Table 1-** Diel vertical migration indices for 27 taxonomic groups (TAXA) identified and sorted by ZooScan at 3 stations off
 294 northern Chile (see **Fig. 1** for acronyms), during the austral spring 2015. Amplitude of migration (Δ DVM) is in meters.
 295 Positive values indicate normal DVM and negative values indicate reverse DVM (see Methods). N: D is Night: day ratio of
 296 abundance. Four groups are defined in according to DVM behaviour. Relative abundances are shown in %.

T5				T3				L6				MIGRANTS
TAXA	Δ DVM	N:D	%	TAXA	Δ DVM	N:D	%	TAXA	Δ DVM	N:D	%	
EC.	225	0.6	14	EC.	-440	8.3	28	EC.	-208	4.1	20	STRONG MIGRANTS > 5%
PT.	-188	12.4	12	EU.	149	0.3	10	AM.	115	0.5	11	
EU.	181	0.2	11	BR.	-129	7.0	8	BR.	-107	3.8	10	
AN.	145	0.3	9	CH.	-114	3.2	7	AC.	-103	7.0	10	
FO.	126	0.6	8	AN.	105	0.5	7	CT.	-94	2.6	9	
DC.	90	0.3	6	AP.	-88	5.9	6	CL.	-87	3.6	8	
CH.	88	0.4	6	GA.	-84	2.3	5	OS.	49	0.7	5	
SIP.	73	0.3	5	LC.	-70	2.3	5	CH.	35	0.6	3	INTERMEDIATE MIGRANTS > 2%
AM.	64	0.5	4	SC.	-70	3.3	5	PT.	29	0.4	3	
CL.	51	0.7	3	NP.	-42	1.6	3	SA.	29	0.5	3	
DL.	50	0.5	3	AM.	42	3.3	3	GA.	-27	1.4	3	
LC.	47	0.7	3	FO.	-38	0.5	2	PL.	26	0.7	2	
HY.	39	0.5	2	SA.	32	1.2	2	DL.	22	0.7	2	
SC.	34	0.5	2	HY.	-27	0.4	2	FO.	-22	0.7	2	
NP.	24	0.6	2	PT.	-23	1.5	1	HY.	21	1.1	2	WEAK MIGRANTS > 1%
SA.	23	0.7	1	SIP.	-15	1.2	1	LC.	19	0.7	2	
BR.	18	0.7	1	CL.	12	1.6	1	SIP.	15	0.8	1	
IC.	-18	0.7	1	OS.	-9	0.8	1	EU.	11	0.7	1	
FE.	49	1.1	3	FE.	-52	1.1	3	FE.	25	0.9	2	NON- MIGRANTS <1%
OS.	16	0.8	1	DL.	-7	1.1	0	NP.	6	0.8	1	
GA.	15	0.8	1	AC.	-	-	-	SC.	-4	1.1	0	
AP.	7	0.8	0	IC.	-	-	-	AN.	-4	1.0	0	
AC.	-	-	-	CT.	-	-	-	AP.	2	1.0	0	
CT.	-	-	-	RA.	-	-	-	RA.	-	-	-	
PL.	-	-	-	DC.	-	-	-	DC.	-	-	-	

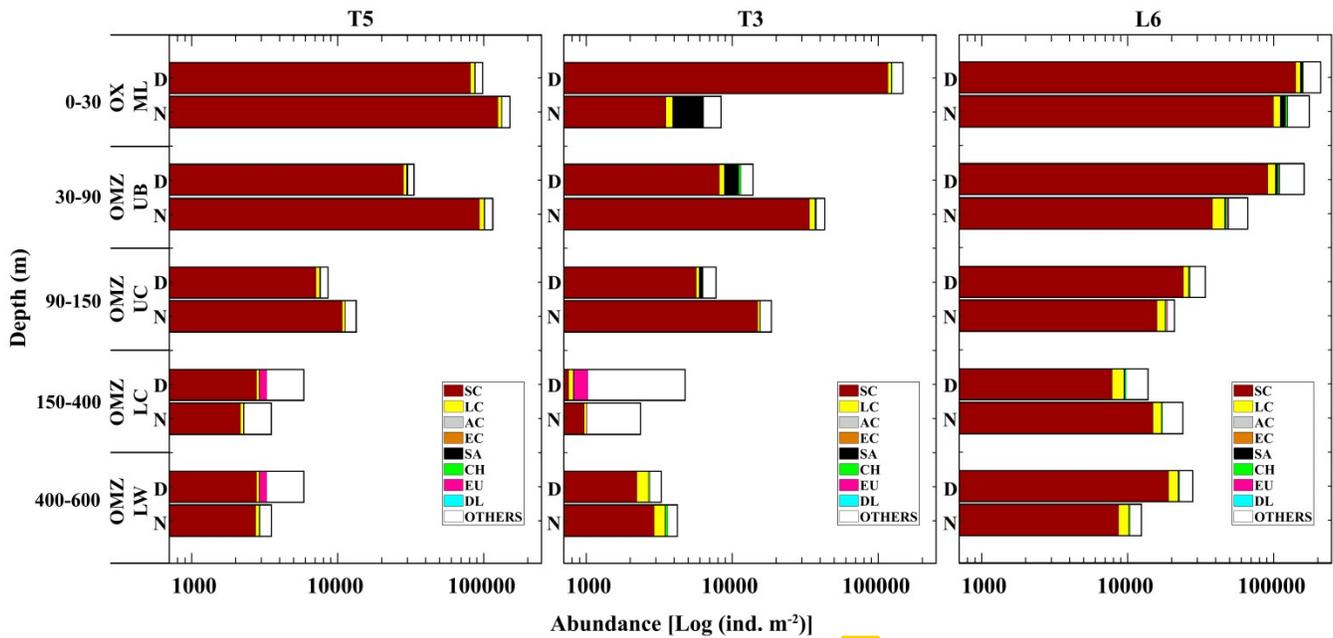
RA.	-	-	-	PL.	-	-	-	IC.	-	-	-	
EL.	-	-	-	EL.	-	-	-	EL.	-	-	-	

