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1	Zooplankton diel vertical migration during Antarctic summer
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25 Abstract

26 Zooplankton diel vertical migration (DVM) during summer in the polar oceans is presumed to be 27 dampened due to near continuous daylight. We analyzed zooplankton diel vertical distribution 28 patterns in a wide range of taxa along the Western Antarctic Peninsula (WAP) to assess if DVM 29 occurs, and if so, what environmental controls modulate DVM in the austral summer. 30 Zooplankton were collected during January and February in paired day-night, depth-stratified 31 tows through the mesopelagic zone along the WAP from 2009-2017, as well as in day and night 32 epipelagic net tows from 1993-2017. The copepod Metridia gerlachei, salp Salpa thompsoni, 33 pteropod Limacina helicina antarctica, and ostracods consistently conducted DVM between the 34 mesopelagic and epipelagic zones. Migration distance for M. gerlachei and ostracods decreased 35 as photoperiod increased from 17 to 22 h daylight. The copepods Calanoides acutus and 36 Rhincalanus gigas, as well as euphausiids Thysanoessa macrura and Euphausia 37 *crystallorophias*, conducted shallow (mostly within the epipelagic zone) DVMs into the upper 38 50 m at night. Rhincalanus gigas, T. macrura, and L. h. antarctica DVM behavior was 39 modulated by chlorophyll a concentration, mixed layer depth, and depth of the subsurface 40 chlorophyll a maximum, respectively. Carnivorous and detritivorous taxa – including the 41 calanoid copepod Paraeuchaeta antarctica, ostracods, chaetognaths, and Tomopteris spp. 42 polychaetes – as well as seasonally migrating copepods, were most abundant in the mesopelagic 43 zone regardless of the diel cycle. Paraeuchaeta antarctica underwent reverse DVM within the 44 top 100 m. The impacts of Antarctic zooplankton summer DVM and the resident mesopelagic 45 assemblage on carbon export should be better quantified.

46 **1. Introduction**

47 Many zooplankton and fishes throughout the world's oceans undergo diel vertical 48 migration (DVM), feeding in productive surface waters at night and seeking refuge from visual 49 predators at mesopelagic depths during the daytime (Hays, 2003). A global estimate suggests 50 \sim 50% of sound-scattering mesopelagic biomass performs DVM (Klevjer et al., 2016). Diverse 51 zooplankton taxa independently evolved DVM behavior, as it optimizes the adaptive balance 52 between feeding and predator evasion (Zaret and Suffern, 1976; Stich and Lampert, 1981; 53 Gliwicz, 1986; Hays, 2003). While predator avoidance is the accepted evolutionary driver for 54 DVM, shifts in downwelling irradiance at sunrise and sunset are the dominant proximate cues for 55 this behavior (Ringelberg and Van Gool, 2003; Cohen and Forward, 2009). Thus, it was assumed 56 that DVM is restricted in polar regions and may cease altogether in mid- winter and summer 57 during 24-h darkness and light, respectively (Blachowiak-Samolyk et al., 2006). 58 DVM studies in the polar oceans show seasonal variability in behavior, with DVM 59 magnitude changing in relation to photoperiod. During moored Acoustic Doppler Current 60 Profiler deployments in the Ross, Lazarev, and Weddell Seas, DVM continued through Antarctic 61 winter but ceased during the period of extended daylight from November to February (Cisewski 62 et al., 2010; Cisewski and Strass, 2016; Picco et al., 2017). Persistent winter DVM occurs as far 63 north as 77°N (Hobbs et al., 2018), although zooplankton DVM is restricted to small-scale (6 to 64 8 m) migrations within the upper 30 m during Arctic winter (Ludvigsen et al., 2018). Therefore, 65 the apparent pause of DVM during Antarctic summer may in fact be due to seasonal changes in 66 DVM amplitude as some species undertake shallower migrations during summer that go 67 undetected by conventional sampling (Flores et al., 2014; Daase et al., 2016). Asynchronous

migrations throughout the diel cycle may also explain why acoustic records do not detect
summer DVM in the Southern Ocean (Cottier et al., 2006).

In addition to photoperiod, other environmental conditions are likely to influence the
amplitude of zooplankton DVM during polar summer. Phytoplankton blooms may halt DVM as
zooplankton remain in surface waters to feed (Cisewski et al., 2010; Cisewski and Strass, 2016).
The depth of the subsurface chlorophyll *a* maximum can also influence zooplankton DVM, as
observed with Arctic copepods (La et al., 2015a). Similarly, vertically migrating Arctic

75 zooplankton concentrate just below the mixed layer (Berge et al., 2014). In the southern

76 California Current, increased light attenuation results in a decreased amplitude of copepod DVM

77 (Ohman and Romagnan, 2016). The interaction between light conditions and phytoplankton

78 distribution is thus likely to be a key driver of Antarctic zooplankton vertical distribution.

79 Few studies address polar mesopelagic zooplankton composition and 80 taxon-specific variability in DVM behavior. Arctic zooplankton demonstrate asynchronous 81 DVM patterns, which are explained by variation in feeding ecology, predation risk, and seasonal 82 migration behavior (Fortier et al., 2001; Cottier et al., 2006; Falk-Petersen et al., 2008). Prior 83 studies assessing zooplankton vertical distribution along the Western Antarctic Peninsula (WAP) 84 have either focused on specific taxonomic groups (e.g., Nordhausen, 1994a; Lopez and Huntley, 85 1995) or lacked the comparable day and night sampling necessary to assess DVM behavior 86 (Marrari et al., 2011). Additionally, there have not been any comprehensive studies resolving polar DVM variability over interannual timescales. 87

We analyzed zooplankton diel vertical distribution patterns along the WAP during
mid-summer as part of the Palmer Antarctica Long-Term Ecological Research (PAL LTER)
program. Zooplankton were sampled at discrete depth intervals through the epi- and mesopelagic

91 zones in paired day and night net tows using a MOCNESS (2009-2017) to assess taxon-specific 92 zooplankton vertical distribution. Day and night epipelagic net tows (1993-2017) throughout the 93 PAL LTER sampling region provided additional information on DVM behavior. We examined 94 environmental controls (e.g., photoperiod, mixed layer depth) on DVM amplitude in taxa 95 showing clear DVM. Our results show diverse zooplankton DVM modes, depth distributions, 96 and responses to phytoplankton biomass and vertical distribution, all of which can affect 97 zooplankton-mediated carbon export during Antarctic summer. 98 2. Materials and Methods

99 2.1. Study region

100 The PAL LTER study region ranges from Anvers Island (64.77°S, 64.05°W) in the north to Charcot Island (69.45°S, 75.15°W) in the south, extending from the WAP coast to the 101 102 continental slope (Ducklow et al., 2012) (Fig. 1). The PAL LTER research grid is composed of 103 sampling lines running perpendicular to the Peninsula every 100 km, and standard grid stations 104 within each line are separated by 20 km (Waters and Smith, 1992). From 1993-2008, the 105 sampling plan included all stations on grid lines 600 to 200. In more recent years, the study area 106 has expanded to include lines 100, 000, and -100, with sampling resolution reduced to three 107 stations per line. As in previous studies, three latitudinal sub-regions were designated to 108 represent hydrographic, sea ice, and ecological gradients (Martinson et al., 2008; Stammerjohn et 109 al., 2008; Steinberg et al., 2015) (Fig. 1), with regional boundaries along sampling grid lines as 110 follows: 'North' (lines 600 to 400), 'South': (lines 300 and 200), and 'Far South' (lines 100 to 111 -100).



Figure 1. PAL LTER study area along the Western Antarctic Peninsula. 'North,' 'South,' and
'Far South' regions are indicated. Circles indicate epipelagic sampling stations (1993-2017).
Stars indicate paired day-night MOCNESS sampling locations on the shelf (2009-2017). Cross

116 indicates paired day-night MOCNESS sampling station on the slope (2017). Shading indicates

117 bathymetry. An: Anvers Island; Ad: Adelaide Island; MB: Marguerite Bay; Ch: Charcot Island.

118 2.2. Zooplankton collection

Zooplankton were sampled during austral summer (02 January to 13 February) on annual
PAL LTER research cruises aboard the MV *Polar Duke* (1993-1997) and ARSV *Laurence M. Gould* (1998-2017). Zooplankton sampling was conducted using two types of gear as described
below.

