1 Using biological traits and environmental variables to characterize two Arctic

# 2 epibenthic invertebrate communities in and adjacent to Barrow Canyon

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- 11 ABSTRACT

12 The Arctic's Barrow Canyon, located in the northeastern Chukchi and western Beaufort seas, supports a 13 rich and diverse benthic ecosystem and is often termed an ecological "hotspot" of productivity. Within

- 14 and adjacent to Barrow Canyon, the epibenthic invertebrate communities vary, with biomass and
- 15 taxonomic distributions related to habitat variation. Here we asked if the patterns observed are due to
- 16 Barrow Canyon's variation in near-seafloor physical hydrography, and whether differences in taxonomic
- 17 distribution also reflect differences in functional properties of the epibenthic invertebrate community.
- 18 Data were collected using a standardized 83-112 bottom trawl during two surveys in and adjacent to
- 19 Barrow Canyon: the northeast Chukchi Sea survey in 2013 and the western Beaufort Sea survey in 2008.
- 20 A portion of the Beaufort Sea survey also used a liner to retain smaller organisms. A suite of nine
- 21 environmental variables were examined, that included depth, bottom water temperature, bottom
- 22 hardness as measured by acoustics, and circulation model hindcast current speed. They explained 18-
- 23 47% of observed variance for each of the three data sets (Chukchi Sea, Beaufort Sea lined net (LN),
- 24 Beaufort Sea unlined net (UN)). In the Chukchi Sea, bottom hardness and depth were significant
- 25 variables. In the Beaufort Sea LN hauls, depth, bottom temperature, and the mean current speed on the
- 26 day of sampling were significant variables and in the Beaufort Sea UN hauls, depth was the only

27	significant variable. Of the 150+ collected taxa from each survey, ~20 made up 90% of the total biomass
28	in the Beaufort and Chukchi Seas, and six of the 20 taxa were common to both study areas. We used
29	biological traits analysis (BTA) of body morphology, trophic, and reproductive traits to further
30	characterize the epibenthos at the head of Barrow Canyon in the Chukchi Sea and into Barrow Canyon in
31	the Beaufort Sea. Although the Chukchi and Beaufort seas differed taxonomically in abundance and
32	distribution, they were functionally similar based on the biological traits we examined. A traits analysis
33	can advance knowledge of a community of organisms; however, it is most informative if used as a
34	complement to a taxonomic composition analysis of abundance and distribution.
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39 Keywords: Chukchi Sea; Beaufort Sea; Barrow Canyon; epibenthic invertebrates; biological traits

#### 40 1. Introduction

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42 Barrow Canyon (Fig. 1) is an important marine geologic feature situated in the northeast 43 Chukchi Sea and is frequently referred to as the gateway to the Pacific Arctic Basin. The head of Barrow 44 Canyon starts approximately 150 km southwest of Point Barrow and extends into the Beaufort Sea with 45 depths up to 300 m. Several water masses transiting from the Bering and Chukchi continental shelves 46 flow northward into the Arctic Basin through Barrow Canyon (Weingartner et al., 2005a; Gong and 47 Pickart, 2015). Upwelling in Barrow Canyon causes locally elevated primary production (Hill and Cota, 48 2005) which supports large numbers of both pelagic- and benthic- feeding seabirds (Kuletz et al., 2015; 49 Lovvorn et al., 2015) and marine mammals (Moore et al., 2010) during the summer months. Arctic cod 50 (Boreogadus saida) are also common, both within the vicinity of Barrow Canyon and in the canyon itself, 51 where abundance is highest in colder and more saline near-bottom waters (Logerwell et al., this issue). 52 In the areas adjacent to Barrow Canyon, the Chukchi and Beaufort Sea shelves, epibenthic communities 53 have been well described (Feder et al., 1994; Bluhm et al., 2009; Blanchard et al., 2013a, b; Ravelo et al., 54 2014; Grebmeier et al., 2015a, b; Ravelo et al., 2015), including a macrofaunal benthic "hotspot" that 55 has persisted for decades, just northwest of the canyon on the Chukchi shelf (Grebmeier et al., 2015). 56 However, despite the oceanographic and biological importance of Barrow Canyon, few studies have 57 explicitly examined the benthic invertebrate fauna within the canyon. In this study, we examined two 58 communities that together encompass most of Barrow Canyon; the Chukchi study area is located at the 59 head of Barrow Canyon, extends onto the Chukchi shelf, and is relatively shallow. The Beaufort study 60 area is located north of the Chukchi study area, in the deeper portion of Barrow Canyon and extends onto the Beaufort Shelf. 61

