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1	Cross-shelf structure and distribution of mesozooplankton communities in the East-
2	Siberian Sea and the adjacent Arctic Ocean
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12	Abstract The East-Siberian Sea (ESS) plays a significant role in circulation of the surface

water and biological production in the Arctic, yet due to its remote location and historically 13 difficult sampling conditions remains the most understudied of all Arctic shelf seas, with 14 15 even baseline information on biological communities absent in literature. We contribute to 16 such a baseline by describing the distribution and community structure of 17 mesozooplankton in the ESS and the adjacent Arctic Ocean based on recent (September 18 2009, 2015) as well as historical (August-September 1946, 1948) data. We found that the 19 overall biomass and abundance during our studies were significantly lower than in the 20 adjacent Chukchi Sea, but higher than historical estimates from ESS, around 25-35 mg DW 21 m⁻³. The diversity was low and characteristic for other Arctic shelf seas, with increasing 22 number of species in deeper waters. Biomass was highest at the shelf break, where it 23 approached 70 mg DW m⁻³, and was mainly composed of the large copepod *Calanus* 24 glacialis. On the shelf, abundance and biomass were low (10-20 mg DW m⁻³) and dominated 25 by small copepods and chaetognaths. Several distinct assemblages of zooplankton were 26 identified and related to the physical properties of the water masses present. A striking 27 result was the presence of both Atlantic and Pacific expatriates in offshore waters close to 28 the shelf break, but generally not on the shelf. Tracking these advected organisms could be 29 a useful tool in determining the pathways, extent and transit time of Atlantic and Pacific 30 water entering the Arctic.

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32 Keywords: Arctic Ocean, zooplankton, pelagic ecosystems, climate change

33 Introduction

34 The East-Siberian Sea (ESS) is located east of the Laptev Sea and west of the Chukchi 35 Sea, bordered by the New Siberian Islands on the west and Wrangel Island on the East (Fig. 36 1). It is the largest, broadest and shallowest of all Arctic shelf seas, widely open to the Arctic 37 Basin. Nearly 70% of the shelf of the ESS is shallower than 50 m, with most of the area 38 dominated by depths of 20-25 m (Williams and Carmack 2015). Oceanographically, it 39 interacts both with the adjacent Atlantic-influenced Laptev Sea, and the Pacific-influenced 40 Chukchi Sea and is heavily influenced by river runoff from large Siberian rivers Kolyma and 41 Indigirka (Semiletov et al. 2005). The water exchange between the ESS and neighboring 42 Laptev and Chukchi Seas is mostly determined by atmospheric circulation varying 43 significantly year to year. Eastward winds keep riverine water from the Laptev Sea close to 44 the coast and lead to the development of the Siberian Coastal Current, which carries low 45 salinity water eastward through the Long Strait into the Chukchi Sea. In contrast, prevailing westward winds cause fresh surface water to be transported off the shelf, and the direction 46 47 of the prevailing currents is reversed, resulting in advection of Pacific-origin water from the Chukchi Sea (Weingartner et al. 1999). Historically, the ESS has been the most heavily 48 49 ice-covered shelf within the Eurasian Arctic, characterized by extensive pack ice formation that reached 300-500km from the shore (Dobrovolskii and Zalogin 1982). 50

51 Despite much effort being devoted to Arctic research in recent decades, mostly it has 52 been concentrated in relatively easily accessible regions within the European and north-53 American sectors (e.g. Kassens et al. 1999: Stein et al. 2003: Flint et al. 2010: Grebmeier 54 and Maslowski 2014). The large knowledge gaps remain primarily along the Siberian shelf, 55 despite its high significance for sea ice formation and ocean circulation within the Arctic. Due to its remote location, shallow depths inaccessible to large research vessels, and 56 57 historically servere ice conditions, the ESS remains the most understudied of all Arctic 58 shelves, even compared to other Russian Arctic seas, with the few existing studies in 59 western literature limited to oceanography and biogeochemistry (i.e., Münchow et al. 60 1999; Semiletov et al. 2005; Anderson et al. 2011; Pipko et al. 2011). Limited information on the East Siberian shelf pelagic biological communities collected in the 1940's. and 61 1980's was published in Russian "grey" literature (Brodsky 1957; Pavshtiks 1994; Pinchuk 62

63 1994). It is not easily accessible even in Russian, and unavailable in English. No published
64 zooplankton studies have been conducted in this region since 1986.

65 As the sea ice extent, duration and thickness continues to decline in the Arctic, it has 66 become common in recent years for the entire East-Siberian shelf to become ice-free 67 during the ice minimum (Nghiem et al. 2006; Kwok et al. 2009). Similar to other areas of the Arctic (Buchholz et al. 2012; Ershova et al. 2015a; Vihtakari et al. 2018), this is 68 69 expected to result in significant shifts in plankton production patterns and community 70 composition. In order to be able to detect the ongoing changes in the pelagic ecosystem of 71 this region, within this study we aimed to provide baseline information on the structure of 72 the ESS zooplankton communities using net-based data collected in the ESS and adjacent 73 Arctic Ocean in September 2015. We also complement our data with other available 74 datasets collected on the ESS shelf: 2009, when the area was similarly nearly ice-free, and 75 August-September 1946-1948, when it was covered with pack ice (Pavshtiks 1994). This is 76 the first study in western literature providing description of the species composition, 77 spatial distribution, abundance, biomass estimates and community structure analysis of the 78 ESS zooplankton.