297

298 **3.2 Vertical Distribution and DVM of dominant groups.**

299 Vertical distributions of zooplankton were assessed for 5 taxonomic groups, which represented 82% of total abundance in
300 average: Salps (SA.); Copepods represented by small Copepods (SC.), Large Copepods (LC.), Eucalanidae Copepods (EC)
301 and *Acartia* Copepods (AC); Chaetognaths (CH); Euphausiids (EU) and Decapods Larvae (DL); as well as their patterns of
302 strata–station–abundance relationships are illustrated in **Fig. 4**. The abundance of these zooplankton groups regarding depth
303 strata was significantly different (ANOSIM, $p < 0.05$) at each station, and therefore representing distinctive microhabitats
304 characterized by specific depth and oxygen concentration. In general, the higher abundance (>50%) was found in the
305 shallower strata and well oxygenated layers (**OX-ML and OMZ UP**) (>250 $\mu\text{mol O}_2 \text{kg}^{-1}$), and then it decreased rapidly in
306 the strata associated with the OMZ core (**OMZ-UC and OMZ-LC**). Below this stratum a slight peak in abundance was
307 present in the **OMZ-LW** at all stations, occurring between 400-600 m, both daytime and night conditions (**Fig. 4**).

308



309

310 **Fig. 4.** Vertical distribution of dominant taxonomic groups of zooplankton sampled at 3 stations off northern Chile: off
311 Iquique at 21° S (Stations T5 and T3) and off Coquimbo at 29° S (Station L6). Abundances are mean values from samples
312 obtained during 2 consecutive days under night and day conditions in the austral spring 2015. Sampled layers represent

313 different conditions in terms of oxygen levels (see Methods) and the taxonomic groups and their acronyms are detailed in
314 Supplemental Material **Table S3**.

315 As expected, Copepods dominated numerically the zooplankton community both within and outside the OMZ (**Fig. 4**). Small
316 Copepods (SC) were the most abundant, followed by large Copepods (LC), whereas the Copepods *Acartia* (AC) and
317 Eucalanidae (EC) showed the lowest abundances among Copepods. The largest aggregation of Copepods (pooled data)
318 altogether during the entire study period was observed at the offshore station St. T5 (87%), where abundances reached
319 192088 ind. m⁻². At onshore station St. T3 percentage of contribution of copepods was 79% and 69% at the St L6 (**Table S5**
320 **Supplemental Material**). Off Iquique, the highest abundance and biomass were in the shallower strata (**OX-ML**) at St.T5
321 (46%) and at St. T3 (47%), and these values were reduced in the core of the OMZ at St. T5 (4 to 1%) and at T3 (8 to 1%)
322 between 90-400 m, where oxygen had the lowest concentrations (< 20 µM to 1µM). Abundances showed slight increases in
323 the **OMZ-LW** stratum during daytime, where oxygen levels increased after the extremely low levels within the OMZ. At the
324 onshore station off 29°S (St. L6), having a weaker and less extensive OMZ, the abundance showed similar vertical
325 distribution. However, the abundance of copepods was lowest in this station (in about 69%) in comparison with stations off
326 Iquique, in the core of the OMZ the percentage was between 5 to 3%.

327 DVM of Copepods was pronounced at onshore stations (Stations T3 and L6), but the strength of migration was higher
328 overall at St. T3 off Iquique, as reflected by the migration indices (WMD and ΔDVM) (**Table 1**).The WMD of these taxa
329 showed a broad range (17–500 m), which varied significantly among copepods groups and stations, both in day and night
330 samples (p <0.05) (**Fig. 3**). During the night, at the offshore station (St. T5) most Copepods exhibited normal DVM and they
331 were concentrated mainly in the oxic surface strata (**OX-M**) and **OMZ-UB** (40-60 m) without interacting with the OMZ;
332 except for Eucalanidae which concentrated deeper in the **OMZ-LC** stratum, associated with the lower core of the OMZ and
333 showing a high ΔDVM (225m). During the day these four groups of Copepods tended to remain deeper in the stratum
334 associated with the lower core of the OMZ (**OMZ-UC**) and lower O₂ gradient (**OMZ-LW**), except for small Copepods that
335 remained at the **OMZ-UB** stratum and showed a smaller ΔWMD (34 m). At the offshore stations (Stations T3 and L6) most
336 Copepods exhibited reverse DVM, except for large Copepods (LC) which showed slightly normal DVM at St. L6 off 29° S.
337 At night Copepods tended to concentrate deeper in the stratum associated with the lower core of OMZ (**OMZ-UC**) and
338 lower O₂ gradient (**OMZ-LW**), particularly Eucalanidae which exhibited a strong DVM with high ΔWMD of 440 m (St. T3)
339 and 208 m (St. L6) and *Acartia* Copepods with 103 m (St. L6) (**Table 1**), whereas at St. L6 small Copepods (SC) were
340 caught in abundance at the **OMZ-UB** stratum down to 82–90 m depth, respectively (**Fig. 3**). During the day, Copepods
341 remained shallower than at night, although they concentrated at different depths. Small Copepods tended to concentrate in
342 the oxic surface strata **OX-ML** (St. T3) and remained in the upper boundary of the OMZ (St. L6) without detectable DVM,
343 as judging by the small difference between their day- and night-time distributions (ADVM ca. 4 m). Large Copepods (LC),
344 as expected, showed a normal migration and tended to stay inside the OMZ and concentrated in the **OMZ-UC** stratum (St.
345 L6) and **OMZ-UB** (St.T3). Finally, Eucalanidae, which exhibited a strong DVM tended to distribute in the **OMZ-UC**
346 (St.T3) and the **OMZ UB** (St. L6) (**Fig. 3**).