123 2.2.1. Multiple discrete depth sampling through the epi- and mesopelagic zones 124 A 1.4-m² frame, 500-µm mesh Multiple Opening/Closing Net and Environmental 125 Sensing System (MOCNESS) (Wiebe et al., 1985) was used to collect meso- and 126 macrozooplankton in discrete depth intervals from 2009-2017. Each year, paired day (10:07-127 15:15 local start time) and night (23:03-01:46) MOCNESS tows were carried out in coastal or 128 shelf waters in the 'North', and in most years also the 'South' and 'Far South' (Fig. 1). Sampling 129 time and location were used to calculate solar elevation at the start of tows (Meeus, 1998); 130 daytime solar elevation was $> 35.51^{\circ}$ and nighttime solar elevation $< 0.56^{\circ}$. The MOCNESS was 131 towed obliquely at a speed of 2-2.5 knots, with a typical tow duration of 2.25-3 h. On average, 132 790 m³ (range 161-1900 m³) of water was filtered within a single depth interval as measured by a flow meter mounted on the system. 133

Eight discrete depth intervals were sampled during the upcast as follows: 500-400, 400-300, 300-250, 250-200, 200-150, 150-100, 100-50, and 50-0 m. Occasionally, the deeper intervals were not sampled when towing in waters shallower than the deep target depths. On four occasions, two depth intervals were combined due to net sampling errors. In these cases, the taxon density in each interval was assumed to equal the density calculated for the combined interval. Sea ice conditions occasionally prevented MOCNESS sampling in the South and Far South regions, therefore, the sample size n = 7 for paired day/night tows in the North, while n =

141	6 for the South, and $n = 4$ for the Far South from 2009-2015. Data from additional paired
142	MOCNESS tows in 2016 and 2017 are presented for euphausiids and salps only (for these taxa, n
143	= 10 for the North, $n = 8$ for the South, and remains $n = 4$ for the Far South).
144	A pair of day (09:54 local start time) and night (22:03) tows was also carried out over the
145	continental slope in 2017 (Fig. 1). In this case, eight discrete depth intervals were sampled during
146	the upcast as follows: 1000-750, 750-500, 500-400, 400-300, 300-200, 200-100, 100-50, and
147	50-0 m (100-0 m for day, due to a net sampling error). Salp data from this slope sample are
148	presented independent of the coastal and shelf data.
149	2.2.2. Epipelagic sampling
150	Macrozooplankton were also collected from 1993-2017 throughout the PAL LTER study
151	area using a 2 x 2 m square, 700-µm mesh Metro net towed obliquely to 120 m (Ross et al.,
152	2008, Steinberg et al., 2015) (Fig. 1). The net depth and tow profile were monitored with a depth
153	sensor linked to the conducting hydro wire. Average volume filtered was 9023 m ³ (range 1715-
154	71929 m ³), calculated using a General Oceanics flow meter suspended in the net opening. Ship
155	speed was 2–2.5 knots while towing, and typical tow duration was 30–35 minutes.
156	Epipelagic samples were designated night tows when the sun was below the horizon,
157	accounting for atmospheric refraction (calculated solar elevation \leq -0.833° at the start of the tow)
158	(Atkinson et al., 2008; Steinberg et al., 2015). Sample size varied by taxon (day $n = 966-1071$
159	and night $n = 181-198$), as not all taxa were identified during shipboard processing throughout
160	the time series.
161	2.3. Taxonomic composition

2.3.1. Discrete depth samples

163	All euphausiids and salps collected in MOCNESS tows were identified and quantified at
164	sea. Whole samples were then preserved in sodium borate-buffered 4% formaldehyde and
165	shipped to the Virginia Institute of Marine Science (Gloucester Point, VA, USA) for further
166	taxonomic analysis to quantify all non-euphausiid or salp taxa. Samples were size-fractionated
167	using a 5-mm mesh, with all individuals in this larger size fraction identified and counted. This
168	size-fractionation step removed large, abundant taxa (i.e., salps and euphausiids) from the
169	microscopic analysis. At least 1/64 of the < 5-mm size fraction was counted under a stereo
170	dissecting microscope after dividing the sample with a plankton splitter. A minimum of 100
171	individuals of the most abundant species was enumerated in this smaller size fraction.
172	Discrete depth analyses focused on abundant taxonomic groups. Five common calanoid
173	copepod species were included: Metridia gerlachei, Calanoides acutus, Calanus propinquus,
174	Rhincalanus gigas, and Paraeuchaeta antarctica. Identification included adults and conspicuous
175	copepodites. Discrete depth-stratified data were only analyzed for the smaller, but abundant,
176	euphausiids Thysanoessa macrura and Euphausia crystallorophias, because the larger Antarctic
177	krill Euphausia superba was underrepresented due to avoidance of the 1.4-m ² MOCNESS
178	(Nordhausen, 1994b). The pelagic tunicate Salpa thompsoni was included. The thecosome
179	(shelled) pteropod Limacina helicina antarctica was analyzed individually while the gymnosome
180	(shell-less) pteropods Clione antarctica and Spongiobranchea australis were grouped together
181	(as in Thibodeau et al., 2019). Tomopteris spp. polychaetes were combined into a single group
182	including T. carpenteri. Other groups were analyzed by major taxa, such as ostracods,
183	amphipods (including the hyperiids Themisto gaudichaudii, Cyllopus lucasii, Hyperiella
184	macronyx, Hyperoche medusarum, Primno macropa, Vibilia stebbingi, Scina spp., and the

185 gammarid *Eusirus* spp.), and chaetognaths (inclusive of large, conspicuous *Pseudosagitta* 186 gazellae and *P. maxima*).

187 2.3.2. Epipelagic samples

188 Grid-wide epipelagic tows were sorted at sea as reported in Steinberg et al. (2015). All 189 above taxa were included in the analysis of epipelagic samples except for the calanoid copepods 190 and ostracods. The Antarctic krill Euphausia superba was included in analysis of epipelagic 191 samples.

192 2.4. Vertical structure

193 Night to day ratios (N:D) were calculated to identify diel changes in surface abundance of 194 each taxon. For paired day and night MOCNESS samples, abundance was integrated to 150 m 195 and to 50 m (individuals m⁻²) when a taxon was present in both the day and night tows (Steinberg 196 et al., 2008). These values are referred to as MOCNESS 150 m N:D and MOCNESS 50 m N:D, 197 respectively. MOCNESS N:D data typically ranged across multiple orders of magnitude and 198 were positively skewed. A relatively few large values were influential on the mean MOCNESS 199 N:D values, typically resulting in large mean values compared to the median (Supplemental 200 Tables 1 and 2). Therefore, the median was used to describe the central tendency of MOCNESS 201 N:D data. Additionally, mean day abundance and mean night abundance were calculated from 202 epipelagic (0-120 m) samples to calculate grid-wide N:D ratios for 1993-2017. These values are 203 referred to as grid-wide 120 m N:D.

204

The vertical distribution of taxa in the MOCNESS discrete depth samples was quantified 205 using weighted mean depth (WMD). WMD (m) is calculated as follows:

206
$$WMD = \sum (n_i \times z_i \times d_i) / \sum (n_i \times z_i)$$

207 where for depth interval *i*, d_i is the midpoint (m), z_i is the interval thickness (m), and n_i is

abundance (no. m⁻³) (Andersen et al., 2001). WMD was only calculated for tows reaching 500 m.

209 Night WMD was subtracted from day WMD to determine the amplitude of diel migration

210 (ΔWMD, m). Data used in the analyses are available at: <u>https://pal.lternet.edu/data</u>.

211 2.5. Environmental controls

212 The environmental water column data used in this analysis was collected at sampling 213 stations where paired day-night MOCNESS tows were conducted. Discrete chlorophyll *a* (chl-*a*) 214 measurements were made fluorometrically (Parsons et al., 1984). Primary productivity rates were measured with 24-h incubations of ¹⁴C uptake at various light levels (Steemann Nielsen, 1952; 215 216 Schofield et al., 2018). Both chl-a and primary production were depth-integrated to 100 m. The 217 depth of the subsurface chl-a maximum (Z_{SCM}) and euphotic zone defined by the 1% isolume 218 $(Z_{1\%})$ were determined with a fluorometer and a photosynthetically active radiation (PAR) 219 sensor, respectively, mounted on the CTD rosette. Mixed layer depth (MLD) was calculated as 220 the depth of maximum buoyancy frequency from the same CTD casts (Carvalho et al., 2017). 221 Photoperiod (hours) was calculated for all day-night MOCNESS tow pairs using latitude and day 222 of year (Kirk, 2011).

223 *2.6. Statistical analyses*

Single-factor ANOVA was used to test for differences in Δ WMD and log-adjusted MOCNESS N:D ratios among the North, South, and Far South sub-regions. The significance level (α) was set at 0.05. There was no significant difference among latitudinal sub-regions for any taxa for MOCNESS 150 m N:D or MOCNESS 50 m N:D (ANOVA; p > 0.06). All statistical tests were conducted with R version 3.3.2 (R Core Team, 2016).

Differences between day and night surface abundance (0-150 m and 0-50 m) from MOCNESS pairs were tested using the Wilcoxon signed-rank test. This non-parametric test does not require transformation of non-normal data and gave comparable results to the paired *t*-test using log-transformed data (Supplemental Tables 1 and 2). Differences between unpaired day and night grid-wide epipelagic abundance (0-120 m) were tested using the Wilcoxon rank-sum test.