In both the northeast Chukchi and western Beaufort seas, benthic invertebrate communities are
 characterized by moderate species diversity, compared to other Arctic shelf seas (Piepenburg et al.,

64 2011), and high biomass (~5000 kg/km<sup>2</sup>) when compared to the corresponding benthic fish communities 65 (Rand and Logerwell, 2011). Within each community, patchiness of both species distribution and biomass are common (Ravelo et al., 2014, 2015). Several features of the local habitat influence the 66 67 structure of Arctic benthic communities, such as currents (Grebmeier et al., 2006), nutrient fluxes and 68 food availability (Grebmeier et al., 1989; Cusson and Bourget, 2005; Dunton et al., 2005), and geological 69 characteristics (e.g., sediment, geological structure; Feder et al., 1994; Cusson and Bourget, 2005). The 70 study region's currents within and adjacent to Barrow Canyon consists of swift flows associated with the 71 Alaskan Coastal Current (ACC), the Beaufort Gyre, the Beaufort shelfbreak jet, and wind-driven events 72 (including upwelling), that cause meanders and eddies to disrupt the background circulation field 73 (Pickart et al., 2013). The ACC typically flows from the Chukchi Shelf along the axis of Barrow Canyon toward the continental slope where the Chukchi and Beaufort seas meet (Fig. 2). The Barrow Canyon 74 75 mean flow opposes the prevailing winds that blow from the northeast but the winds are often strong 76 enough to reverse the circulation here and that of the shelfbreak jet, as well as cause upwelling of 77 subsurface waters within Barrow Canyon and along the Beaufort Slope (Mountain et al., 1976; 78 Weingartner et al., 1998; Pickart et al., 2013; Danielson et al., 2016). Elevated levels of turbulent mixing 79 within the canyon is likely biologically important (Shroyer, 2012). The waters leaving Barrow Canyon, in general, tend to follow topographic isobaths eastward and transition into the Beaufort shelfbreak jet 80 81 (Nikolopoulos et al., 2009; von Appen and Pickart, 2012; Gong and Pickart, 2015). The complex 82 bathymetry of the canyon and slope intersection routinely causes eddies to form near the canyon mouth (Pickart et al., 2005; Watanabi and Hasumi, 2009) which propagate into the deep adjoining basin 83 84 or get caught in the westward-flowing Beaufort Gyre (Watanabe et al., 2011). The currents here evolve 85 over time as a function of the wind and the various water densities, together determining the structure 86 of the flows in and adjacent to Barrow Canyon. This energetic flow environment - and associated 87 fronts, advection, convergence, turbulent mixing, and particulate export - sets the stage for areas of

locally elevated biological activity and benthic production in and near Barrow Canyon. How the currents
and thermohaline properties influence the structuring of the local benthic invertebrate community is
complex (Day et al., 2013) and not well understood in detail.

91 While the Chukchi and Beaufort Sea shelves benthic and pelagic communities are reasonably 92 well characterized with respect to species diversity (e.g. richness and evenness), biomass, relative 93 abundance, and trophic structure (Feder et al., 1994; Bluhm et al., 2009; Blanchard et al., 2013a, b; 94 McTigue et al., 2014; Ravelo et al., 2014; Divine et al., 2015; Grebmeier et al., 2015a, b; Ravelo et al., 95 2015), other functions performed by dominant benthic organisms in this region are less well 96 characterized. At the most basic level, functional ecology is defined as the study of ecological processes, 97 patterns, and underlying mechanisms within an ecological community. Further, a functional or 98 biological trait (herein the same) infers the underlying processes of that trait, such as feeding or growth. 99 The degree of overlap between species diversity and functional diversity is different among systems. A 100 system that may be rich in species diversity could have little functional diversity (i.e. most species feed 101 similarly) or very high functional diversity (e.g. several different feeding strategies; Hewitt et al., 2008). 102 Thus, functional community structure can inform us about the energy flow and resource partitioning in a 103 system and provide insight into a system's resilience to change. Incorporating ecological function into a 104 diversity analysis is especially important in the benthic marine system because of the strong relationship 105 between habitat variability and taxonomic diversity and distribution (Hewitt et al., 2008).