347 Following Copepods, Salps were the second most abundant taxa of zooplankton, showing the largest aggregation at the
348 onshore station St. L6 (~3%), where abundances made up to $10347 \pm 219 \text{ ind. m}^{-2} \text{ d}^{-1}$. The WMD of Salps showed a range
349 between 54 and 25 m depth (**Table 1**) and appeared to aggregate deeper at all stations, also present in the OMZ (**Fig. 3**).
350 Similarly to Salps, most Chaetognaths showed the largest aggregation at the onshore station St. L6 (~2%), where their
351 abundances reached up to $4755 \pm 1038 \text{ ind. m}^{-2} \text{ d}^{-1}$. The abundance and biomass of this group increased in the upper boundary
352 of the OMZ (**OMZ-UB**) during day and night. No DVM was discernible for this group in this station, because of the slight
353 difference between their day- and night-time distributions (**Fig. 4**). By contrast, off Iquique they appeared to perform a
354 strong DVM between the **OMZ-UB** and the **OMZ-UC** strata, as indicated by the migration indices (WMD- Δ DVM) (**Table**
355 **1**). However at the onshore station (St.T3) they showed a reverse DVM.
356 Euphausiids exhibited greater abundances at the onshore station L6 (<1%), where they reached up to $4755 \pm 1038 \text{ ind. m}^{-2} \text{ d}^{-1}$,
357 although regarding vertical distribution they were mainly concentrated in the **OMZ-UB** and **OMZ-LC** in St. T5 and St. T3
358 at day and at night, respectively. No DVM was detectable in the onshore station (St. L6), judging by the small difference
359 between their day- and night-time distributions (**Fig. 4**). However, at the offshore station St. T5 they appeared to perform a
360 strong DVM with a vertical range between 236 and 56 m and a mean Δ DVM of 181 m (**Table 1**). Decapods larvae were
361 abundant at the **OMZ-UB** stratum during the day and at the **OMZ-UC** at layer at night in the offshore station (St.T3), where
362 they reached up to $292 \pm 62 \text{ ind. m}^{-2} \text{ d}^{-1}$, with a weak reverse DVM. However, at the offshore station St. T5 they appeared to
363 perform a strong normal DVM with a vertical range between 120 and 30 m and a mean Δ DVM of 90 m (**Table 1**).
364 The other main taxa, Appendicularian, showed the largest aggregation at the onshore station T5 (5%), where abundances
365 reached up to $21848 \text{ ind. m}^{-2}$. The abundance and biomass of this group were mainly found in the oxic surface strata OX-ML
366 at St. T5 and St. L6 both at day and at night. No DVM was detected in these stations, judging by the small difference
367 between their day- and night-time distributions, while at the onshore station (St. T3) they appeared to have a strong reverse
368 DVM into the OMZ-UC stratum at night as indicated by the migration indices (WMD- Δ DVM) (Table 1). The WMD at this
369 station T3 showed a range between 18 and 106 m depth and Δ DVM of 88 m depth (**Table 1**).

370 **3.3.3 Others groups with vertical distribution associated to OMZ UC**

371 The remaining 19 groups that only constituted 12% (in St.T5) 17% (in St.T3) and 26% (in St.L6) in abundance, showed
372 variable DVM behaviours at each station: Normal DVM (in St.T5) and reverse DVM (in Sts.T3-L6) (**Table 1**). These groups
373 clearly exhibited different daytime and night depths associated with the **OMZ core (OMZ UC-LC)**. Overall, they tended to
374 reside deeper by day and shallower by night in St. T5 than at the other sites (**Fig. 3**).

375 **3.4 Vertical distribution of zooplankton biomass**

376 Estimates (mean \pm SD) of biomasses of the taxonomic groups integrated by depth strata are summarized in **Table 2**. These
377 data, averaged from day and night measurements, contrast with the numerical abundances, which were dominated by
378 copepods. In this case, the bulk of zooplankton biomass was dominated by different groups depending on Stations. In terms

379 of biomass, Salps, Copepods, Chaetognaths, Euphausiids and Decapods accounted, more or less equally, for >70% in the
 380 whole area (**Table 2**). The bulk of biomass at Station T5 was dominated by Decapods, Salps, Copepods and Siphonophores
 381 >70% in this station, while Salps largely dominated the bulk of biomass at Station T3 (>60%) followed by Chaetognaths
 382 (12%). At the onshore Station L6, Copepods and Chaetognaths accounted for about 40% of total biomass (**Table 2**).
 383

384 **Table 2.** Mean and Standard Deviation (\pm) of integrated biomass (mg C m^{-2}) by taxonomic groups identified and sorted by
 385 ZooScan during daytime/ night-time conditions at three stations (T5, T3 and L6) sampled off northern Chile, during the
 386 austral spring 2015. Mean \pm SD are from $n=8$ for Stations T5 and T3, and $n=4$ for Station L6.

TAXA	T5		T3		L6	
	Integrated Biomass		Integrated Biomass		Integrated Biomass	
Salps	652.37	\pm 262.0	9904.93	\pm 1906.6	963.60	\pm 245.6
Copepods						
Large Copepods	552.91	\pm 137.4	231.05	\pm 46.9	838.28	\pm 139.6
Small Copepods	419.12	\pm 96.9	271.77	\pm 48.4	2500.51	\pm 400.3
Eucalanidae Copepods	0.02	\pm 0.0	0.00	\pm 0.0	0.44	\pm 0.2
Acartia Copepods	0.71	\pm 0.3	8.75	\pm 3.2	50.89	\pm 16.3
Chaetognaths	215.71	\pm 72.1	1800.82	\pm 744.1	2261.81	\pm 510.6
Euphausiids	296.39	\pm 69.0	436.24	\pm 115.4	919.43	\pm 250.6
Decapoda Larvae	21.83	\pm 5.4	1074.62	\pm 321.2	549.82	\pm 160.0
Hydrozoa	275.97	\pm 106.9	363.61	\pm 128.2	796.03	\pm 182.7
Annelidae	71.17	\pm 9.8	105.60	\pm 15.4	1044.62	\pm 268.9
Siphonophores	452.64	\pm 106.5	363.37	\pm 103.3	124.70	\pm 29.9
Decapods	652.79	\pm 279.4	0.00	\pm 0.0	0.00	\pm 0.0
Appendicularian	30.30	\pm 10.1	14.79	\pm 5.9	543.98	\pm 232.7
Ctenophores	0.62	\pm 0.3	119.55	\pm 53.5	102.50	\pm 31.7
Ostracods	11.92	\pm 1.3	16.51	\pm 1.1	185.56	\pm 35.4
Amphipods	10.62	\pm 2.2	51.17	\pm 15.8	30.73	\pm 6.4
Fish Eggs	17.64	\pm 3.0	10.64	\pm 1.4	47.24	\pm 9.1
Platyhelminthes	0.00	\pm 0.0	0.00	\pm 0.0	62.04	\pm 26.7
Nauplii	3.11	\pm 0.8	1.41	\pm 0.4	25.53	\pm 7.6
Ichthyoplankton	6.98	\pm 2.1	18.83	\pm 8.4	0.00	\pm 0.0
Pteropods	5.58	\pm 2.4	0.89	\pm 0.2	4.27	\pm 1.8
Foraminifera	1.14	\pm 0.4	0.14	\pm 0.0	9.13	\pm 2.5
Gastropods	0.94	\pm 0.2	0.56	\pm 0.1	3.59	\pm 0.4
Cirripedia Larvae	0.25	\pm 0.1	0.14	\pm 0.0	4.10	\pm 1.1
Bryozoa	0.80	\pm 0.2	0.20	\pm 0.0	1.35	\pm 0.4
Radiolarian	0.04	\pm 0.0	0.08	\pm 0.0	1.03	\pm 0.4
Echinoderm Larvae	0.01	\pm 0.0	0.00	\pm 0.0	0.04	\pm 0.0