235 Multiple linear regression was used to identify environmental controls on Δ WMD for 236 taxa that made DVMs from the mesopelagic zone into the epipelagic zone. Only M. gerlachei 237 and ostracods were included in this analysis; Δ WMD was not a sensitive metric for L. h. 238 antarctica, because it was concentrated in the epipelagic zone, and Salpa thompsoni was 239 excluded due to an insufficient sample size. Δ WMD did not differ among latitudinal sub-regions 240 for *M. gerlachei* or ostracods (ANOVA; p > 0.96). Therefore, data for the analysis were 241 combined across the entire sampling region. 242 MOCNESS 50 m N:D was a sensitive metric for taxa that were concentrated in the 243 epipelagic zone or conducted DVM within the epipelagic zone. These taxa included L. h. 244 antarctica, C. acutus, R. gigas, T. macrura, and E. crystallorophias. Generalized linear models 245 with a gamma distribution and log link function were used to identify environmental controls on 246 MOCNESS 50 m N:D. The gamma distribution is appropriate for ratios, because it is constrained 247 to positive, continuous values. The log link function also ensures positive fitted values. 248 Significant model fits were not achieved for the shallow migrators C. acutus and E. 249 crystallorophias. 250 A suite of nine models was fitted for each individual taxon included in Δ WMD analysis

251 (multiple linear regression) and in MOCNESS 50 m N:D analysis (generalized linear model with

252 gamma distribution and log link function). The water column properties investigated in this study 253 were correlated with one another and therefore were not included in the same models to avoid 254 problematic collinearity. For example, as MLD deepened so did Z_{SCM} (Pearson's r = 0.69; p =255 0.0004). $Z_{1\%}$ deepened as depth-integrated chl-*a* decreased (Pearson's r = -0.60; p = 0.004) and 256 as Z_{SCM} deepened (Pearson's r = 0.44; p = 0.044). None of the water column properties were 257 correlated with photoperiod. Therefore, the nine models included each explanatory variable 258 individually (i.e., photoperiod, chl-a, Z_{SCM} , $Z_{1\%}$, and MLD) as well as photoperiod paired with 259 each of the water column properties. Model selection statistics are presented in Supplemental 260 Tables 3-7. Final models were selected according to the lowest Akaike Information Criterion 261 value corrected for small sample size (AICc) (Hurvich and Tsai, 1989) using the model.sel 262 function in the MuMIn package (Barton, 2016). Presented models satisfied assumptions as 263 verified by plotting residuals versus fitted values and explanatory variables.

3. Results

265 *3.1. Environmental conditions*

Mean photoperiod during MOCNESS sampling was 20 h 11 min (range: 17 h 41 min to 267 21 h 47 min), and mean $Z_{1\%}$, was 47 m (range: 16-81 m). Mean depth-integrated chl-*a* was 126 268 mg m⁻² (range: 13-517 mg m⁻²), and mean depth-integrated primary production was 2489 mg C 269 m⁻² d⁻¹ (range: 605-5354 mg C m⁻² d⁻¹). Mean MLD was 28 m (range: 5-79 m), and mean Z_{SCM} 270 was 20 m (range: 4-60 m).

271 *3.2. Diel vertical depth distributions by taxon*

3.2.1. Calanoid copepods

The calanoid copepod *Metridia gerlachei* was the most abundant taxon in MOCNESS
tows and a strong diel vertical migrator (Fig. 2), with a median MOCNESS 150 m N:D of 8.0

275 (Wilcoxon signed-rank test p = 0.002) (Table 1). Much of the *M. gerlachei* population did not 276 migrate and resided between 300-500 m, particularly in the North (Fig. 2a). The *M. gerlachei* 277 depth distribution was more even in the South and Far South (Fig. 2b-c). Calanoides acutus was 278 the second-most abundant calanoid and although its abundance from 0-150 m did not differ 279 significantly between day and night (Wilcoxon signed-rank test p = 0.64) (Table 1), it was more 280 abundant during night tows from 0-50 m with a median MOCNESS 50 m N:D of 2.3 (Fig. 3d; 281 Table 2) (Wilcoxon signed-rank test p = 0.001). Like *M. gerlachei*, *C. acutus* vertical distribution 282 also varied with latitudinal sub-region. Calanoides acutus was distributed relatively evenly with 283 depth in the North and South (Fig. 3a-b) but was concentrated between 250-400 m in the Far 284 South (Fig. 3c), where it was also an order of magnitude more abundant at this depth zone 285 compared to the other sub-regions.

286 In contrast to M. gerlachei and C. acutus, Calanus propinguus, Rhincalanus gigas, and 287 Paraeuchaeta antarctica were an order of magnitude less abundant and did not vary appreciably 288 with latitudinal sub-region. Calanus propinguus was most abundant in the surface 50 m (Fig. 4a) 289 unlike other calanoid copepods, which had peak abundances in the mesopelagic zone (day and 290 night). Epipelagic C. propinguus abundance did not differ between day and night (Tables 1 and 291 2). Rhincalanus gigas was most abundant from 250-300 m during the day and from 200-250 m at 292 night (Fig. 4b), and abundance in the surface 50 m was significantly greater at night than during 293 the day (Wilcoxon signed-rank test p = 0.004). Median R. gigas MOCNESS 50 m N:D was 36.6 294 (Table 2). Paraeuchaeta antarctica was most abundant from 300-500 m and mostly remained 295 resident in the mesopelagic zone during day and night (Fig 4c). Although scarce in the epipelagic 296 zone, *P. antarctica* was significantly more abundant from 0-50 m during the day (Wilcoxon 297 signed-rank test p = 0.012) (Table 2), which suggests this species conducted reverse DVM.

298	Table 1. Diel, depth-integrated zooplankton abund	ance (0-150 m) from MOC	CNESS tows along the WAP of	continental shelf. For
			Ũ	

- euphausiids and *Salpa thompsoni*, Day and Night n = 22 (samples from 2009-2017). For all other taxa, Day and Night n = 17 (samples
- 300 from 2009-2015). Night:Day *n* varies because taxa were not always present in both day and night paired tows. *p*-values are for the
- 301 Wilcoxon signed-rank test comparing paired day-night abundance values when a taxon was present in both tows.

	Day (individuals m ⁻²)		Night (in	dividuals m ⁻²)	Night:Day			
						25% - 75%		
Taxon	Median	Range	Median	Range	Median	Quantiles	р	n
Calanoid copepods								
Metridia gerlachei	32.7	0.1 - 1975.0	249.5	3.1 - 2860.8	8.0	6.2 - 32.0	0.002	17
Calanoides acutus	37.1	4.2 - 230.6	51.6	7.3 - 269.1	1.0	0.8 - 1.5	0.64	17
Calanus propinquus	4.8	0.0 - 165.4	7.3	1.0 - 75.7	1.7	0.4 - 4.4	0.67	16
Rhincalanus gigas	4.2	0.3 - 20.3	3.9	0.0 - 14.8	1.2	0.5 - 2.3	0.82	16
Paraeuchaeta antarctica	0.9	0.0 - 4.5	1.4	0.0 - 7.1	1.1	0.6 - 2.3	0.68	12
Euphausiids								
Thysanoessa macrura	5.0	0.2 - 60.4	9.9	0.1 - 74.9	1.4	1.1 - 2.5	0.003	22
Euphausia crystallorophias	0.2	0.0 - 18.3	0.1	0.0 - 61.6	0.7	0.5 - 5.6	0.58	11
Other crustaceans								
Ostracoda	3.8	0.0 - 38.2	20.1	1.2 - 219.1	2.9	1.7 - 7.4	0.0003	16
Amphipoda	1.9	0.0 - 8.5	1.3	0.0 - 16.4	0.9	0.6 - 2.1	0.56	15
Gelatinous zooplankton								
Salpa thompsoni [†]	0.0	0.0 - 19.4	0.0	0.0 - 174.4	9.0	4.5 - 9.0	0.75	3
Limacina helicina antarctica	1.2	0.0 - 53.2	3.4	0.0 - 56.1	1.7	1.2 - 3.6	0.021	13
Gymnosomata	0.5	0.0 - 25.3	0.6	0.0 - 3.3	0.8	0.5 - 1.3	0.34	12
Chaetognatha	24.0	0.7 - 118.6	21.8	1.7 - 84.1	1.0	0.6 - 1.6	0.75	17
Tomopteris spp.	0.04	0.0 - 1.1	0.4	0.0 - 1.6	1.2	1.1 - 6.7	0.30	9

[†] See Figure 7b for *Salpa thompsoni* data from the continental slope.