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80 Methods

81 Zooplankton collection and processing

Zooplankton samples were collected in September 2015 from the R/V "Akademik 82 Tryoshnikoy" at 16 stations in the northern ESS, on two transects extending from the shelf 83 (~35m depth) onto the basin (>2000m) (Fig. 1). Mesozooplankton was collected using a 84 85 closing Juday net with a mesh size of 180 µm and opening diameter of 37 cm. At each station, stratified samples were taken at depth intervals of $\sim 0.25, 25-65, 65-130, 130-260$. 86 87 and 260-450 m. No samples were collected deeper than 450 m due to the limitations of the 88 research vessel. The net was towed vertically with a wire speed of 0.5 m/sec, and closed at 89 each designated depth with a messenger, which was propelled down the wire as the net 90 ascended. The volume of the water sampled was calculated from the height of each tow; 91 100% filtering efficiency was assumed, as there were no observed cases of clogging of the 92 nets. Zooplankton samples were preserved using 10% formalin (4% formaldehyde) for 93 later processing in the laboratory.

94 In the laboratory, each sample was scanned under stereomicroscope for large and 95 uncommon species, which were identified to the lowest taxonomic level and measured. The 96 rest of the sample was split using a Folsom splitter until there were ~100 individuals of the 97 most common species in the terminal split. Increasingly larger splits were scanned to 98 obtain counts for rarer taxa; a total of 400-600 individuals were examined from each 99 sample. All organisms were measured using a computer measurement system (ZoopBiom software, Roff and Hopcroft 1986) and the DW of each specimen was predicted from a 100 101 length-weight regression relationship known for the same species, or a morphologically 102 similar organism (Ershova et al. 2015b). Copepods were staged and identified to species; 103 copepodite stages within some genera, which are morphologically undistinguishable (i.e. 104 *Pseudocalanus* spp.) were pooled together by stage. Meroplankton was grouped to the 105 macrotaxa or to the family level (in the case of shrimp larvae).

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107 Sea ice conditions, oceanography and Chlorophyll-a

108 Sea ice concentrations were obtained for each sampling location from the Nimbus-7 109 SMMR and DMSP SSM/ISSMIS Passive Microwave Data set, available through the NSIDC 110 archives (Cavalieri et al. 1996). In addition to sea ice concentration, the distance to the 111 nearest ice edge, defined as 15% ice concentration, was calculated for each station (with 112 positive values indicating open water stations, and negative values indicating ice-covered 113 stations). Temperature and salinity data were collected with a Seabird SBE911plus CTD 114 (SeaBirdTM Electronics Inc.) equipped with a dissolved oxygen sensor, svstem 115 transmissometer, fluorometer, and turbidity sensor with data binned into 1-m intervals 116 during post processing. The water column was divided into water masses based on the 117 definitions for the Arctic Ocean in Rudels (2008). Chlorophyll samples were collected using 118 Niskin bottles attached to the CTD casts at depths approximately corresponding to 3, 10, 119 20, 30, 40 and 50 meters. Typically, 500 ml of sample water was filtered onto GF/F glass 120 fiber filters, extracted in 90% acetone and analyzed fluorometrically. All samples were 121 processed at sea.

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123 Comparison to other datasets

124 Our data on zooplankton distribution was compared to published and unpublished data 125 from other expeditions collected during the same time period (September) in 2009 126 (Ershova et al. 2015b), as well as 1946 and 1948. The 1946 data, collected from the Soviet 127 ice-breaking vessel "Temp", is found in a brief publication about ESS zooplankton by 128 Pavshtiks (1994); the 1948 data, sampled from the ice-breaker "Severnyj Poljus" in the 129 Chukchi and East-Siberian Sea, is available from an archive compiled for the Arctic regions 130 by Markhaseva et al. (2005). To our knowledge, this joint dataset represents all publically 131 available zooplankton data for this region, with the exception of the expedition to Chaun 132 Bay in 1986 (Pinchuk, 1994). The latter was excluded due to being restricted only to the 133 inner bay and not extending onto the shelf. The listed expeditions have little spatial 134 overlap (Fig. 1), but together cover a wide area of the ESS shelf. The samples during the 135 2009 expedition were collected by a 150-µm double ring net of 60-cm mouth diameter, 136 with flowmeters attached at the mouth, towed vertically from ~ 5 m off the seafloor to the 137 surface at 0.5 m/sec. While the wider mouth of the net may have affected the sampling 138 efficiency, the similar mesh size makes the datasets partially comparable. During the two 139 historical cruises, 1946 and 1948, zooplankton were sampled with a closing 168-µm Juday 140 net (mouth diameter 37 cm), in a manner identical to ours. The species lists produced for the two historical datasets are very detailed for some groups (i.e. copepods), with 141 142 identification done to species and stage level, but very coarse for others, with just the broad 143 taxonomic categories (i.e. cnidarians, amphipods) listed. The taxonomy during all years 144 was cross-checked using the Arctic Register of Marine species (Sirenko et al. 2019), and 145 World Register of Marine Species (WORMS Editorial Board, 2019) in order to remove 146 synonyms and suspicious identifications. For comparing years, taxonomic assignments 147 within each dataset were adjusted to the highest common denominator. Abundance data 148 from 1946 and 1948 was converted to biomass by using average dry weights for each taxa 149 based on our own results.