Total	3702	±	805	14796	±	2512	11071	±	1635
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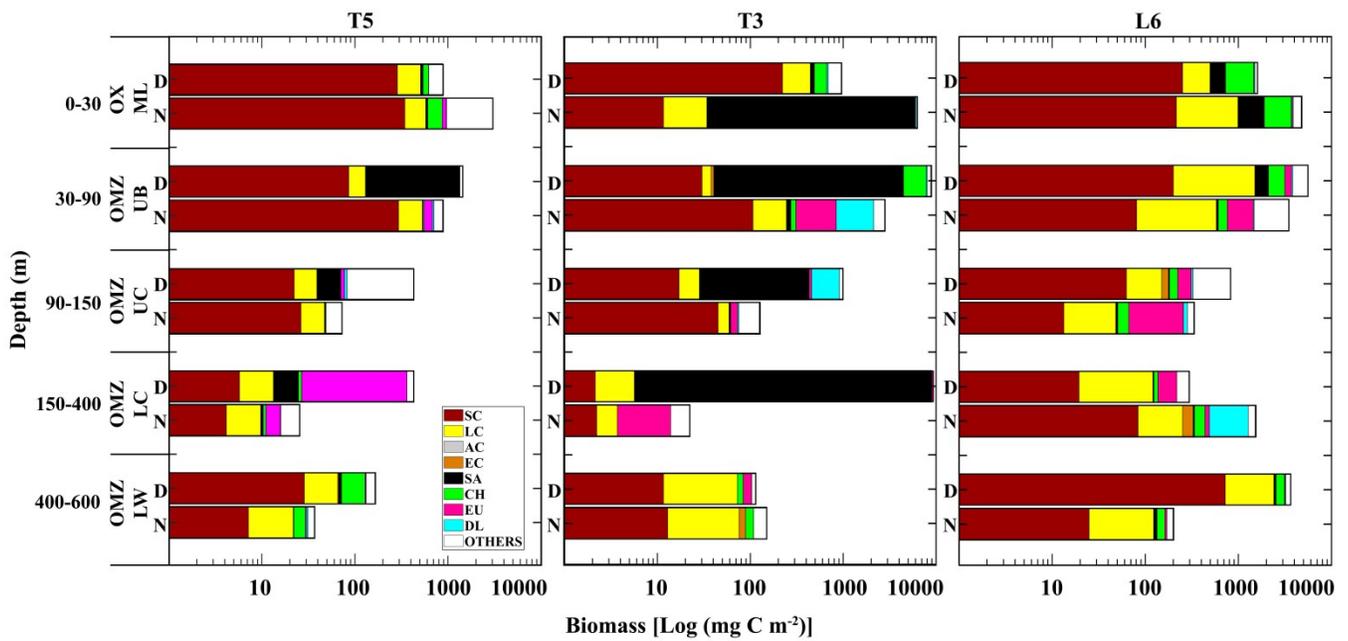
387

388 When assessing the day vs. night vertical distribution of taxonomic groups in terms of their contribution to biomass, different
389 patterns arise compared to numerical abundance. In this case, we used 9 taxonomic categories to examine vertical
390 distribution and DVM in terms of biomass: small Copepods (SC), large Copepods (LC), *Acartia* Copepods (AC),
391 Eucalanidae Copepods (EC), Salps (SA), Chaetognaths (CH), Euphausiids (EU), Decapod larve (DL), and all the other taxa
392 (**Fig. 5**). Contrasting with numerical abundance (**Fig. 4**), the vertical distribution of biomass was more heterogeneously
393 divided among taxonomic groups and DVM patterns vary strongly between stations. Small Copepods continue to dominate
394 in the upper oxic layer (**OX-ML**), but at the Station T3 their biomass decreases sharply within the **OMZ-UB** and within the
395 OMZ (**OMZ-UC** and **OMZ-LC**). This abrupt decrease in biomass coincides with the intense OMZ present at this station T3.
396 Large Copepods tended to distribute more homogeneously in the water column, although they also tend to diminish their
397 biomass during the day in the upper layers, while Euphausiids showed their ascent from deep layers to the upper ones at
398 Station T5, but with lower biomass within the OMZ at the onshore stations T3 and L6. Chaetognaths were another group
399 exhibiting important vertical movements of biomass between day and night across strata, although with not clear night ascent
400 (**Fig. 5**).