303	Table 2. Diel, depth-integrated zooplankton abundance (0-50 m) from MOCNESS tows along the WAP continental shelf. For	

304 euphausiids and *Salpa thompsoni*, Day and Night n = 22 (samples from 2009-2017). For all other taxa, Day and Night n = 17 (samples

from 2009-2015). Night:Day *n* varies because taxa were not always present in both day and night paired tows. *p*-values are for the

306	Wilcoxon	signed-ran	nk test co	omparing p	aired	day-night	abundance	values w	hen a taxon	was present in	1 both tows.
		0		1 01		5 0				1	

	Day (individuals m ⁻²)		Night (individuals m ⁻²)		Night:Day			
						25% - 75%		
Taxon	Median	Range	Median	Range	Median	Quantiles	р	n
Calanoid copepods								
Metridia gerlachei	4.0	0.0 - 153.8	7.6	0.0 - 948.6	1.8	1.4 - 7.0	0.013	13
Calanoides acutus	9.0	0.0 - 76.0	13.8	0.0 - 155.5	2.3	1.6 - 2.9	0.001	12
Calanus propinquus	0.2	0.0 - 159.5	4.0	0.0 - 62.0	2.1	0.9 - 4.2	0.38	8
Rhincalanus gigas	0.3	0.0 - 4.9	13.8	0.0 - 155.5	36.6	18.2 - 112.4	0.004	9
Paraeuchaeta antarctica	4.0	0.0 - 62.0	0.1	0.0 - 1.6	0.2	0.1 - 0.4	0.012	9
Euphausiids								
Thysanoessa macrura	0.2	0.0 - 54.8	3.1	0.1 - 47.1	11.7	3.6 - 19.4	0.006	16
Euphausia crystallorophias	0.0	0.0 - 4.7	0.1	0.0 - 56.3	8.8	4.3 - 13.1	0.047	7
Other crustaceans								
Ostracoda	1.0	0.0 - 23.3	0.9	0.0 - 24.0	1.3	1.0 - 2.1	0.24	11
Amphipoda	0.3	0.0 - 8.0	0.3	0.0 - 3.2	0.7	0.5 - 2.4	0.85	10
Gelatinous zooplankton								
Salpa thompsoni	0.0	0.0 - 5.4	0.0	0.0 - 63.0	1.2	0.6 - 6.5	0.75	3
Limacina helicina antarctica	0.6	0.0 - 28.4	0.9	0.0 - 31.7	2.8	1.7 - 3.6	0.13	9
Gymnosomata	0.0	0.0 - 25.0	0.1	0.0 - 2.5	1.2	0.4 - 1.3	1.00	6
Chaetognatha	1.9	0.0 - 14.2	1.6	0.0 - 10.4	0.9	0.7 - 1.5	0.68	13
<i>Tomopteris</i> spp.	0.0	0.0 - 1.1	0.0	0.0 - 0.8	0.5	NA	NA	1



Figure 2. Mean day (light gray, left) and night (dark gray, right) abundance of the calanoid

310 copepod *Metridia gerlachei* in the North (a), South (b), Far South (c) sub-regions, and full shelf

- 311 sampling region (d) at discrete depth intervals from 0-500 m. Error bars indicate one standard
- 312 error. North n = 5-7; South n = 5-6; Far South n = 2-4; Full n = 12-17.



Figure 3. Mean day (light gray, left) and night (dark gray, right) abundance of the calanoid copepod *Calanoides acutus* in the North (a), South (b), Far South (c) sub-regions, and full shelf sampling region (d) at discrete depth intervals from 0-500 m. Error bars indicate one standard error. North n = 5-7; South n = 5-6; Far South n = 2-4; Full n = 12-17. Note different scaling on x-axes.



320 Figure 4. Mean day (light gray, left) and night (dark gray, right) abundance of the calanoid

- 321 copepods *Calanus propinquus* (a), *Rhincalanus gigas* (b), and *Paraeuchaeta antarctica* (c)
- 322 sampled at discrete depth intervals from 0-500 m for the full shelf sampling region. Error bars
- 323 indicate one standard error. Full n = 12-17. Note different scaling on x-axes.

3.2.2. Euphausiids

325	The abundant krill species Thysanoessa macrura was concentrated in the epipelagic zone
326	during day and night, but ascended at night, especially into the upper 50 m (Fig. 5a). Some
327	degree of DVM by T. macrura was supported by all metrics tested. This species was
328	significantly more abundant in 0-50 m depths at night compared to day (Wilcoxon signed-rank
329	test $p = 0.006$) (Table 2), and <i>T. macrura</i> median MOCNESS 50 m N:D was 11.7. <i>Thysanoessa</i>
330	macrura was also more abundant at night vs. day in the upper 0-150 m (MOCNESS tows;
331	Wilcoxon signed-rank test $p = 0.003$; median MOCNESS 150 m N:D = 1.4) (Table 1) and from
332	0-120 m during nighttime grid-wide epipelagic tows (Wilcoxon rank-sum test $p = 0.0003$; grid-
333	wide 120 m N:D = 1.6) (Table 3). <i>Euphausia crystallorophias</i> was less abundant than T .
334	macrura, but similarly was concentrated in the upper 100 m (Fig. 5b) and migrated into the top
335	50 m at night, as indicated by higher abundance from 0-50 m at night than day (Wilcoxon
336	signed-rank test $p = 0.047$) and a median MOCNESS 50 m N:D of 8.8 (Table 2). <i>Euphausia</i>
337	superba remained in the epipelagic zone through the diel cycle with a grid-wide 120 m N:D of
338	0.93 (Table 3).

339 3.2.3. Other crustaceans

Ostracods migrated nightly into the upper 200 m (Fig. 6a), with significantly higher abundance from 0-150 m at night vs. day (Wilcoxon signed-rank test p = 0.0003) and a median MOCNESS 150 m N:D of 2.9 (Table 1). Mean ostracod abundance peaked in the 200-250 m layer, where they were about 50% more abundant during day than night (Fig. 6a). Most of the ostracod community did not migrate and resided between 200-500 m throughout the diel cycle. Amphipods were an order of magnitude less abundant than ostracods, with two distinct abundance peaks in the mesopelagic zone during day, and highest abundance from 100-200 m at

- 347 night (Fig. 6b). Amphipods were significantly more abundant in nighttime epipelagic tows and
- had a grid-wide 120 m N:D of 2.1 (Wilcoxon rank-sum test $p = 2.6 \times 10^{-12}$) (Table 3).

- **Table 3.** Diel, depth-integrated zooplankton abundance (0-120 m) from grid-wide epipelagic tows across the PAL LTER sampling
- region from 1993-2017. *n* varies because not all taxa were identified consistently throughout the time series. *p*-values are for the
- 351 Wilcoxon rank-sum test.

	Day (individuals m ⁻²) Night (individuals m ⁻²)							
Taxon	Mean	SE	n	Mean	SE	n	Night:Day	р
Euphausiids								
Thysanoessa macrura	22.9	2.0	1063	36.9	6.8	197	1.6	0.0003
Euphausia crystallorophias	2.9	0.38	1067	6.2	2.3	196	2.1	0.77
Euphausia superba	13.7	2.0	1071	12.7	6.1	198	0.9	0.001
Other crustaceans								
Amphipoda	0.36	0.025	1026	0.75	0.13	185	2.1	2.6 x 10 ⁻¹²
Gelatinous zooplankton								
Salpa thompsoni	4.9	1.4	1069	12.5	3.0	197	2.6	7.7 x 10 ⁻¹⁷
Limacina helicina antarctica	7.3	0.62	1056	9.9	1.1	195	1.4	0.004
Gymnosomata	0.22	0.014	1050	0.31	0.039	193	1.4	0.0002
Chaetognatha	1.5	0.16	994	0.88	0.20	183	0.6	0.088
Tomopteris spp.	0.051	0.0049	966	0.095	0.014	181	1.9	0.0002





Figure 5. Mean day (light gray, left) and night (dark gray, right) abundance of the euphausiids *Thysanoessa macrura* (a) and *Euphausia crystallorophias* (b) sampled at discrete depth intervals from 0-500 m for the full shelf sampling region. Error bars indicate one standard error. Full n =17-22. Note different scaling on x-axes.







362 region. Error bars indicate one standard error. Full n = 12-17. Note different scaling on x-axes.

3.2.4. Salps

364	Salpa thompsoni was a strong diel vertical migrator. Mean abundance over the
365	continental shelf was highest from 200-400 m during the day, and from 0-200 m at night (Fig.
366	7a). Salpa thompsoni median MOCNESS 150 m N:D was 9.0, although salps were only present
367	in three day-night MOCNESS tow pairs along the continental shelf (Table 1). Salps were also
368	significantly more abundant at night in epipelagic tows, with a grid-wide 120 m N:D of 2.6
369	(Wilcoxon rank-sum test $p = 7.7 \times 10^{-17}$) (Table 3). Over the continental slope, <i>S. thompsoni</i>
370	migrated mostly from daytime residence depths in the 200-300 m layer into the surface 100 m at
371	night (Fig. 7b), with MOCNESS 100 m N:D = 94.6 and MOCNESS 200 m N:D = 6.3. Salps
372	were relatively scarce below 300 m, although a small, deep peak occurred from 750-1000 m on
373	the slope.