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151 Data analysis

All analyses were carried out in R (R Core Computing Team 2017). Differences in biomass and abundance of key groups were compared between transects (2015) and years using a one-way ANOVA, with values log-transformed to meet ANOVA assumptions. Within the 155 2015 data, only the top 50 m were included in the analysis to exclude depth-based 156 differences. Zooplankton community structure was investigated using cluster analysis and 157 non-metric multidimensional scaling (nMDS) using the R package "vegan" (Oksanen 2013). 158 Species matrices containing abundance and biomass data were fourth-root transformed; 159 rare groups, which failed to contribute at least 5% to at least one stations were excluded. 160 Key copepod species (*Calanus* spp., *Metridia longa*) were separated by developmental stage 161 into "early juveniles" (CI-CIII) and "late juveniles/adults" (CIV-CVI) as separate categories. 162 Hierarchical cluster analysis was carried out using average linkage: significant clusters 163 were identified using the SIMPROF routine ($\alpha = 0.05$) (Clarke and Gorley, 2005). The 164 resulting clusters were confirmed by separation on a 2-dimensional nMDS ordination. The 165 taxa responsible for discriminating the groups were identified by correlating species abundances to the ordination via the *envfit* function, as well as by the function *IndVal* 166 167 (Dufrêne and Legendre 1997), which finds "indicator" species that are significantly 168 associated with specific groups based on the relative frequency and relative average 169 abundance within each cluster. The relationship of community structure to the 170 environment was explored by correlating environmental parameters to the nMDS 171 ordination using the *envfit* function, with significance established via a permutation test (*n* 172 = 10000). Additionally, the best set of physical parameters that describes community 173 structure was identified via the BIOENV routine (Clarke and Ainsworth 1993), which 174 selects the set of variables that produce the highest correlation between two data matrices. 175 Significance of these correlations was established using a permutation test (n = 10000) at p 176 < 0.05.

177 Zooplankton community structure was also examined within the context of the historical data, similar to the process above, with all sampling years pooled. Only 178 179 abundance data was used; questionable taxa, or taxa which were not 180 enumerated/identified during all of the expeditions were excluded from analysis. Depth 181 integrated data was used for 2015. The resulting clusters were overlain on an MDS 182 ordination and spatial map of the sampling locations. Taxa responsible for community 183 separation were similarly identified using the *envfit* function.

184

185 **Results**

186 Oceanographic conditions during September 2015

187 Sea ice conditions varied from completely open water to 45-60% ice cover at the 188 deeper stations of each transect (Fig. 2). The water column was well stratified along both 189 transects (Fig. 2) with distribution of water masses typical for the Arctic Ocean (Rudels 190 2008 water mass T-S characteristics defined therein). The Polar mixed layer (PML) 191 characterized by a low salinity (<29) and temperature ranging from -1 to 0 °C occupied the top \sim 20 m. This layer was underlain by the Arctic Halocline, with a higher salinity (>32.5) 192 193 and colder temperature generally below 0°C. Warm and salty Atlantic Water (AW) (T > 0194 °C) was found below depths >250 m at the basin stations, and as shallow as 150m at 195 stations along the slope, indicating possible upwelling of AW onto the shelf. A local oxygen 196 minimum, as well as a sharp salinity gradient was observed within the halocline, 197 particularly on Transect 1, possibly indicating two sources of halocline water (with upper 198 halocline of Pacific origin, 32.5 < S < 33.5 and lower halocline with 33.5 < S < 34.5, Shimada 199 et al. 2005; Rudels 2008). Chl-a concentrations were low throughout the entire sampled 200 area being highest on the shelf on Transect 1 and reaching up to 2 μ g L⁻¹. Within the surface 201 mixed layer they rarely exceeded 0.2-0.4 µg L⁻¹. The chlorophyll maximum was found 202 around \sim 50m in the basin and near bottom at the shallow shelf stations (Fig. 2).

203

204 Cross shelf-structure of zooplankton communities

205 A total of 70 taxonomic categories were identified within the mesozooplankton (Table 206 2), including 41 crustacean species (35 species of copepods, 5 amphipods, 1 ostracod), 14 207 cnidarian species (12 hydromedusae, 1 scyphomedusae, 1 siphonophora), 2 ctenophores, 2 208 pteropods, 2 larvaceans, 3 chaetognaths, and 7 meroplankton taxa (Table 2). However, the 209 number of species at each station and each sampled depth stratum was low, usually not 210 exceeding 20-30 taxa. Diversity increased with depth and away from the shelf margin, with 211 the highest species number observed within the deepest layers at the offshore stations (Fig. 212 3a). Most of the recorded taxa belonged to typical Arctic neritic and Arctic epi- and 213 mesopelagic species, but the species list also included Atlantic (*Calanus finmarchicus*) and 214 Pacific (Metridia pacifica, Eucalanus bungei) expatriates. Other than Eurytemora herdmani, 215 which was observed at one shallowest station, no euryhaline or brackish-water species, characteristic for the shallow river-influenced Siberian shelf seas (Kosobokova et al., 1998;
Lischka et al, 2001; Hirche et al, 2006) were observed.