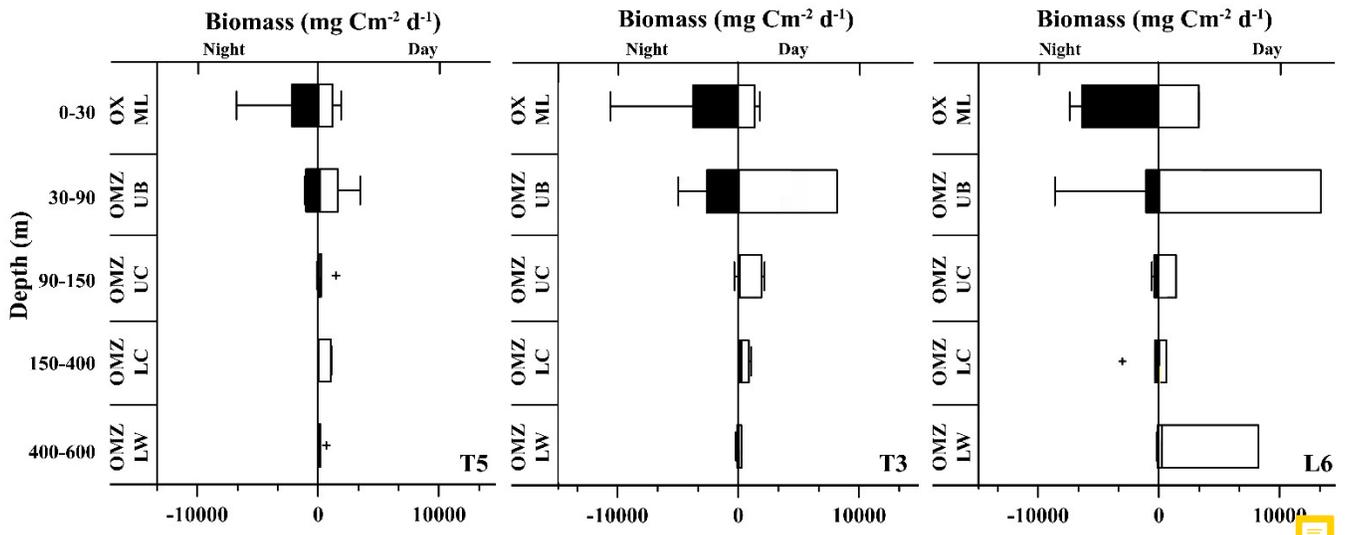
401 Total added biomass of zooplankton revealed more clearly DVM behaviour of the whole zooplankton community (**Fig. 6**).
402 The vertical distribution and daytime vs. Night-time variability of zooplankton biomass showed distinctive features
403 associated with the OMZ structure, with significant differences ($p < 0.05$) between strata for both daytime and night-time
404 samplings, as based on the ANOSIM test ($p < 0.05$). In the whole area most of the biomass was concentrated in a narrow band
405 within the OX-ML and OMZ UB strata, associated with more oxygenated surface waters, with reduced values in deeper
406 waters associated with the OMZ core, in special at the onshore station off Iquique (St. T3) (Fig. 6). Overall, we observed
407 that highest values of biomass were present during the night at the shallower sampling stratum (Ox-ML) and in the
408 subsurface during the day. There was also an important increase in biomass at the deepest stratum (OMZ-LW) during the
409 day.

410

411



412
 413 **Fig. 5.** Daytime vs. Night-time vertical distribution of biomass of different taxonomic groups at 3 stations off northern
 414 off Iquique (Stations T5 and T3) and off Coquimbo (Station L6). Data are from night and day replicated samples during two
 415 consecutive days in the austral spring 2015. Values represent means from sampling size n=4 for St. T5 and St. T3, and n=2
 416 for St. L6. Layers were defined from variable levels of oxygen concentration (see Methods).
 417



418
 419 **Fig. 6.** Vertical distribution of total zooplankton biomass during daytime and night-time conditions at 3 stations off northern
 420 Chile: off Iquique (Stations T5 and T3) and off Coquimbo (Station L6) during two consecutive days in the austral spring

421 2015. Some outliers are plotted individually (+). Error bars are standard deviations from sampling size (n=4 for St. T5 and
 422 St. T3 and n=2 for St. L6). Layers were defined from variable levels of oxygen concentration (see Methods).

423 3.4.1 Migrant biomass of the zooplankton taxa

424 The migrant biomass of the zooplankton taxa and the rate of migration (RM), represented by the proportion of biomass (%)
 425 being vertically moved daily from the upper 90 m are shown in **Table 3**. Most groups showed a high rate of migration as
 426 reflected in the RM. In terms of migrant biomass, Copepods, Chaetognaths, Euphausiids, Hydrozoa and Decapod larvae
 427 accounted for a large proportion of total migrant biomass (>80%), although high estimates of migrant biomasses were also
 428 associated with high standard deviations, indicating a strong variation among replicated samples (**Table 3**). Presence of zero
 429 values in **Table 3** represents absence of a given taxonomic group in the upper 90 m layer or extremely low values of biomass
 430 under both conditions' day and night, so that such groups did not contribute or they had a non-significant contribution to
 431 total migrant biomass.

432
 433 **Table 3.** Migrant biomass (mg C m⁻²) and rate of migration (RM) (%) for taxonomic groups of zooplankton sampled at
 435 northern Chile at 3 stations: off Iquique (Stations T5 and T3) and off Coquimbo (Station L6) during the austral spring 2015.
 436 RM represents the proportion (%) of migrant biomass with respect to total biomass found at night in the 0-90 m for a given
 437 taxonomic group. SD= Standard deviation of the migrant biomass estimated from n=4 (St. T5 and St. T3) and n=2 (St. L6).

TAXA	ST-T5			ST-T3			ST-L6		
	Migrant Biomass	SD	RM %	Migrant Biomass	SD	RM %	Migrant Biomass	SD	RM %
Salps	0.00	0.00	0	2296.90	3248.31	54	314.46	444.71	60
Copepods									
Large Copepods	202.92	143.03	41	0.00	0.00	0	261.35	369.61	44
Small Copepods	266.81	67.20	41	0.00	0.00	0	0.00	0.00	0
Eucalanidae Copepod	0.00	0.00	0	0.00	0.00	0	0.00	0.00	0
<i>Acartia</i> Copepod	0.04	0.06	100	0.00	0.00	0	0.00	0.00	0
Chaetognaths	243.92	344.96	93	0.00	0.00	0	651.37	921.18	59
Euphausiids	209.63	288.02	58	530.47	616.61	99	212.29	67.48	48
Decapoda Larvae	27.76	21.73	79	1261.69	1784.30	100	6.54	9.25	14
Hydrozoa	427.18	603.78	48	0.00	0.00	0	41.90	59.25	77
Annelidae	51.36	31.95	72	20.92	29.59	80	182.49	57.38	36
Siphonophores	113.28	140.89	34	71.80	101.55	27	32.45	45.89	46
Decapods	1296.57	1833.63	100	0.00	0.00	0	0.00	0.00	0
Appendicularia	0.00	0.00	0	0.00	0.00	0	1041.36	1471.54	52
Ctenophores	0.00	0.00	0	239.10	338.14	100	149.20	211.01	50
Ostracods	9.07	5.29	81	0.06	0.09	2	28.57	8.78	50