In addition to characterizing the epibenthic community structure in these communities, we
applied a biological traits analysis (BTA) to better understand the influence of bottom hardness and
hydrography on the ecological functions of epibenthos in and adjacent to Barrow Canyon. This approach
has been used most recently in other Arctic communities (Oug et al., 2012; Krumhansl et al., 2016),
including the Barents Sea (Cochrane, et al., 2012; Wlodarska-Kowalczuk et al., 2012), and the Canadian

111 Arcitc (Krumhansl et al., 2016). A BTA approach (Bremner et al., 2003, 2006) is a tool used to examine a 112 suite of biological traits simultaneously across the dominant taxa in a study system to aid in identifying 113 those habitat characteristics that may influence the selection of traits (Bremner et al., 2003). Our 114 approach in this study followed the general framework of several earlier studies that used biological 115 traits to characterize invertebrates in freshwater (Jackson, 1993; Usseglio-Polatera et al., 2000; Haybach 116 et al., 2004; Bonada et al., 2007; Conti et al., 2014), lagoons (Sigala et al., 2012), estuaries (Alves et al., 117 2014), and marine ecosystems (Bremner et al., 2006; Hewitt et al., 2008; Pacheco et al., 2010; Paganelli 118 et al., 2012; Krumhansl et al., 2016). To our knowledge, a BTA approach has not been used to examine 119 the epibnethic community structure in both the Chukchi Sea or US Beaufort Sea.

120 In this study, we assessed both the taxonomy and biological traits of two adjacent epibenthic 121 communities in a hydrographically complex region of the Pacific Arctic shelf; the Chukchi community 122 that sits at the head of Barrow Canyon and the Beaufort community that sits in Barrow Canyon and onto 123 the Beaufort shelf. Because the Chukchi and Beaufort portions of Barrow Canyon share similar water 124 masses, we expected that the epibenthic invertebrate communities shared both taxonomic and 125 biological trait similarities. Specifically, we 1) used environmental variables to explain the spatial 126 variability in benthic invertebrate taxa that comprise the top 90% of the biomass, and further we 2) 127 described each community using a suite of biological traits to identify patterns in their distribution and 128 how these relate to the regional oceanographic characteristics. Finally, we discuss how a biological 129 traits analysis could be potentially useful in monitoring the effects of change within the Arctic epibenthic 130 invertebrate communities.

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132 2. Methods

#### 134 2.1. Data collection in the Chukchi Sea

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136 Chukchi Sea samples were taken during the 2013 SHELFZ (Shelf Habitat and Ecology of Fish and 137 Zooplankton) survey in the northeast Chukchi from Wainwright (160° W) to Point Barrow (155° W) and 138 encompassed the southern portion of Barrow Canyon from 17 August to 5 September 2013 (Fig. 1). 139 Epibenthic invertebrates were collected from 29 bottom hauls. The standardized 83-112 eastern otter 140 trawl net (Stauffer, 2004) was used for all sampling and has a 25.3-m headrope and a 34.1-m footrope 141 with a 10.16 cm mesh in the wings and body, and an 8.89 cm mesh in the intermediate and codend. The 142 net towed at constant speed (3 knots) on a relatively straight tow path for 15 min, maintaining constant 143 bottom contact as measured by the HOBO Pendant G Acceleration Data Logger. The measured net 144 opening was approximately 2 m from the head rope to the foot rope and 18-20 m in width, from wing to 145 wing. The width and spread of the net were measured using wing and head rope sensors by Marport 146 Deep Sea Technologies (Milford, NH) to insure the net maintained consistent specifications, and bottom contact measured distance fished. All hauls were conducted during daylight hours. The catch was 147 148 brought onboard the vessel, sorted to the lowest taxonomic level, counted and weighed in its entirety 149 on a motion compensated Marel scale. The invertebrate portion of the catch was quantitatively 150 subsampled for species composition. The invertebrate subsample was identified to the lowest possible 151 taxonomic level and within the lowest taxonomic level, counted and weighed to obtain a total for each 152 taxonomic group in the subsample. Total invertebrate catch compositions, counts and weights were 153 extrapolated by weight from the subsample.