218 The highest abundance of zooplankton was also found at stations along the slope and 219 was concentrated in the surface layer (Fig. 3b); the highest abundances on Transect 1 220 coincided with the highest surface temperatures (Fig. 2a). Abundance was significantly 221 higher in the surface waters (above 65m depth) of Transect 1 than on Transect 2 (ANOVA, 222 $F_{1,14} = 8.93$, p = 0.009). Differences between deeper layers (below 65m) were insignificant. 223 Small copepods *Pseudocalanus* spp. and *Oithona similis* dominated the abundance at all but 224 the farthest offshore stations. Appendicularians Fritillaria borealis and Oikopleura 225 vanhoeffeni were the second most significant contributors to abundance, particularly at the 226 shelf stations (Fig. 4).

227 Zooplankton biomass ranged from 10 to 100 mg DW m⁻³, and was highest at stations at 228 the shelf break on both transects, at depths of 150-250 m (Fig. 3c), mainly driven by the 229 large copepods Calanus glacialis, C. hyperboreus and Metridia longa. At the shallow shelf 230 stations, biomass was lower, not exceeding 10-20 mg DW m⁻³, with *Pseudocalanus* spp. 231 being the main contributors. Chaetognaths represented the next largest contributor to 232 biomass, composing 20-30% of total biomass at all stations. *Parasagitta elegans* dominated 233 on the shelf and slope and *Eukhronia hamata* largely replaced it at the deeper sampling 234 locations (Fig. 4).

235 Multivariate analysis separated zooplankton communities by depth and distance from 236 shore according to both abundance (Fig. 5) and biomass data (Online Resource 1), with 237 surface shelf communities being distinct from surface slope communities, as well as deeper 238 communities at the shelf break. Based on abundance, the community was homogenous in 239 the surface and subsurface layers (0-25 and 25-65 m) at the Transect 1 stations; on 240 Transect 2, on contrary, most stations contained distinct communities in these two layers 241 (Fig 5; Online Resource 1). Based on both the IndVal analysis (Online Resource 2) and 242 correlations to the nMDS ordination (only significant correlations shown), the separation of 243 the **surface shelf communities** was driven by the increased abundance and biomass of the 244 hydromedusae *Calycopsis birulai*, *Aeginopsis laurentii*, the copepods *Microsetella norvegica*, 245 Acartia longiremis, as well as adult females of Calanus glacialis, and the larvacean 246 *Oikopleura vanhoeffeni*. The **off-shelf surface communities**, mainly observed on the slope,

247 were distinguished by high numbers of the juvenile *Pseudocalanus* spp., juvenile *C. glacialis* 248 and meroplankton: echinoderm, bivalve and polychaete larvae. A few stations on in the 249 surface waters on Transect 2 also clustered as a separate community type, driven by the 250 high abundance of juvenile Aglantha digitale, Fritellaria borealis and Oithona similis. The 251 **sub-surface communities** along the slope were distinguished by the mesopelagic *Triconia* 252 borealis, Metridia longa (adults and older juveniles), P. minutus (adults and older juveniles), 253 Microcalanus pygmaeus, Paraeuchaeta glacialis, as well as the Pacific expatriates Eucalanus 254 *bungii* and *Metridia pacifica*. These stations also contained high biomass of the chaetognath 255 *Parasagitta elegans.* Finally, the **deep communities** in the waters at the shelf break were 256 distinguished by a number of Arctic Basin species, as well as the Atlantic expatriate *Calanus* 257 finmarchicus. Biomass data separated the 25-65 m layer as a distinct group only at the 258 slope stations of both transects (Online Resource 1).

259 The MDS ordination was largely determined by sampled depth strata, which distributed 260 stations along Axis 1. It was also strongly and significantly (p < 0.01) correlated with 261 physical factors, with salinity and bottom depth having the highest correlations to the 262 ordination (Fig. 5). Chl-*a* concentration, sea ice concentration and temperature were also 263 significantly correlated to the ordination, but explained less of the variability. There was no correlation to distance from ice edge or oxygen concentration. The BIOENV routine 264 265 predicted layer depth as the most significant factor correlated with community structure (ρ 266 = 0.64), while the inclusion of salinity and bottom depth increased Spearman's correlation 267 to $\rho = 0.7$.

268

269 Comparison with other datasets

270 Overall mean abundance and biomass values were similar between years (Table 3), 271 although some significant differences between sampling years were observed in overall 272 zooplankton abundance (ANOVA, $F_{3,44} = 5.71$, p = 0.00216) and biomass (ANOVA, $F_{3,44} =$ 273 5.94, p = 0.00345), as well as abundance of *Calanus glacialis* (ANOVA, $F_{3,44} = 9.51$, p < 274 0.0001). Abundance and biomass were significantly higher in 2009 than during the other 275 sampling years; abundance of *C. glacialis* was significantly higher in 2009 and 2015 than in 276 the two early years (Table 3, Fig. 6). The diversity in 2009 and 2015 was also much higher, 277 due to the absence of Arctic Basin species in the shallow regions explored in 1946 and

1948, as well as overall lower taxonomic resolution during the processing of 1946 and1948 samples.