Amphipods	0.12	0.18	9	35.60	20.17	56	38.74	54.78	98
Fish Eggs	7.50	4.07	61	0.00	0.00	0	1.11	1.56	14
Platyhelminthes	0.00	0.00	0	0.00	0.00	0	2.12	2.99	100
Nauplii	4.42	2.75	84	0.00	0.00	0	4.05	5.73	42
Ichthyoplankton	0.06	0.08	0	0.00	0.00	0	0.00	0.00	0
Pteropods	0.00	0.00	0	0.91	1.29	100	8.10	2.57	99
Foraminifera	0.05	0.05	80	0.00	0.00	0	0.00	0.00	0
Gastropods	0.76	1.07	87	0.00	0.00	0	0.12	0.17	31
Cirripedia Larvae	0.38	0.26	97	0.06	0.08	100	0.00	0.00	0
Bryozoan	1.50	1.13	99	0.00	0.00	0	1.10	1.55	82
Radiolarian	0.00	0.00	0	0.00	0.00	0	0.00	0.00	0
Echinoderm Larvae	0.03	0.04	100	0.00	0.00	0	0.00	0.00	0
Total	2863	1177.94		4457	4227.70		2977	116.66	

438

439

440 **4 Discussion**

441 Studies on zooplankton DVM and the active transport of C mediated by zooplankton have been documented previously for
442 the Pacific Ocean and for other areas of the world's ocean, as summarized in **Table 4**. However, downward C flux due to
443 DVM in highly productive upwelling regions, such as northern Chile, which is also characterized by severe subsurface
444 hypoxic conditions upon presence of a shallow OMZ, is still poorly understood. Some studies have shown that hypoxic
445 conditions can interfere with DMV of many meso- and macrozooplankton species (Wishner *et al.*, 2013; Ekau *et al.*, 2010;
446 Escribano *et al.*, 2009; Apablaza and Palma, 2006; Antezana, 2002; Escribano, 1998). These studies have shown that small
447 differences in oxygen concentration can make a large difference for zooplankton behavior, physiology and adaptation
448 (Wishner *et al.*, 2018; Kiko *et al.*, 2016; Seibel, 2011; Gonzalez and Quiñones, 2002; Escribano and McLaren, 1999).
449 Therefore, it seems that the OMZ can play a very significant role influencing vertical distribution, DVM and ultimately the
450 downward C flux mediated by zooplankton.

451 Our approach to assess downward C flux into the Oxygen Minimum Zone, based on estimates of the migrant biomass and
452 our proposed migration indices, allowed us on one hand to examine the contribution that different zooplankton can have for
453 vertical flux of C and hence export production, and in the other hand to assess zooplankton responses (e.g. vertical
454 distribution and DVM performance) to changes in environmental conditions over the vertical gradient, such as temperature,
455 water density and the abrupt changes in oxygenation levels. In this subtropical upwelling region, vertical gradients are much
456 stronger than in temperate upwelling zones. For example, the coastal zone in this region is more stratified and with a very
457 shallow OMZ (<50 m) with a weak seasonal signal and moderate upwelling throughout the year (Paulmier and Ruiz-Pino,
458 2009; Fuenzalida *et al.*, 2009; Escribano *et al.*, 2004). This means that zooplankton must cope with hypoxic conditions
459 during their entire life cycle, except for some species that may reside in near surface water (<30 m), such as *C. chilensis* and

460 *C. brachiatus* which have been reported as mostly restricted to the upper layer without performing any substantial DVM
461 (Escribano *et al.*, 2012, 2009; Escribano and Hidalgo, 2000; Escribano, 1998).

462 The vertical distribution and diurnal variability of zooplankton biomass seem to be disturbed by the OMZ, such that high
463 biomass aggregates above the oxycline in a narrow band within the OX-ML and OMZ-UB layers, associated with more
464 oxygenated surface waters, whereas extremely low biomass reside in deeper waters, in particular within the OMZ core. This
465 condition was more evident in the coastal station off Iquique (St.T3), characterized by the most intense OMZ in the whole
466 study area. In the eastern tropical north Pacific, biomass distribution seemed different, exhibiting a secondary peak at depth
467 during the daytime within the upper oxycline or OMZ core (Wishner *et al.*, 2013).

468 Regarding the estimates of biomass for each of the taxonomic groups, our approaches can certainly introduce variation,
469 depending on selected regressions and conversion factors from highly diverse body shapes and body densities of the
470 zooplankton taxa affecting the estimates of biovolume, dry weight and C content. Various approaches have been adopted for
471 converting sizes to body masses. For example, Lehette and Hernández-León (2009) provided some general regression
472 equations for subtropical and Antarctic zooplankton describing the relationship between scanned area and body mass (C
473 content). These authors also proposed two separate regressions for crustacean and gelatinous zooplankton, because of
474 different body densities. In our samples there was a high diversity of taxonomic groups as identified by ZooScan, such that
475 unique regressions for crustacean and gelatinous organisms may lead to strong biases in body mass estimates, because of
476 high variability in C content, which is the key component of body mass needed to estimate C flux. Therefore, following
477 Yamaguchi *et al.* (2014) and Sato *et al.* (2015), we converted biovolumes into dry weights using a mean body density and
478 mean water content of zooplankton, estimated across taxa by Wiebe (1975), but then taxa-specific Carbon: dry weight ratios
479 collected from published works. In any case, the use of single conversion factors between body volume and mass and wet
480 weight and dry weight among taxa must be considered as source of variability in the estimates of taxonomic biomasses.

481 Despite the apparently hostile oxygen-deficient habitat, associated with the OMZ, we found that most taxa were able to
482 perform DVM in the upwelling zone withstanding severe hypoxia. Even, several zooplankton groups are strong migrants,
483 exhibiting large DVM amplitude (~500 m). Among them, an important migrant group is comprised by the copepods
484 Eucalanidae which have been described as even being able to enter the core of the OMZ, and then migrate downward to the
485 lower limit of the OMZ, which is slightly more oxygenated (Hidalgo *et al.*, 2005). In our study however, their contribution to
486 total migrant biomass was too small (ca. 2.6 mg C m⁻² d⁻¹), as compared to the estimate made by Hidalgo *et al.* (2005). In
487 fact, the migrant biomass and rate of migration of this group was non-significant when considering DVM between the upper
488 90 m and below, suggesting a little or no contribution to downward flux of C for this group of copepods. However it seems
489 that Eucalanidae remains below the oxycline or nearby the base of the oxycline day and night, as shown by their weighted
490 mean depth (WMD) and therefore suggesting that they may still contribute to vertical flux by feeding at the base of the
491 oxycline at night and then migrating into the OMZ during the day.