374 3.2.5. Pteropods

375 The thecosome (shelled) pteropod Limacina helicina antarctica was concentrated in 376 surface waters but also migrated from 150-250 m during the day into the upper epipelagic zone 377 at night (Fig. 8a). This result is supported by their higher abundance in the upper 150 m at night 378 from MOCNESS tows (Wilcoxon signed-rank test p = 0.021) (Table 1) and from 0-120 m at 379 night in epipelagic tows (Wilcoxon rank-sum test p = 0.004) (Table 3). Median L. h. antarctica 380 MOCNESS 150 m N:D was 1.7 (Table 1), and grid-wide 120 m N:D was 1.4 (Table 3). 381 Gymnosome (shell-less) pteropods were less abundant than L. h. antarctica in the epipelagic 382 zone and were distributed relatively evenly with depth, with highest mean gymnosome 383 abundance from 0-50 m during the day (Fig. 8b). However, DVM by gymnosomes is indicated 384 grid-wide, with significantly higher abundance at night in the epipelagic zone and a grid-wide 385 120 m N:D of 1.4 (Wilcoxon rank-sum test p = 0.0002) (Table 3).



Figure 7. Mean day (light gray, left) and night (dark gray, right) abundance of *Salpa thompsoni* from the full continental shelf sampling region (a) and a single pair of tows on the continental slope (b) sampled at discrete depth intervals from 0-500 m and 0-1000 m, respectively. Error bars indicate one standard error. Full Shelf n = 17-22; Slope n = 1. Note different scaling on yaxes.

392





intervals from 0-500 m for the full shelf sampling region. Error bars indicate one standard error.

397 Full n = 12-17. Note different scaling on x-axes.

398 3.2.6. Gelatinous carnivores

399 Chaetognaths and *Tomopteris* spp. polychaetes were mostly resident in the mesopelagic 400 zone and relatively scarce from 0-100 m (Fig. 9). Mean abundance of both taxa was highest from 401 200-250 m during the day and from 150-200 m at night (Tomopteris spp. also had a second night 402 peak at 300-400 m) (Fig. 9). Median chaetognath MOCNESS 150 m N:D was 1.0 (Table 1) and 403 although grid-wide 120 m N:D was 0.6, epipelagic abundance did not differ between day and 404 night (Tables 1-3), suggesting chaetognaths did not undergo DVM. However, *Tomopteris* spp. 405 polychaetes did appear to undergo DVM as they were significantly more abundant during night 406 epipelagic tows and had a grid-wide 120 m N:D of 1.9 (Wilcoxon rank-sum test p = 0.0002) 407 (Table 3).



409 **Figure 9.** Mean day (light gray, left) and night (dark gray, right) abundance of chaetognaths (a) 410 and *Tomopteris* spp. polychaetes (b) sampled at discrete depth intervals from 0-500 m for the full 411 shelf sampling region. Error bars indicate one standard error. Full n = 12-17. Note different 412 scaling on x-axes.

3.3. Environmental controls on DVM

414	For strong migrators traveling between the mesopelagic and epipelagic zones, migration
415	distance (i.e., Δ WMD) was sensitive to photoperiod and vertical water column structure. For the
416	copepod <i>M. gerlachei</i> , photoperiod and Z_{SCM} best explained ΔWMD (Table 4), with <i>M</i> .
417	gerlachei making shorter vertical migrations as photoperiod grew longer and when Z_{SCM} was
418	shallower (Fig. 10a). Similarly, photoperiod and MLD best explained ostracod Δ WMD (Table
419	4). Ostracods made shorter DVMs as photoperiod grew longer and when MLD was deeper (Fig.
420	10b; note- an outlier that was excluded prior to model selection for ostracods is included for
421	visualization in this figure).
422	For taxa making shorter-distance DVMs mostly within the epipelagic zone, the
423	magnitude of DVM into the surface layer (i.e., MOCNESS 50 m N:D) was best explained by
424	phytoplankton abundance and distribution. The final models each included a different, single
425	explanatory variable for the copepod R. gigas, euphausiid T. macrura, and pteropod L. h.
426	antarctica. Rhincalanus gigas DVM into the surface 50 m decreased as depth-integrated chl-a
427	increased (Fig. 11a; Table 5). Thysanoessa macrura DVM decreased when MLD was deeper
428	(Fig. 11b), and similarly L. h. antarctica DVM decreased when Z_{SCM} was deeper (Fig. 11c – a
429	finding robust to the inclusion of an outlier value – see inset) (Table 5).

Table 4. Statistics from multiple linear regression models assessing the impact of environmental variables on zooplankton Δ WMD

	431	from MOCNESS tows	(0-500 m) along the	WAP	continental	shelf from	2009-201
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Variable	n	Coefficient	SE	t	р	Partial R ²
<i>Metridia gerlachei</i> Δ WMD (adjusted R ² = 0.76; <i>p</i> = 0.001)	11					
Photoperiod		-19.8	4.6	-4.3	0.003	0.70
Depth of chl- <i>a</i> maximum		2.27	0.47	4.8	0.001	0.75
Intercept		400.0	92.5	4.3	0.003	
Ostracoda Δ WMD (adjusted R ² = 0.79; <i>p</i> = 0.002)	10					
Photoperiod		-12.1	3.0	-4.1	0.005	0.70
Mixed layer depth		-0.666	0.23	-2.9	0.022	0.55
Intercept		283.8	59.7	4.8	0.002	

Table 5. Statistics from generalized linear models (with a gamma distribution and log link function) assessing the impact of

434 environmental variables on zooplankton 50m N:D from MOCNESS tows along the WAP continental shelf from 2009-2015 (for

435	Rhincalanus gigas and Limacina helicina antarctica) and 2009-2017 (for Thysanoessa macrura).	

Variable	n	Coefficient	SE	t	р
<i>Rhincalanus gigas</i> 50m N:D (deviance explained = 35%)	8				
Chl-a concentration		-0.0088	0.0031	-2.8	0.031
Intercept		5.15	0.41	12.6	1.5 x 10 ⁻⁵
<i>Thysanoessa macrura</i> 50m N:D (deviance explained = 27%)	15				
Mixed layer depth		-0.035	0.011	-3.1	0.008
Intercept		3.67	0.42	8.7	8.8 x 10 ⁻⁷
<i>Limacina helicina antarctica</i> 50m N:D (deviance explained = 40%)	8				
Depth of chl- <i>a</i> maximum		-0.055	0.020	-2.7	0.035
Intercept		2.06	0.61	3.4	0.015



Figure 10. Environmental controls on Δ WMD. (a) Depth of chlorophyll *a* maximum versus *Metridia gerlachei* Δ WMD. (b) Photoperiod versus Ostracod Δ WMD. Solid lines indicate the linear regression with all *M. gerlachei* data points and without the ostracod outlier value, indicated by an open diamond. *M. gerlachei*: n = 11, p = 0.048, $R^2 = 0.30$; Ostracod (without outlier): n = 10, p = 0.005, $R^2 = 0.60$.



445 Figure 11. Environmental controls on 50 m N:D. (a) Depth-integrated chlorophyll *a*446 concentration versus *Rhincalanus gigas* 50 m N:D. (b) Mixed layer depth versus *Thysanoessa*

447 *macrura* 50 m N:D. (c) Depth of chlorophyll *a* maximum versus *Limacina helicina antarctica* 50

448 m N:D. Inset includes an outlier *L. h. antarctica* 50 m N:D value indicated by an open diamond.

449 Solid line indicates generalized linear model fit. Dashed lines indicate one standard error. *R*.

450 *gigas*: n = 8, p = 0.031, deviance explained = 35%; *T. macrura*: n = 15, p = 0.008, deviance

451 explained = 27%; *L. h. antarctica* (without outlier): n = 8, p = 0.035, deviance explained = 40%.

453 **4. Discussion**

454 *4.1. Zooplankton DVM modes*

455

4.1.1. DVM between epipelagic and mesopelagic zones

456 Four taxa performed consistent DVM between the mesopelagic zone during day and 457 epipelagic zone at night. The copepod *Metridia gerlachei* migrated into the upper 100 m at night, 458 consistent with results from prior studies in the northern WAP (Hopkins, 1985; Lopez and 459 Huntley, 1995; King and LaCasella, 2003). Similarly, a portion of the ostracod community made 460 relatively extensive (~100 m) DVMs resulting in a 21% decrease in abundance from 200-300 m 461 and a 3.5-fold increase in abundance from 0-200 m at night. Ostracod DVM is well-documented 462 in the Atlantic and Pacific Oceans (Angel, 1979; Steinberg et al., 2008). Population-wide DVM 463 by Salpa thompsoni from 300 m into surface waters supports previous observations throughout 464 the Southern Ocean during summer (Piatkowski, 1985; Casareto and Nemoto, 1986; Perissinotto 465 and Pakhomov, 1998; Pakhomov et al., 2011). Although rarely encountered deeper than 300 m 466 during summer, the pteropod *Limacina helicina antarctica* underwent DVM between the 467 epipelagic and upper mesopelagic zones. In the Lazarev Sea, L. h. antarctica also conducted 468 DVM from November to February (Hunt et al., 2008). Collectively, M. gerlachei, ostracods, S. 469 thompsoni, and L. h. antarctica constitute an assemblage of strong vertical migrators along the 470 WAP during summer.