280 Nevertheless, some species were observed during these earlier expeditions that were 281 not recorded (Table 2, Fig. 6). These include the brackish water species Pseudocalanus 282 major and Drepanopus bungei, as well as juvenile and adult euphausiids, which were 283 common in the eastern part of the region during the three expeditions prior to 2015. 284 Multivariate analysis (cluster analysis and MDS) grouped stations from all the four 285 expeditions into 5 groups with two outliers (Fig. 7). The most distinct group was the 286 stations west of Chaun Bay (coral color on Fig. 7), characterized by the dominance of 287 brackish *P. major* and *D. bungei*, which were observed nowhere else, as well as extremely 288 low abundance and biomass of all other species. The stations near Chaun Bay formed a 289 distinct group (green color on Fig. 7), separated by the presence of brackish *P. major* and 290 Limnocalanus macrurus. The remainder of the shelf stations belonged to two groups: one 291 common to all expeditions (vellow color on Fig. 7), with the separation driven by the 292 abundance of hydrozoan jellyfish, harpacticoid copepods, appendicularians, and 293 meroplankton. The other group (pink color on Fig. 7) was unique to 2009 and was 294 characterized by the higher abundances of the hydrozoan jellyfish *Aeginopsis laurentii* and 295 the chaetognath *Parasagitta elegans*. All stations deeper than 50 m (all from 2015) grouped 296 together, characterized by typical Arctic Basin communities. Pacific expatriates, with the 297 exception of euphausiids, were generally only observed during the 2015 expedition at 298 stations deeper than the shelf break.

299

300 Discussion

301 The patterns of zooplankton abundance and diversity across the shelf as well as 302 through the vertical column in the northern region of the East Siberian Sea during our 303 study in 2015 generally followed those observed earlier in other parts of the Arctic, with 304 highest zooplankton abundance and biomass found in the surface layers, and increasing 305 diversity with depth (Fig. 3) (Kosobokova et al. 1998; Ashjian et al. 2003; Hirche et al. 306 2006; Kosobokova and Hirche 2009). A marked hotspot of both abundance and biomass 307 was observed close to the shelf break, with much lower values observed both farther on-308 and off- the shelf. The biomass peak at the shelf break was mainly driven by *Calanus* 309 glacialis, which similarly concentrated at the outer shelf and slope in other parts of the 310 Arctic, substantially contributing to the "carbon belt" of elevated zooplankton biomass 311 around the entire Eurasian slope (Kosobokova and Hirche 2009; Kosobokova 2012). The 312 vertical structure of zooplankton communities in the northern ESS was fairly typical for 313 other regions of the Arctic, with distinct communities in the surface, sub-surface and 314 mesopelagic layers (Kosobokova and Hopcroft 2010; Kosobokova et al. 2011). The upper 315 limit of the range and mean biomass we observed in the northern ESS (10-100mg DW, 316 mean \sim 30 mg DW) was higher than previously reported for the ESS, similar to the Laptev 317 and Kara Seas (Kosobokova et al. 1998; Fahl et al. 1999; Abramova and Tuschling 2005; 318 Arashkevich et al. 2018). They were, however, lower than in the adjacent Chukchi Sea 319 (Ouestel et al. 2013; Ershova et al. 2015b), which experiences a much longer productive 320 period and a high inflow of nutrients and allochtonous biomass from the North Pacific.

321 Semiletov et al. (2005) suggested that oceanographically the East Siberian Sea can 322 be separated into two distinct regions: the river-influenced western region and the Pacific-323 influenced eastern region, separated by a frontal zone which position fluctuates year-to-324 year between 160° and 170° E. While our combined dataset is mostly concentrated west of 325 this front, our data suggests distinct regional differences in zooplankton community structure between these two regions. The inner western stations (all sampled in 1948), 326 327 heavily influenced by river outflow, are characterized by overall low abundance, diversity, 328 and dominance of brackish species, which were not observed elsewhere. The outer shelf 329 stations (sampled in 2015), although also located far enough to the west, did not capture 330 any of this brackish signature, which is likely a consequence of the river-influenced domain 331 being restricted only to the inner shelf. Alternatively, it is possible that during 2015 the 332 frontal zone separating the two domains was located farther to the west. It is noteworthy 333 that all remaining shelf stations were similar in zooplankton community structure, both in 334 present times and 75 years ago, and contained typical Arctic neritic fauna, which is shared 335 with adjacent Siberian Arctic seas (Kosobokova et al. 1998; Lischka et al. 2001; Hirche et al. 336 2006; Flint et al. 2010). Our data suggest that Long Strait and the area around Wrangel 337 Island may represent yet another community type, possibly influenced by increased flow 338 from the Chukchi Sea (Coachman and Rankin 1968; Pisareva et al. 2015). And, finally, at the 339 shelf break, the zooplankton communities transitioned to typical Arctic Basin type

340 containing a much higher diversity compared to the shallow ESS shelf, and characteristic
341 Arctic Basin type of vertical structure (Kosobokova and Hopcroft 2010; Kosobokova et al.
342 2011).