492 Other taxa, such as Euphausiids, *Acartia* spp., other copepods, Ctenophores, Decapods, Annelidae, Bryozoa, Pteropods and
493 Chaetognaths tended to concentrate their populations inside the OMZ core showing a strong link to the OMZ with important

494 movement throughout the water column. Antezana (2010) showed that *E. mucronata*, an endemic and abundant euphausiid
 495 in the coastal upwelling zone off Chile, is a well-adapted species to vertically migrate into the core of the OMZ. In fact, the
 496 euphausiids studied here showed a large DVM amplitude (~250 m), descending into the core of the OMZ and below 250 m
 497 each day. In general, all strong migrants taxa showed a strong interaction with the core of OMZ, remaining there either
 498 temporarily or permanently during the day or night condition, contributing in this way to the release of C below the
 499 thermocline, despite presence of hypoxic conditions.

500 Our estimates of DVM-mediated C flux showed that migrant biomass ($3433 \pm 889 \text{ mg C m}^{-2} \text{ d}^{-1}$) and C flux estimates
 501 ($674 \pm 291 \text{ mg C m}^{-2} \text{ d}^{-1}$) of the major taxa performing DVM, were greater than those reported for the Pacific Ocean, both in
 502 oligotrophic, such as Hawaii, and mesotrophic waters, as the subarctic North Pacific (Steinberg *et al.*, 2008), and even
 503 greater than that informed by Yebra *et al.* (2005) within eddies with enhanced biological production. Most of these previous
 504 estimates however have not been done in regions with severe hypoxia or anoxia at mid water depths (e.g. Kiko *et al.*, 2016),
 505 such as the highly productive upwelling region of the coastal zone off northern of Chile, where the oxygen concentrations
 506 may fall below $<1 \mu\text{mol}$ in the core of OMZ (Paulmier and Ruiz-Pino, 2009). Moreover, only few works have considered the
 507 whole zooplankton community (**Table 4**). High productivity and strong aggregation of zooplankton in coastal areas of this
 508 region (Escribano *et al.*, 2000; Escribano and Hidalgo, 2000) may promote greater amounts of migrant biomass. This
 509 requires however that DVM should not be majorly constrained by presence of the OMZ and that most migrant taxa are
 510 tolerant to low-oxygen. On the other hand, our estimates of downward C flux were substantially lower than previous ones
 511 reported off northern Chile by Hidalgo *et al.* (2005) for *Eucalanus inermis* alone ($14.1 \text{ mg C m}^{-2} \text{ d}^{-1}$) and for copepods and
 512 euphausiids by Escribano *et al.* (2009) ($7200 \text{ mg C m}^{-2} \text{ d}^{-1}$) (**Table 4**). Although, such previous estimates may be too high,
 513 considering the level of primary production in the upwelling zone of Chile ($\sim 10000 \text{ mg C m}^{-2} \text{ d}^{-1}$, the maximum estimated
 514 value) (Daneri *et al.*, 2000), suggesting that previous works may have overestimated active transport of C. Although the
 515 work of Escribano *et al.* (2009) was based on samples obtained with a Tucker Trawl net, which can be more efficient in
 516 capturing large-sized zooplankton or macrozooplankton (Escribano *et al.*, 2007), as compared with the vertically towed
 517 Multinet. This means that our estimates mainly based on mesozooplankton may not include the contribution of some
 518 macrozooplankton, and therefore such values may be greater.

519
 520 **Table 4.** Comparison of active transport of carbon (AC) ($\text{mg C m}^{-2} \text{ d}^{-1}$) by vertically migratory taxa in Pacific Ocean. Diel
 521 vertically migratory taxa (DVM), productivity primary (PP) ($\text{mg C m}^{-2} \text{ d}^{-1}$), migrant biomass (MB) (mg C m^{-2}), respiratory
 522 loss (R) ($\text{mg C m}^{-2} \text{ d}^{-1}$), faecal pellets production (F) ($\text{mg C m}^{-2} \text{ d}^{-1}$) and mortality (M) ($\text{mg C m}^{-2} \text{ d}^{-1}$). Where provided by
 523 authors, estimated passive export (POC) is listed. Fluxes refer to carbon export beneath the epipelagic zone (150–200 m
 524 depth, depending on the study) in $\text{mg C m}^{-2} \text{ d}^{-1}$

Location	Taxa	PP	MB	AC	R	F	M	% POC	References
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N.Hawaii ALOHA	DVM Zooplankton		108–216	7.1	2.6–4.8			12–18	Al-Mutairi and Landry (2001)
N. Hawaii ALOHA			157.9	3.2–13.6	3.7			18	Steinberg <i>et al.</i> (2008)
N.W. Pac.	DVM Metridia	418	144	9	3	5	1	23.1–61.8	Kobari <i>et al.</i> (2008)
N.W. Pac.	DVM copepods			8				22.3	Takahashi <i>et al.</i> (2009)
N.E. Pac.	Mesopelagic fishes	170			23.9				Davison <i>et al.</i> (2013)
Eastern Equator			96.0± 25.2		4.2± 1.2		2.9±0.8	18.4	Zhang and Dam (1997)
Eastern Equator	DVM zooplankton		154.8± 32.4		7.3± 1.4		5.4±1.1	25.4	Zhang and Dam (1997)
Central Equator (HNLC)			52.9		6		–	4	Rodier and Le Borgne (1997)
Western Equator			46.9		3		–	6	Rodier and Le Borgne (1997)
E. Eq. Pac.	DVM Zooplankton		1214	7.1	7.1			204	Rodier and Le Borgne (1997)
Western Equator	DVM Zooplankton		144–447	23.53-9.97	7.3–19.1		2.6–4.4	13–35	Hidaka <i>et al.</i> (2002)
Equator divergence			2.8–21.8		0.9–1.2			<1-2	Roman <i>et al.</i> (2002)
Oligotrophic area			30.2–33.8		1.3–1.7			4	Roman <i>et al.</i> (2002)
E.S.Pac. N.Chile		5503							Gonzalez <i>et al.</i> (1998)
E.S.Pac. N.Chile		10000							Daneri <i>et al.</i> (2000)
E.S.Pac. N.Chile	DVM Eucalanus		8.0 -34	14.1					Hidalgo <i>et al.</i> (2005)
E.S.Pac. N.Chile	DVM zooplankton		37810	7200		670	0		Escribano <i>et al.</i> (2009)
E.S.Pac. N.Chile	DVM zooplankton	2833±1155	3433 ±889	674±291					This study