In addition to processing the bottom haul catch, water column hydrographic profiles were
 collected. Conductivity, temperature, and depth (CTD) data were collected using a Seabird SBE-19
 equipped with a fluorometer and photosynthetically available radiation sensor. Data were processed

157 using a combination of the manufacturer's software (Seabird Electronics, 2012) and custom Matlab 158 software (version 14a, 2014, Mathworks) designed for visual inspection and despiking of the profile 159 data. Raw data were binned to 1 decibar pressure levels (approximately 1 m depth intervals). The 160 measured environmental variables used in the analysis of taxonomic distribution in the Chukchi Sea 161 survey were bottom temperature (°C), bottom depth (m), bottom salinity, and bottom hardness, as 162 measured by acoustics. Acoustic measurements for bottom hardness were collected using a Simrad 163 ES60 echosounder operating at 38 kHz (12° beam angle between half power points). The echosounder 164 was calibrated prior to the survey. Water column and seabed backscatter (i.e. reflected echoes) were 165 recorded continuously along survey transects, sampling at a rate of 1 Hz. All transects were divided into 166 100 m horizontal bins. Bottom types were quantified using substrate hardness metrics (unitless) on a 167 scale of 1 (least) to 10 (most) by integrating the first and second bottom reflections in Echoview 168 software (v6.1). The entire second bottom echo was integrated for the bottom hardness metric based 169 on the acoustic impedance mismatch between the seabed and the water column (Chivers et al., 1990). 170 To estimate bottom hardness at the Chukchi Sea bottom trawl stations, approximately 20 bottom 171 hardness data points, as measured with acoustics, were selected using ESRI ArcGIS version 10.3 at or 172 within the vicinity of the bottom trawl location. These selected points were then averaged within 173 ArcGIS to give a single estimate of bottom hardness for that station. The number of data points selected 174 (~20) was based on the average distanced fished by the bottom trawl.

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176 2.2. Data collection in the Beaufort Sea

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Epibenthic invertebrates were surveyed in the western Beaufort Sea in 2008. The survey extended from Point Barrow (155°W) east to 152°W (Fig. 1). The survey also used an 83-112 Eastern otter trawl; however, in a portion of the hauls a small mesh liner was added to the codend (mesh liner

181 was 3.8 cm). The portion of the survey using a lined net is termed Beaufort LN and the portion of the 182 survey using an unlined net is termed Beaufort UN. Because the net and possibly the catchability of the net were altered with the mesh liner, the data were analyzed separately and the two gear types were 183 184 considered two surveys; Beaufort Sea LN and Beaufort Sea UN hereafter. Full details of the survey and 185 the effects of the different gear types can be referenced in Rand and Logerwell (2011). The invertebrate 186 catch was quantitatively subsampled, counted and weighed using the same procedure outlined in the 187 Chukchi Sea survey. All invertebrates were identified to the lowest possible taxonomic level and 188 counted and weighed at that level to obtain a total for each taxon in the subsample. For both regions, 189 field identifications were confirmed in the lab from voucher material with the assistance of the 190 taxonomic experts listed in the acknowledgments and taxonomic names were standardized to the World 191 Register of Marine Species. 192 The water column profile characteristics were measured using the Seabird 19-Plus at or near the

**193** bottom haul stations, and processed following the procedures described for the Chukchi Sea survey.

The measured environmental variables used in both the Beaufort Sea LN and Beaufort Sea UN
hauls were the same as those used in the Chukchi Sea survey with the exception of bottom hardness.
Although an acoustic-trawl survey took place in the Beaufort Sea in 2008, the second bottom echo data
were not collected, precluding bottom hardness as an environmental variable in the Beaufort Sea
analysis.

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200 2.3. Epibenthic invertebrate density

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The catch-per-unit effort (CPUE) in kg wet weight km<sup>-2</sup> was estimated for both the Chukchi and Beaufort Sea surveys for all taxa. To calculate CPUE, the net width and distance fished was used to estimate the area swept by the net for each haul and estimated catch weight (kg) was divided by area

swept (km<sup>2</sup>). This gives an estimate of biomass density for each species or taxa in each haul. To
estimate the average CPUE for each taxa in each survey, zero catches were also included at each station
for taxa that were not present in the haul. All analyses were done on individual taxa, by survey, (Table
1) and all three surveys were treated independently. To reduce skewness that is common with CPUE
data, CPUE estimates were cube-root transformed preceding analyses.