343 As an interior arctic shelf sea (Williams and Carmack 2015), the ESS has limited 344 exchange with other oceans as well as adjacent regions of the Arctic when compared to 345 regions of inflow/outflow (the Barents Sea, Chukchi Sea), but advection still plays an 346 important role in structuring its pelagic communities. It is noteworthy that the ESS is one of 347 the few locations in the Arctic influenced by inflow both from the Atlantic and Pacific 348 oceans. One striking result of our study in this context is the record of both Atlantic and 349 Pacific expatriates (the copepods *Calanus finmarchicus, Eucalanus bungii, Metridia pacifica*) in our study area in 2015, albeit in fairly low numbers, and the Pacific expatrites 350 351 euphausiids Thysanoessa spp. in 1946, 1948 and 2009 (although we cannot exclude that 352 they represent a local reproducing population). The advected copepods were generally 353 found in the water layers below 50m, and all observed specimens were adults/sub-adults. 354 indicating that they probably have been in transit for an extended period. A period of 5-6 355 years is required for inflowing Atlantic water to reach the East Siberian Sea, (E. Carmack, 356 pers. comm.). Although lifespans of pelagic copepods have never been measured in natural 357 conditions, it is generally believed that few are capable of surviving for 5+ years (Hirche 358 2013). Tracking the advected organisms could be a useful tool in determining the 359 pathways, extent and transit time of Atlantic and Pacific water entering the Arctic.

360 While both shelf and off-shelf regions of the ESS may be influenced by Pacific water 361 (Semiletov et al. 2005), the taxonomic composition of Pacific expatriates is very different in 362 the two regions. On the inner shelf, euphausiids Thysanoessa spp., which are also not 363 presumed to reproduce in the Arctic and are not found farther to the west, were observed 364 during nearly every sampling year (1947, 1948, 2009), but Pacific copepods were absent or 365 overlooked. At the off-shelf stations, the reverse was true, with the Pacific copepods 366 Eucalanus bungii and Metridia pacifica being prominent members of the communities 367 (2015). This is suggestive of two distinct sources of Pacific water present in the region: the 368 more "short-term" water entering the ESS through Long Strait from the Chukchi Sea 369 (Semiletov et al. 2005), containing entrapped euphausiids, and the "older" Pacific water, 370 which has been in transit for at least one season (Bluhm et al., 2015), entering from the

371 Canadian Basin away from the shelf. It is unclear why we also do not observe Pacific 372 copepods in the Long Strait region. As they are oceanic species, originating on the Bering 373 Sea slope by the Gulf of Anadyr, perhaps they cannot survive the low salinity conditions of 374 this region, which is influenced by the fresh waters of the Siberian Coastal Current 375 (Weingartner et al. 1999). It is also quite possible that part of the *C. glacialis* that we 376 observe in the area belongs to the north Bering Sea population that is advected through the 377 Bering Strait (Nelson et al. 2009). The latter point is impossible to clarify without genetic 378 studies.

379 Historically, the ESS has been one of the most heavily ice covered regions of the 380 Arctic (Nghiem et al. 2006), but has in recent decades transitioned into part of the seasonal 381 ice zone, with extended areas of open water during the summer months (Maslanik et al. 382 2007). The decline in sea ice extent and duration, as well as increasing Atlantification 383 (Polyakov et al. 2017) of the Eastern Arctic are shifting the conditions in to resemble those 384 in the western Eurasian Basin, and such a shift is expected to be reflected in the structure 385 and/or productivity of biological communities. Although the data we collected is 386 insufficient to document change, the significantly higher abundance of zooplankton, 387 particularly *Calanus glacialis*, that we observed during recent years (2009, 2015), may 388 suggest that this Atlantification is already having an effect on the region. Of course, such 389 conclusions must be reached with caution, given the different spatial coverage and 390 sampling methods during the different expeditions. However, these changes would not be 391 entirely surprising, given the trends of increasing of zooplankton advection into the Arctic 392 (Johannesen et al. 2012; Gluchowska et al. 2017), including the adjacent Chukchi Sea 393 (Ershova et al. 2015a). We expect our study to provide the important information on the 394 present state of the plankton communities of this understudied arctic sea and will serve as 395 a valuable baseline to document further change within this region.

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- 549 Figures
- 550
- **Figure 1** Sampling locations in the East Siberian Sea during September 2015 and sampling
- locations of other expeditions included in the dataset (1946, 1948 and 2009)
- **Figure 2** Sea ice concentration, salinity, temperature, oxygen and Chl-*a* distributions along
- Transect 1 (a) and Transect 2 (b). Dashed lines delineate presumed water masses

Figure 3 Depth-stratified distribution of abundance (ind m⁻³), biomass (mg DW m⁻³), and
 diversity (number of species) along Transects 1 and 2 in the East-Siberian Sea

557 **Figure 4** Relative contribution of most common species to (a) biomass and (b) abundance

in September 2015 across Transect 1 in the East Siberian Sea (Transect 2 followed similar

trends where the depth ranges overlapped). The diameter of the circles at the top of the

- 560 columns indicate total abundance/biomass.
- 561 **Figure 5** (a) results of non-metric multidimensional scaling and cluster analysis of fourth

root-transformed zooplankton abundance in the East-Siberian Sea during September 2015

and (b) locations of clusters along the sampled transects. Each symbol represents one

sample; symbol types represent sampled depth strata; colors indicate significant clusters.