525 Differences of our estimates with previous works may also be accounted by strong variability of zooplankton abundance in
526 the upwelling zone. In fact, our estimates of migrant biomasses of the different taxonomic groups based on 2 days of
527 sampling and two replicates for each condition (day and night) are strongly variables, as shown by the standard errors in
528 **Table 3** which can be as much as 100% from the mean value. Therefore, comparisons must take caution upon strong time-
529 space variation when assessing zooplankton abundance.

530 Even although the OMZ did not greatly prevent DVM migration, zooplankton behaviour appeared disrupted or exhibited
531 reversed patterns, depending on vertical distribution of OMZ and on the taxonomic group being considered. This behaviour
532 was more evident in the onshore stations (Stations T3 and L6), but in particular in the station off Iquique (St. T3) that also
533 showed a higher migration rates (60%). According to Ekau *et al.* (2010), other indirect effects could also be caused by the
534 hypoxia conditions, such as changes in prey availability, prey size or predation risk, as well as changes in species

535 composition, the strength of which depends on the duration and intensity of the hypoxic events. This could explain why
536 individuals within a single population can perform reverse, normal, or non DVM, apparently depending on the more
537 important source of mortality: predation by nocturnal feeding, normally migrating carnivorous zooplankton, or visually
538 hunting planktivorous fish (Ohman, 1990). These kind of DVM behaviors can only be better assessed and understood when
539 looking at the population level, although again time-space variation in zooplankton abundance in a highly heterogeneous
540 upwelling zone should be kept in mind.

541 Concerning C fluxes, our estimates of active transport of carbon by zooplankton were greater than estimates of passive C
542 sinking obtained off northern Chile at 60 m depth off Antofagasta (23°S) by Gonzalez *et al.* (1998) based on sediment traps
543 (125 to 176 mg C m⁻² d⁻¹). Regarding the question on how efficient is active C transport mediated by DVM, we obtained
544 satellite-based (<http://science.oregonstate.edu.ocean.productivity>) estimates of net primary production (monthly means for
545 November-December 2015) for the coastal area (Stations T3 and L6) and the coastal transition zone (Station T5), averaged
546 for the months of November and December 2015. Our estimates of downward C flux represented a mean of 25 % of export
547 of Carbon resulting from net primary production in the upwelling region, estimated in the range of 1500-3500 mg C m⁻² d⁻¹
548 (**Table 5**). If we consider this is accounted only by mesozooplankton, then an important fraction of freshly produced C might
549 be taken downward by zooplankton, and this DVM-mediated C flux ought to be taking into account when analysing and
550 modelling the C budget in the upwelling zone.

551

552 **Table 5.** Mean net primary production rate and estimates of daily downward C flux due to passive sinking and mediated by
553 diel vertical migration (DVM) of mesozooplankton at three stations (T5, T3 and L6) in the coastal upwelling region off
554 northern Chile during the austral spring 2015. Primary production represents satellite-based estimates of monthly mean
555 (November–December 2015) at the 3 sampling stations. Passive C flux is a mean value estimated from sediment traps by
556 González et al. (2000) off Antofagasta (northern Chile, 23 °S) for January 1997. Total biomass and epipelagic biomass are
557 mean observed values from day-night conditions after 2 consecutive days of sampling.
558

Station	T5	T3	L6	Mean ± SD
Primary Production mg C m ⁻² d ⁻¹	1500	3500	3500	2833 ± 1155
Passive Carbon Flux mg C m ⁻² d ⁻¹				151 ± 36
Integrated Abundance ind. m ⁻² (0–600 m)	221735	127085	371235	240018 ± 123097
Total biomass mg C m ⁻² d ⁻¹ (0–600m)	3702	14796	11071	9856 ± 5646
Epipelagic biomass (mg C m ⁻²) (0–90m)	3126	9391	7679	6732 ± 3238
Migrant biomass (mg C m ⁻² d ⁻¹)	2863	4457	2977	3433 ± 889
Rate of Biomass migration (%)	51	26	37	38 ± 12
Active Carbon Flux (mg C m ⁻² d ⁻¹)	428	996	598	674 ± 291
Total Active Carbon exported (%)	29	28	17	25 ± 7

559

560 5. Conclusions

561 In the coastal upwelling zone off northern Chile the presence of a subsurface oxygen minimum zone (OMZ) can impose an
562 important constraint for diel vertical migration of zooplankton and so influencing the downward C export mediated by
563 zooplankton. We found that most of the zooplankton biomass aggregates above the oxycline, associated with more
564 oxygenated surface waters and this was evident upon presence of a more intense OMZ. Some taxonomic groups however,
565 were found closely associated with the OMZ and several taxa were able to perform DVM in the upwelling zone withstanding
566 severe hypoxia. Also strong migrants, such as large sized copepods and copepod of the group Eucalanidae and Euphausiids,
567 can exhibit a large migration amplitude (~500 m), remaining either temporarily or permanently during the day or night
568 condition within the core of the OMZ, and so contributing to the release of C below the oxycline (and thermocline). Our
569 estimates of DVM-mediated C flux suggested that a mean migrant biomass of 3.4 g C m⁻² d⁻¹ may contribute with about 0.6g
570 C m⁻² d⁻¹ to the OMZ system through respiration, mortality, and production of fecal pellets at depth, accounting for ca, 25%
571 of the net primary production, and so implying the existence of a very efficient mechanism to incorporate freshly produced C
572 into the OMZ. This downward C flux mediated by zooplankton DVM is however strongly depending on the taxonomic
573 structure due to variable migration amplitude and DVM behavior. These estimates should also consider the strong temporal-
574 spatial variation in zooplankton abundance in the upwelling zone for comparison purposes.

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