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211 2.4. Biological Traits

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213 To focus on those invertebrates and the biological traits that dominated each of the three data 214 sets, only the taxa in the top 90% by total biomass estimated from CPUE were used in all analyses. This 215 reduced the number of taxa whose biological traits we examined for each system from >100 to less than 216 20 (Table 1). The taxa from the top 90% by total biomass were selected for each survey, independent of 217 the other surveys. Any invertebrates in the trawls that were generally considered infaunal (e.g. 218 polychaetes) were further removed from the top 90%. We chose the taxa that made up the top 90% 219 total biomass for each survey to address the basic ecological function of the epibenthic invertebrates 220 that dominate areas in and adjacent to Barrow Canyon. Detailed life history information for even the 221 dominant invertebrate taxa in the sub-arctic and Arctic is sparse at best. The 10% of invertebrate taxa 222 not used in the analysis were even more difficult to collect life history information on and often times 223 would reflect the life histories of those taxa found in the top 90%, since the family level was often used 224 in assigning trait modalities. 225 Ten biological traits that summarize basic biological characteristics (size, shape) as well as

fundamental trophic (feeding type and mechanisms), reproductive and life history ecology (dispersal and larval types) were selected and divided into 31 trait modalities (Table 2). The selected traits are common to several recent studies using biological traits analysis (Bremner et al., 2003; Haybach et al.,

2004; Hewitt et al., 2008; Paganelli et al., 2012), and maximize some of the fundamental differences in
the biology and ecology of species. The only biological trait that was quantitatively measured was Size,
determined as the average weight per individual animal for each taxon calculated from the catch. An
extensive literature search was used to assign each taxon to a category in each of the ten trait groups.
When little or no information on a species' life history could be found, information from the genus level
or species within the same family was used.

Three matrices were produced for the analyses: 1) a haul by taxon biomass (CPUE) matrix, 2) taxon by trait matrix, and 3) haul by trait matrix. The third matrix was a product of the haul by taxon biomass matrix and taxon by trait matrix (Bremner et al., 2003). This was constructed by multiplying trait modalities for each taxon present in a haul by its CPUE in that haul, and then summing the biomass across each trait modality over all taxa present in that haul to obtain a single value for each trait modality in each haul. Essentially, this weights the occurrence of a biological trait or trait modality across taxa at the haul level (Charvet et al., 1998).

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243 2.5. Numerical model integration

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245 In order to depict the mean regional circulation field for the purpose of providing additional 246 environmental context to the biological analyses and interpretations, we compiled results from a 3-247 dimensional ocean and ice circulation numerical hindcast model (Curchitser et al., 2013; Danielson et al., 248 2016b), which was integrated over 2005 to 2011 within the Regional Ocean Modeling System (ROMS) 249 framework (Fig. 2). Quantitative comparisons of model output to year-long in situ mooring-based 250 current velocity data show that the model reproduces (at the 95% confidence level) the observed mean 251 velocity vector components at nearly all of the 19 Beaufort and Chukchi evaluation sites (Curchitser et 252 al., 2013). Results from the new integration (Danielson et al., 2016b) and this study are driven with the

253 same initial conditions, boundary conditions and atmospheric forcing described in Curchitser et al. 254 (2013). A number of improvements were implemented in the present version including extending the 255 Pacific side southern boundary from Bering Strait to south of the Bering Sea, an improved sea-ice 256 formulation, and more realistic coastal river discharges (Danielson et al., 2016b). Model horizontal 257 resolution is about 6 km in the Barrow Canyon region and there are 50 terrain-following layers in the 258 vertical dimension. Several model-derived variables were included in the initial analyses for all three 259 surveys: speed of the near-bottom current averaged over the day of sampling (Speed Day of Sample), 260 eastward velocity component averaged over the day of sampling (East Velocity), northward velocity 261 component averaged over the day of sampling (North Velocity), year-long mean near-bottom speed at 262 this sampling site (Mean Year Speed), and year-long mean near-bottom speed variance at this sampling 263 site (Mean Year Speed Var).

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265 2.6. Statistical Analysis

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The taxa in the top 90% by biomass were analyzed in two ways: 1) a canonical correspondence analysis (CCA) using taxon biomass estimates and environmental variables (Cajo and Braak, 1986; McGarigal et al., 2000), and 2) a fuzzy correspondence analysis (FCA) to quantify the variation in biological trait distribution for each survey (Chevenet et al., 1994).