Evidence for amphipod, gymnosome pteropod, and *Tomopteris* spp. polychaete DVM was less consistent than for the above species, but these taxa were each more abundant at night compared to day in grid-wide epipelagic tows. All amphipod species were grouped together, but DVM is likely species-specific. For example, the hyperiid amphipod *Cyllopus lucasii* was more abundant through the upper 200 m at night during summer, autumn, and winter in the Lazarev

476 Sea where there was no evidence for DVM by the hyperiella dilatata and Primno 477 macropa (Flores et al., 2014). Themisto gaudichaudii (synonym Parathemisto gaudichaudii) is 478 abundant along the WAP (Steinberg et al., 2015), and this amphipod made DVMs from ~200 m 479 to the surface 50 m in the Atlantic Ocean (Williams and Robins, 1981). Prior evidence for Clione 480 antarctica and Spongiobranchia australis DVM is inconsistent (Hunt et al., 2008), but our 481 epipelagic day-night abundance data suggest these gymnosome pteropods conduct DVM, likely 482 to feed on their primary prey L. h. antarctica (Lalli and Gilmer, 1989; Van der Spoel and Dadon, 483 1999) in the epipelagic zone at night. At night, *Tomopteris* spp. polychaete abundance decreased 484 37% from 200-300 m and increased 3-fold in the surface 0-200 m. We suggest a portion of the 485 amphipod, gymnosome, and *Tomopteris* spp. assemblage conducted DVM to feed in the upper 486 200 m while other individuals remained at depth.

487

4.1.2. DVM within epipelagic zone

488 The copepods *Calanoides acutus* and *Rhincalanus gigas* made shallow DVMs from the 489 50-100 m layer into the upper 50 m at night. Shallow DVMs within the upper 70 m for *C. acutus*

490 and upper 90 m for *R. gigas* were also reported in January near South Georgia (Atkinson et al.,

491 1992a, 1992b). A study in the Drake Passage and northern Antarctic Peninsula found no *C*.

492 *acutus* DVM during December to March (Huntley and Escritor, 1991), but was limited to vertical

493 resolution of 0-100 and 100-200 m, making it unlikely to detect shallow DVM.

494 The krill species *Thysanoessa macrura* and *Euphausia crystallorophias* performed

shallow DVM. DVM within the epipelagic zone was reported during spring and autumn further

- 496 north of our study site for *T. macrura* (Loeb and Shulenberger, 1987; Nordhausen, 1994a) and *E.*
- 497 crystallorophias (using acoustics; Everson, 1987). Summer surveys in the northern WAP
- 498 (Nordhausen, 1992) and Amundsen Sea (La et al., 2015b) did not detect DVM by T. macrura

and *E. crystallorophias*, respectively, possibly due to limitations of sampling methods in
detecting shallow DVM. Net avoidance by the larger, faster *E. crystallorophias* was not apparent
in the northern WAP during winter (Nordhausen et al. 1994b) but was during autumn (Everson,
1987), which could exaggerate the shallow DVM signal and contribute to higher nighttime
abundance depicted in Figure 5b.

504 4.1.3.

4.1.3. DVM within mesopelagic zone

505 Chaetognaths and the copepod R. gigas both undertook a modest DVM within the 506 mesopelagic zone between 150 and 300 m, possibly indicating predator-prey coupling. The 507 chaetognaths *Eukronia hamata* and *Sagitta gazellae* predominately fed on the copepods C. 508 acutus, C. propinguus, and M. gerlachei in the upper 200 m of the Weddell Sea in autumn 509 (Hopkins and Torres, 1989). Chaetograths along the WAP mainly remained in deeper layers 510 during summer to feed on abundant copepod prey. Although less numerous, R. gigas is a larger 511 (Gleiber, 2014), and perhaps preferable, copepod prey item compared to *M. gerlachei* or *C.* 512 acutus.

513 *4.1.4. Reverse DVM*

The copepod *Paraeuchaeta antarctica* made relatively short reverse DVMs, from 0-50 m during the day to 50-100 m at night. The primarily carnivorous *P. antarctica* is the largest copepod in this study (mean adult prosome length 7 mm; Gleiber, 2014), making it particularly vulnerable to visual predators in surface waters (Aksnes and Giske, 1993; Ohman and Romagnan, 2016). Reverse DVM is adaptive for species susceptible to predators that undertake normal DVM (Ohman et al., 1983). The reverse DVM of *P. antarctica* is likely used to avoid vertically migrating visual predators.

521 *4.2. Non-migrating zooplankton*

4.2.1. Epipelagic non-migrators

The copepod *Calanus propinquus* was concentrated in the upper 50 m and did not consistently undertake DVM. *Calanus propinquus* feeds omnivorously (Atkinson, 1998; Pasternak and Schnack-Schiel, 2001) and was concentrated in the upper 100 m year-round in the Scotia Sea (Atkinson and Sinclair, 2000). Therefore, it appears *C. propinquus* typically remains resident in surface waters on both seasonal and diel time scales.

528 The negligible difference between night and day E. superba abundance in our 120 m 529 tows was expected since this depth was selected to collect Antarctic krill across its main summer 530 depth range (Ross et al., 1996). Acoustic studies have documented sporadic DVM within the upper 100 m during summer while DVM is more pronounced during spring and autumn 531 532 (Everson, 1983; Godlewska and Klusek, 1987; Demer and Hewitt, 1995; Ross et al., 1996). The 533 daytime formation of larger schools and nighttime dispersal into smaller schools may be a more 534 consistent predator avoidance behavior for E. superba in the summer (Everson, 1983; Zhou and 535 Dorland, 2004; Tarling et al., 2018).

536

4.2.2. Mesopelagic carnivores and detritivores

537 The copepod *P. antarctica*, ostracods, chaetognaths, and *Tomopteris* spp. polychaetes 538 were concentrated in the mesopelagic zone regardless of the diel cycle, and together compose a 539 deep carnivorous and detritivorous assemblage. The carnivorous P. antarctica (synonym 540 *Euchaeta antarctica*) preved mainly on other copepods in the Weddell Sea (Hopkins and Torres, 541 1989) and near South Georgia where feeding continued through winter (Øresland and Ward, 542 1993). Thus, *P. antarctica* likely remains resident in the mesopelagic zone where metazoan prey 543 is sufficiently abundant throughout the year. While vertically migrating ostracods feed in 544 productive surface waters, the more numerous mesopelagic residents feed as carnivores or

545	detritivores (Angel, 1972; Lampitt et al., 1993; Vannier et al., 1998). Elevated ostracod
546	abundance from 200-500 m was also observed in the northern WAP during summer
547	(Blachowiak-Samolyk and Żmijewska, 1997) and in Marguerite Bay during autumn (Marrari et
548	al., 2011). Chaetognaths along the WAP likely remained in deeper layers during summer to feed
549	on abundant copepod prey as previously discussed. Chaetognaths were the numerically dominant
550	macrozooplankton in the mesopelagic zone throughout the year in the Lazarev Sea (Flores et al.,
551	2014) and are similarly important in the WAP mesopelagic zone. Small Tomopteris spp.
552	polychaetes in the epipelagic zone fed on phytoplankton in the northern WAP during summer
553	(Phleger et al., 1998) and during autumn in the Weddell Sea (Hopkins and Torres, 1989).
554	However, Tomopteris spp. polychaetes in the mesopelagic zone are carnivores or detritivores
555	(Steinberg et al., 1994; Jumars et al., 2015). In particular, individuals larger than 20 mm are
556	primarily carnivorous (Jumars et al. 2015), and large Tomopteris spp. individuals in our study
557	exceeded 60 mm, further supporting their role as carnivores in the mesopelagic zone.
558	Amphipods and gymnosome pteropods were distributed throughout the water column,
559	with a substantial portion of the population residing in the mesopelagic zone during day and
560	night. A diet study in the Weddell Sea during March found multiple amphipod species were
561	feeding mainly on copepods, larval euphausiids, and gelatinous zooplankton (Hopkins and
562	Torres, 1989). Therefore, carnivory explains increased amphipod density below 100 m where
563	metazoan prey is abundant. Although the highest gymnosome abundance was in surface waters,
564	their consistent mesopelagic presence suggests gymnosomes consume other prey in addition to
565	the shelled pteropod L. h. antarctica. A genetic diet analysis of the Arctic Clione limacina found
566	this species fed on amphipods and calanoid copepods in addition to shelled pteropods (Kallevik,

567 2013). Similar to other mesopelagic zooplankton, amphipods and gymnosome pteropods likely568 have multiple feeding modes.