565 Vectors show correlations of species abundances and physical parameters to the

566 ordination (with length reflecting the correlation coefficient *r*)

Figure 6 Distribution of (a) overall zooplankton abundance, (b) overall zooplankton
biomass, and (c-e) select zooplankton species in the East-Siberian Sea during 1946, 1948,
2009 and 2015

570 **Figure 7** (a) results of non-metric multidimensional scaling and cluster analysis of fourth

571 root-transformed abundance data for all years pooled, and (b) spatial distribution of

572 clusters. Each symbol represents one sample; symbol types represent sampling year; colors

573 indicate significant clusters. Vectors show correlations of species abundance to the

574 ordination (with length reflecting the correlation coefficient *r*)

ESM1 (a) results of non-metric multidimensional scaling and cluster analysis of fourth logtransformed zooplankton biomass in the East-Siberian Sea during September 2015 and (b)
locations of clusters along the sampled transects. Each symbol represents one sample;
symbol types represent sampled depth strata; colors indicate significant clusters. Vectors
show correlations of species abundances and physical parameters to the ordination (with
length reflecting the correlation coefficient *r*)

581

Compliance with Ethical Standards

- 584 The authors declare that the research was conducted in the absence of any commercial or
- *financial relationships that could be construed as a potential conflict of interest.*





Figurea³ Species diversity

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(b) Abundance (ind m⁻³)



(c) Biomass (mg DW m⁻³)







Table 1. Average abundance (ind m⁻³) and biomass (mg DW m⁻³) of all zooplankton species observed during 1946, 1948, 2009, and 2015 in the ESS. "+" – indicates values <0.1; "obs." – species were observed but not enumerated; "-" – no data.

	1946 (data from Markhaseva et		1948 (Pavshtiks,		2009 (Ershova et		2015 (This			
Species	al. 2005)		1994)		al., 2015a)		study)			
	Abund.	Biom.*	Abund.	Biom.*	Abund.	Biom.	Abund.	Biom.		
Copepoda	Copepoda									
Acartia longiremis	75.9	0.4	19.8	0.1	36.1	0.1	4.8	+		
Aetideopsis rostrata	-	-	-	-	-	-	0.3	+		
Bradyidius similis	-	-	-	-	-	-	4.0	0.1		
Aetideidae	-	-	-	-	-	-	0.2	+		
Calanus finmarchicus	-	-	-	-	-	-	0.6	0.1		
Calanus glacialis	17.9	1.5	106.9	16.0	147.3	10.6	75.1	8.8		
Calanus hyperboreus	-	-	-	-	0.1	0.3	7.0	3.7		
Chiridius obtusifrons	-	-	-	-	-	-	0.8	0.2		
Disco triangularis	-	-	-	-	-	-	0.2	0.3		
Drepanopus bungei	-	-	871.3	2.2	-	-	-	-		
Eucalanus bungii	0.4	0.1	-	-	-	-	0.3	0.1		
Eurytemora herdmani	-	-	-	-	-	-	1.1	+		
Gaetanus brevispinus	-	-	-	-	-	-	+	+		
Gaetanus tenuispinus	-	-	-	-	1.0	0.1	1.2	0.2		
Heterorhabdus norvegicus	-	-	-	-	-	-	1.0	0.1		
Jaschnovia brevis	-	-	-	-	-	-	10.1	0.2		
Jaschnovia tolli	24.7	1.9	31.9	1.0	-	-	0.6	+		
Jaschnovia sp.	-	-	-	-	-	-	1.3	+		
Limnocalanus macrurus	3.7	+	-	-	-	-	-	-		
Metridia longa	-	-	-	-	6.8	0.9	30.3	1.9		
Metridia pacifica	-	-	-	-	-	-	1.2	0.2		
Microcalanus pygmaeus	67.1	0.1	19.8	+	1.3	+	34.2	0.1		
Microsetella norvegica	-	-	-	-	123.1	0.9	4.1	+		
Paraeuchaeta glacialis	-	-	-	-	-	-	0.9	0.6		
Paraeuchaeta barbata	-	-	-	-	-	-	+	+		
Paraeuchaeta polaris	-	-	-	-	-	-	+	+		
Paraheterorhabdus compactus	-	-	-	-	-	-	+	+		
Pseudocalanus newmani	-	-	-	-	-	-	4.0	+		
Pseudocalanus acuspes	-	-	-	-	26.8	0.3	38.6	0.3		
Pseudocalanus major	13.8	0.1	119.4	0.3	-	-	-	-		
Pseudocalanus minutus	-	-	-	-	3.6	0.1	30.8	0.3		
Pseudocalanus sp.	913.7	2.3	285.5	0.7	4331.8	10.3	1140.8	3.4		
Scaphocalanus brevicornis	-	-	-	-	-	-	+	+		
Scaphocalanus magnus	-	-	-	-	-	-	0.4	0.2		
Scolethricella minor	-	-	-	-	-	-	1.1	+		
Spinocalanus antarcticus	-	-	-	-	-	-	+	+		
Spinocalanus longicornis	-	-	-	-		-	5.6	+		
Temorites brevis	-	-	-	-	-	-	+	+		
Other calanoida	-	-	-	-	-	-	1.2	0.1		