569

4.2.3. Seasonal vertical migrators

570 The copepods *M. gerlachei*, *R. gigas*, and *C. acutus* all make seasonal vertical 571 migrations, although M. gerlachei and R. gigas feed through winter while C. acutus enters 572 diapause at depth once it has acquired sufficient lipid reserves (Atkinson, 1998; Pasternak and 573 Schnack-Schiel, 2001; Schnack-Schiel, 2001). Elevated mesopelagic concentrations for these 574 species in our study are more similar to autumn or winter depth distributions farther north in the 575 Scotia Sea (Atkinson and Sinclair, 2000; Ward et al., 2012). High M. gerlachei and R. gigas 576 concentrations from 200-500 m are likely indicative of carnivorous and detritivorous feeding, 577 which may be more important along the WAP where the productive season is shorter vs. lower 578 latitudes. High mesopelagic abundance indicates C. acutus adults were likely in diapause and 579 had not yet fully begun their ascent (Atkinson and Shreeve, 1995) in the Far South where we 580 sampled pre-bloom conditions and mean depth-integrated chl-a was only 38 mg m⁻². Reduced 581 seasonal sea ice coverage coincident with increasing phytoplankton biomass in the PAL LTER 582 study area (Stammerjohn et al., 2008; Montes-Hugo et al., 2009) may result in earlier ascents for 583 seasonally migrating copepods.

Indications that the pteropod *L. h. antarctica* conducts a seasonal vertical migration to feed in WAP surface waters during summer and overwinter at depth include that this species was concentrated from 0-100 m during our sampling but was most abundant from 100-200 m during autumn in Marguerite Bay (Marrari et al., 2011). Furthermore, *L. h. antarctica* is typically absent from a moored sediment trap sampling at 170 m on the WAP shelf during summer but commonly collected from June to October (Thibodeau et al., in review). A seasonal vertical

590 migration for the closely related *Limacina helicina helicina* is also suggested in the Arctic Ocean

591 (Kobayashi, 1974).

592 4.3. Environmental controls on DVM

593 *4.3.1. DVM between epipelagic and mesopelagic zones*

594 The copepod *M. gerlachei* and ostracods made shorter DVMs as photoperiod increased

from 17 to 22 h, consistent with previous work demonstrating the sensitivity of polar

596 zooplankton to seasonal irradiance cycles. Acoustic studies throughout the Southern Ocean

597 (64°S to 74°S) suggest that DVM ceases during summer (Cisewski et al., 2010; Cisewski and

598 Strass, 2016; Picco et al., 2017). In these studies, sound scattering layers remained in surface

599 waters around-the-clock rather than descending during the day. However, our results show some 600 taxa continue to migrate between the epipelagic and mesopelagic zones, responding to relatively 601 small changes in photoperiod during Antarctic summer from 64°S to 70°S.

602 Metridia gerlachei made DVMs over a greater depth range when Z_{SCM} was deeper, likely due to changing phytoplankton availability and predation risk. Deeper Z_{SCM} was associated with 603 604 reduced light attenuation. With a deep Z_{SCM} , *M. gerlachei* encountered maximum phytoplankton 605 concentrations farther from the surface at night, but DVM distance increased as *M. gerlachei* 606 migrated deeper to avoid visual predators during day. This finding is consistent with previous 607 work in the southern California Current, where reduced light attenuation was associated with 608 longer DVM distances – particularly deeper daytime depths – for migrating copepods (Ohman 609 and Romagnan, 2016). Metridia gerlachei (mean prosome length 3 mm; Gleiber, 2014) is within 610 the size range of the strongest vertical migrators in that study, and therefore may be similarly 611 susceptible to visual predators. Z_{SCM} thus influences *M. gerlachei* DVM distance directly by

612 concentrating prey distribution and indirectly by modulating predation threat (i.e., via light613 attenuation).

614 Ostracods made shorter DVMs when MLD was deeper, likely because migrating 615 ostracods ascended from the mesopelagic zone at night until reaching elevated phytoplankton 616 concentrations in the mixed layer. A shallow mixed layer results in a longer nighttime feeding 617 ascent and a longer return to mesopelagic daytime residence depth. In an acoustic study during 618 Arctic autumn, migrating zooplankton sound scattering layers were coincident with, or just 619 below, the MLD at midnight (Berge et al., 2014). Deeper MLD is associated with reduced sea ice 620 coverage along the WAP (Schofield et al., 2018), which may result in shorter ostracod DVMs 621 under future regional climate conditions.

622

4.3.2. DVM within epipelagic zone

623 Depth-integrated chlorophyll a concentration influenced DVM by the copepod R. gigas 624 within the epipelagic zone. *Rhincalanus gigas* remained in the surface 0-50 m during day and 625 night to feed on elevated phytoplankton biomass, as indicated by decreasing N:D with increasing 626 chl-a. Gut content analysis showed R. gigas feeds primarily on phytoplankton during summer 627 (Pasternak and Schnack-Schiel, 2001). Elevated chl-a was also associated with increased light 628 attenuation, reducing the susceptibility of *R. gigas* to visual predators in surface waters and 629 limiting any benefit gained by daytime migration out of the upper 50 m. Future long-term 630 declines in regional sea ice coverage, upper ocean stability, and chl-a concentration in the PAL 631 LTER study region (Montes-Hugo et al., 2009; Brown et al., 2019) may increase the amplitude 632 of R. gigas DVM.

633 The 0-50 m N:D ratio for the euphausiid *T. macrura* and for the pteropod *L. h. antarctica* 634 increased with shallower MLD and Z_{SCM} , respectively, as these taxa appeared to cue on vertical

635 phytoplankton distribution. MLD and Z_{SCM} were positively correlated in our study. We suggest 636 increased 0-50 m N:D indicates a larger portion of the population migrated into the upper 50 m 637 at night when phytoplankton was concentrated near the surface. When MLD and Z_{SCM} were 638 deeper, T. macrura and L. h. antarctica DVM into the upper 50 m likely decreased because 639 phytoplankton availability and predator avoidance were both maximized below the 50 m 640 threshold of our sampling resolution. During autumn, Arctic zooplankton sound scattering layers 641 migrated to the MLD at midnight (Berge et al., 2014). Migrating pteropods and copepods 642 ascended to the Z_{SCM} at night during Arctic summer (Daase et al., 2016). Deeper MLD under 643 reduced sea ice conditions along the WAP (Schofield et al., 2018) may result in deeper nighttime 644 distributions for T. macrura euphausiids and L. h. antarctica pteropods.

645 *4.4. Zooplankton-mediated carbon export*

646 Zooplankton vertical structure and behavior play key roles in mediating carbon export 647 (Steinberg and Landry, 2017; Cavan et al., 2019), and regional, taxon-specific data are needed to 648 accurately model the contribution of zooplankton DVM to the global biological carbon pump 649 (e.g., Aumont et al., 2018; Archibald et al., 2019). DVM between the epipelagic and mesopelagic 650 zones by WAP zooplankton through Antarctic summer likely results in substantial active carbon 651 transport out of the euphotic zone, which may help resolve surprisingly low regional particle 652 export to primary production ratios (Stukel et al., 2015; Ducklow et al., 2018). Zooplankton 653 DVM is an important control on POC export in the Scotia Sea (Cavan et al., 2015; Liszka et al., 654 2019), and likely plays a similar role along the WAP where zooplankton fecal pellets constitute 655 67% of summer POC flux at 170 m (Gleiber et al., 2012). Future work also should estimate 656 active transport of dissolved carbon (i.e., respiration of CO₂ and excretion of DOC at depth). 657 Additionally, seasonal vertical migrators (e.g., C. acutus copepods and L. h. antarctica

658 pteropods) transfer carbon to the ocean interior as they respire and die in the mesopelagic zone, 659 but this "lipid pump" (for copepods) is yet to be quantified in the Southern Ocean (Kobari et al., 660 2008; Jónasdóttir et al., 2015). Abundant mesopelagic zooplankton also consume sinking detritus 661 and produce fecal pellets, which regulates POC availability to mesopelagic and benthic food 662 webs (Wilson et al., 2008; Belcher et al., 2017). Our findings show it will be essential to consider 663 species feeding ecology and variable DVM amplitudes when assessing zooplankton roles in 664 Southern Ocean carbon cycling. Finally, documented long-term changes in WAP zooplankton 665 composition (e.g., Steinberg et al., 2015; Thibodeau et al., 2019) as well as phytoplankton 666 biomass and vertical distribution (e.g., Brown et al., 2019) will alter zooplankton-mediated 667 export pathways.

- 668 Author contributions
- 669 John A. Conroy: Conceptualization, Investigation, Formal analysis, Visualization, Writing -
- 670 original draft. Deborah K. Steinberg: Conceptualization, Methodology, Investigation, Writing -
- 671 review & editing, Funding acquisition. Patricia S. Thibodeau: Conceptualization, Investigation,
- 672 Writing review & editing. Oscar Schofield: Conceptualization, Investigation, Writing –
- 673 review & editing.

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1017	Supplemental Table 1. Diel, depth-integrated zooplankton abundance (0-150 m) from MOCNESS tows along the WAP continental
1018	shelf. For euphausiids and <i>Salpa thompsoni</i> , Day and Night $n = 22$ (samples from 2009-2017). For all other taxa, Day and Night $n = 22$
1019	17 (samples from 2009-2015). Night:Day <i>n</i> varies because taxa were not always present in both day and night paired tows. <i>p</i> -values
1020	are for the paired <i>t</i> -test comparing log-transformed day-night abundance values when a taxon was present in both tows.

	Day (individuals m ⁻²) Night (individuals m ⁻²)		Night:Day					
Taxon	Mean	SE	Mean	SE	Mean	SE	<i>p t</i> -test	n
Calanoid copepods								
Metridia gerlachei	168.7	114.0	494.0	169.3	80.2	45.1	0.00004	17
Calanoides acutus	71.8	17.1	74.5	17.5	1.5	0.5	0.64	17
Calanus propinquus	21.2	9.9	15.5	5.0	3.0	0.9	0.47	16
Rhincalanus gigas	5.9	1.4	5.5	1.1	2.9	1.2	0.68	16
Paraeuchaeta antarctica	1.3	0.3	1.8	0.4	4.0	2.1	0.39	12
Euphausiids								
Thysanoessa macrura	10.3	3.3	15.1	3.5	5.0	2.2	0.024	22
Euphausia crystallorophias	2.0	1.0	3.9	2.8	3.5	1.4	0.56	11
Other crustaceans								
Ostracoda	9.3	2.9	38.4	13.0	7.5	2.7	0.0002	16
Amphipoda	2.4	0.5	3.6	1.1	2.5	1.2	0.78	15
Gelatinous zooplankton								
Salpa thompsoni	1.6	1.1	8.4	7.9	6.0	3.0	0.76	3
Limacina helicina antarctica	7.5	3.6	10.1	3.5	3.5	1.1	0.008	13
Gymnosomata	2.6	1.5	1.0	0.3	1.4	0.5	0.40	12
Chaetognatha	31.7	7.5	28.1	5.4	1.6	0.4	0.90	17
<i>Tomopteris</i> spp.	0.25	0.1	0.5	0.1	5.1	2.7	0.14	9

1022	Supplemental Table 2. Diel, depth-integrated zooplankton abundance (0-50 m) from MOCNESS tows along the WAP continental
1023	shelf. For euphausiids and <i>Salpa thompsoni</i> , Day and Night $n = 22$ (samples from 2009-2017). For all other taxa, Day and Night $n = 22$
1024	17 (samples from 2009-2015). Night:Day <i>n</i> varies because taxa were not always present in both day and night paired tows. <i>p</i> -values
1025	are for the paired <i>t</i> -test comparing log-transformed day-night abundance values when a taxon was present in both tows.

	Day (individuals m ⁻²)		Night (individuals m ⁻²)		Night:Day			
Taxon	Mean	SE	Mean	SE	Mean	SE	<i>p t</i> -test	n
Calanoid copepods								
Metridia gerlachei	29.2	10.9	127.4	64.4	889.5	884.0	0.044	13
Calanoides acutus	19.2	5.4	34.7	10.6	2.4	0.3	0.000	12
Calanus propinquus	15.6	9.6	10.4	4.2	14.7	12.6	0.25	8
Rhincalanus gigas	1.2	0.4	1.6	0.6	79.4	30.3	0.000	9
Paraeuchaeta antarctica	10.4	4.2	0.4	0.1	0.3	0.1	0.004	9
Euphausiids								
Thysanoessa macrura	3.4	2.5	8.1	2.3	17.3	5.1	0.000	16
Euphausia crystallorophias	0.3	0.2	3.2	2.5	9.9	3.2	0.034	7
Other crustaceans								
Ostracoda	4.1	1.7	5.0	1.8	2.3	0.9	0.53	11
Amphipoda	0.9	0.5	0.7	0.2	1.5	0.4	0.66	10
Gelatinous zooplankton								
Salpa thompsoni	0.5	0.3	2.9	2.9	4.3	3.7	0.80	3
Limacina helicina antarctica	3.5	1.7	6.0	2.3	8.1	5.9	0.21	9
Gymnosomata	1.7	1.5	0.5	0.2	0.9	0.3	0.35	6
Chaetognatha	2.9	0.9	2.6	0.8	1.8	0.6	0.44	13
<i>Tomopteris</i> spp.	0.06	0.1	0.1	0.1	0.5	NA	NA	1

1027 Supplemental Table 3. Summary of model selection statistics from multiple linear regression models assessing the impact of

- 1028 environmental variables on Metridia gerlachei ΔWMD from MOCNESS tows (0-500 m) along the WAP continental shelf from 2009-
- 1029 2015. Statistics AICc: corrected Akaike Information Criterion; ΔAICc: difference from lowest AICc; AICc weight relative model
- 1030 support or probability. Explanatory variables Photoperiod: day length; Z_{SCM}: depth of subsurface chlorophyll *a* maximum; MLD:
- 1031 mixed layer depth; Chl: chlorophyll *a* concentration depth-integrated to 100 m; $Z_{1\%}$: depth of 1% surface irradiance.

Explanatory variables	AICc	ΔAICc	AICc weight
Photoperiod + Z _{SCM}	101.3	0.00	0.967
Z _{SCM}	109.3	7.95	0.018
Photoperiod	111.2	9.83	0.007
MLD	113.9	12.60	0.002
Chl	114.1	12.72	0.002
Photoperiod + MLD	114.2	12.85	0.002
Z _{1%}	114.3	12.99	0.001
Photoperiod + Z _{1%}	116.4	15.05	0.001
Photoperiod + Chl	116.4	15.07	0.001

1033 Supplemental Table 4. Summary of model selection statistics from multiple linear regression models assessing the impact of

1034 environmental variables on Ostracoda Δ WMD from MOCNESS tows (0-500 m) along the WAP continental shelf from 2009-2015.

Explanatory variables	AICc	ΔAICc	AICc weight
Photoperiod + MLD	85.2	0.00	0.666
Photoperiod	87.1	1.96	0.250
MLD	91.3	6.13	0.031
Photoperiod + Chl	92.1	6.91	0.021
Photoperiod + Z _{SCM}	92.8	7.69	0.014
Photoperiod + Z _{1%}	93.0	7.84	0.013
$Z_{1\%}$	97.2	12.09	0.002
Z _{SCM}	97.3	12.18	0.002
Chl	97.4	12.22	0.001

1035 Statistics and explanatory variables as defined for Supplementary Table 3.

- 1037 Supplemental Table 5. Summary of model selection statistics from generalized linear models (with a gamma error distribution and
- 1038 log link function) assessing the impact of environmental variables on Rhincalanus gigas MOCNESS 50 m N:D along the WAP
- 1039 continental shelf from 2009-2015. Statistics and explanatory variables as defined for Supplementary Table 3.

Explanatory variables	AICc	ΔAICc	AICc weight
Chl	95.7	0.00	0.386
Z _{SCM}	96.6	0.88	0.248
Z _{1%}	97.3	1.56	0.177
MLD	97.9	2.22	0.127
Photoperiod	99.7	3.99	0.052
Photoperiod + Chl	105.0	9.29	0.004
Photoperiod + Z _{SCM}	105.8	10.10	0.002
Photoperiod + Z _{1%}	106.0	10.32	0.002
Photoperiod + MLD	106.7	11.00	0.002

- 1041 Supplemental Table 6. Summary of model selection statistics from generalized linear models (with a gamma error distribution and
- 1042 log link function) assessing the impact of environmental variables on *Thysanoessa macrura* MOCNESS 50 m N:D along the WAP
- 1043 continental shelf from 2009-2017. Statistics and explanatory variables as defined for Supplementary Table 3.

Explanatory variables	AICc	ΔAICc	AICc weight
MLD	117.8	0.00	0.449
Photoperiod + MLD	119.2	1.44	0.218
Z1%	120.7	2.95	0.103
Chl	121.7	3.96	0.062
Photoperiod	122.0	4.19	0.055
Z _{SCM}	122.4	4.64	0.044
Photoperiod + Chl	123.0	5.19	0.033
Photoperiod + Z _{SCM}	124.0	6.24	0.020
Photoperiod + Z _{1%}	124.0	6.59	0.017

1045 Supplemental Table 7. Summary of model selection statistics from generalized linear models (with a gamma error distribution and

- 1046 log link function) assessing the impact of environmental variables on *Limacina helicina antarctica* MOCNESS 50 m N:D along the
- 1047 WAP continental shelf from 2009-2015. Statistics and explanatory variables as defined for Supplementary Table 3.

Explanatory variables	AICc	ΔAICc	AICc weight
Z _{SCM}	35.8	0.00	0.627
Photoperiod	39.4	3.55	0.106
Chl	39.7	3.87	0.091
MLD	39.8	3.96	0.087
$Z_{1\%}$	40.0	4.19	0.077
Photoperiod + Z _{SCM}	44.2	8.41	0.009
Photoperiod + MLD	48.4	12.57	0.001
Photoperiod + Chl	48.5	12.70	0.001
Photoperiod + Z _{1%}	48.7	12.86	0.001