Calanoida nauplii	1170.4	0.1	418.8	0.1	989.7	0.6	189.8	0.3
Harpacticoida	209.5	0.7	17.3	0.1	13.3	0.1	0.4	0.1
Oithona similis	411.9	0.6	225.7	0.3	1836.7	2.7	653.0	0.7
Triconia borealis	60.8	0.1	-	-	0.2	+	37.7	0.1
Cyclopoida	-	-	-	-	-	-	0.5	+
Cnidaria		•	•					
Aeginopsis laurentii	2.5	0.1	2.6	0.1	8.2	0.2	3.9	0.1
Aglantha digitale	-	-	-	-	3.7	0.3	6.0	0.9
Calycopsis birulai	-	-	-	-	1.2	3.0	0.4	0.1
Cyanea capillata	-	-	-	-	-	-	0.2	+
Dimophyes arctica	-	-	-	-	-	-	0.9	1.2
Euphysa flammea	0.4	0.1	-	-	-	-	-	-
Halitholis cirratus	0.5	0.3	-	-	0.1	0.5	0.1	0.4
Melicertum octocostratum	-	-	-	-	+	0.1	-	-
Obelia longissima	-	-	-	-	0.6	+	-	-
Paragotoea elegans	-	-	-	-	-	-	+	+
Plotocnide borealis		-	-	-	2.7	0.2	0.1	+
Rhabdon reesi	-	-	-	-	-	-	+	+
Sarsia tubulosa	0.3	0.3	-	-	0.8	2.0	-	-
Sminthea arctica	-	-	-	-	-	-	0.1	+
Other cnidaria	0.3	+	-	-	-	-	-	-
Amphipoda								
Apherusa glacialis	-	-	-	-	0.5	0.9	0.2	0.3
<i>Hyperia</i> sp.	0.6	0.6	-	-	-	-	-	-
Hyperiidae	1.0	0.2	-	-	0.6	0.5	0.3	+
Hyperoche sp.	0.2	0.3	-	-	0.8	0.7	+	+
Themisto abyssorum	-	-	-	-	-	-	0.5	0.9
Themisto libellula	-	-	1.2	1.2	0.1	1.1	0.2	3.1
Scina borealis	-	-	-	-	-	-	+	+
Ctenophora								
Beroe cucumis	-	-	3.5	0.4	obs.	obs.	obs.	obs.
Mertensia ovum	-	-	-	-	0.6	1.0	0.2	0.2
Ostracoda						-		-
Boroecia maxima	-	-	-	-	-	-	2.6	+
Pteropoda								
Clione limacina	-	-	-	-	-	-	0.3	0.1
Limacina helicina	-	-	-	-	12.9	0.1	10.0	0.1
Chaetognatha								
Eukrohnia hamata	-	-	-	-	-	-	4.1	0.8
Parasagitta elegans	9.9	2.7	28.4	4.3	132.3	19.9	63.2	6.3
Pseudosagitta maxima	-	-	-	-	-	-	0.1	0.2
Larvacea		I		I				I
Fritillaria borealis	232.5	0.1	-	-	165.6	0.1	53.5	+
Oikopleura vanhoeffeni	277.8	14.6	118.8	11.9	73.2	0.2	75.9	0.5
Mysidae		I		I				I
Mysis oculata	0.7	0.7	-	-	+	0.6	-	-
Euphausiidae								

Thysanoessa sp.	0.6	2.8	45.4	22.7	1.7	9.3	-	-
Meroplankton								
Ceriantharia	-	-	-	-	2.3	+	1.4	+
Bivalvia	-	-	4.1	+	10.5	+	4.3	+
Cirripedia	-	-	9.9	+	18.4	0.3	5.7	0.1
Decapoda	0.3	+	-	-	0.1	0.4	0.1	+
Fish larvae	-	-	-	-	0.1	3.3	-	-
Nemertina	-	-	-	-	-	-	0.3	+
Echinodermata	301.9	0.1	17.7	+	114.8	0.2	7.6	+
Polychaeta	168.3	0.8	20.0	0.1	16.7	0.2	6.4	0.1

Table 2 Pearson's correlations between log-transformed abundance data and environmental variables in the East Siberian Sea in 2015 (as presented by the BIOENV routine). S - salinity

Variables	<u># of variables</u>	ρ	
Sampled depth layer		1	0.63
S			0.47
Bottom depth			0.42
Sampled depth layer + Bottom depth		2	0.66
Sampled depth layer + S			0.62
S + Bottom depth			0.57
Sampled depth layer + Bottom depth + S		3	0.69

Table 3 Total zooplankton abundance, biomass, diversity and mean abundance of *Calanus* glacialis in the East Siberian Sea in 1946, 1948, 2009 and 2015 (mean \pm sd); significant interactions via the Tukey HSD test: significant interactions between years at p < 0.05 listed in **bold**; non-bold interactions indicate significance at p < 0.1. Basin stations with depths >200m were excluded from analysis

	1946 (<i>n</i> = 17)	1948 (<i>n</i> = 9)	2009 (n = 9)	2015 (<i>n</i> = 13)	Signif. interactions
Total	3189 ± 2013	1939 ± 2116	7091 ± 6940	2908 ± 1950	2009 > 1946, 1948
abundance					
(ind m^{-3})					
Total	25 ± 24	28 ± 44	54 ± 23	35 ± 16	2009 > 1946, 1948 ; 2015 >
biomass (mg					1948
DW m ⁻³)					
Calanus	14 ± 21	83 ± 231	147 ± 181	163 ± 172	2009, 2015 > 1946, 1948
glacialis					
(ind m^{-3})					
Number of	28	20	41	70	
taxa					
recorded					

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