For the taxonomic distribution and environmental variables analysis, the taxon biomass matrix was ordinated using a constrained ordination (i.e. a CCA), and displays only the variation that can be explained by the constraining variables (e.g. temperature, depth) using Chi-square distances. Further, the haul scores are constrained to be linear combinations of the constraining variables. Because of the small sample size in the Beaufort Sea LN and Beaufort Sea UN surveys, the environmental variables examined were limited to those that were not tightly correlated. For example, density and salinity are

277 tightly correlated, therefore we chose a single variable, salinity, for the analysis. Even though 278 intercorrelated variables do not affect the CCA, a reduced number of variables can aid in interpretability, 279 especially when the number of variables approaches the number of samples, as in the case of the 280 Beaufort LN and UN hauls (McGarigal et al. 2000). Also, if the ordination contains as many variables as 281 samples, the ordination is no longer "constrained" by the environmental variables, also termed 282 overfitting. The initial CCA analysis was completed for each survey and included all variables (9 for the 283 Chukchi and 8 for the Beaufort LN and UN surveys). Initially, we performed a variance inflation factor 284 (VIF) test, which measures the extent of multicollinearity between variables. This aided in informing 285 which variables would be used in the final CCA analysis and which variables could be eliminated (i.e. 286 high VIF). Additionally, variables whose arrows were in close proximity (same length and direction), a 287 single variable was selected for further analysis. In all three surveys, salinity and bottom depth were 288 tightly correlated along with year-long mean near-bottom speed at the sampling site (Mean Year Speed) 289 and year-long mean near-bottom speed variance at the sampling site (Mean Year Speed Var); the variables used in the model were bottom depth and Mean Year Speed. In the Chukchi survey, bottom 290 291 hardness and Mean Year Speed were tightly correlated, bottom hardness was the variable used in the 292 model for further analysis. After the variables were selected, the final CCA was performed on the taxon 293 biomass and the selected variables (termed the "full model"). The first two axes from the final CCA were 294 permutated (x=999) and an ANOVA-like test ("pseudo-F statistic") was used to determine those 295 variables that were most significant at p<0.05 (Dray and Dufour, 2007a, b). This analysis tested the 296 reduced model's results after removing a variable against the full model's results. 297 For the biological traits analysis, the haul by traits matrix was ordinated using fuzzy coding 298 correspondence analysis (FCA) (Chevenet et al., 1994). It is used to assign multiple modalities to a single

300 history stages. For example, a gastropod species may be both predator (70% of the time) and scavenger

taxon using percentages. This process accounts for differences in trait modalities within and among life

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301 (30% of the time) during the adult phase. In the absence of this detailed information for many of the 302 Arctic species in this analysis, we instead chose a binary coding procedure where each taxon is assigned 303 to one category within each biological trait based on what the taxon exhibits a majority of the time. For 304 example, the species Neptunea heros was classified as a predator under the trait modality "feeding 305 mechanism"; however, this species may also be an "opportunist/scavenger" at times. FCA is an 306 extension of correspondence analysis (CA) that incorporates discrete variables such as biological traits 307 that are either coded as binary (i.e., 0,1) or fuzzy (i.e., 0.3, 0.7). FCA is able to correct for the fact that 308 each biological trait (e.g. Body Design) has multiple trait modalities (e.g. "soft", "hard shell") and the 309 sum of all trait modalities within a biological trait can be no more than one. Even though biological 310 traits data in this analysis were binary and not coded as fuzzy, this sets up the framework for 311 incorporating additional trait information should it become available. The results of an FCA explain the 312 amount of variation in trait distribution on each axis. To interpret the results, the relationship of hauls 313 (rows) can only be compared to other hauls (rows) and columns (traits) to other columns (traits). Hauls 314 that have similar FCA coordinates and are close to one another on the ordination plot are similar relative 315 to the frequency of traits. Since the haul by traits matrix table contained the same trait modalities for all 316 three surveys, a Kruskal-Wallis test was used to determine if the first axis scores from the FCA ordination 317 for all three surveys differed from one another. The first axis scores account for the greatest amount of 318 variation. No statistical difference would suggest that the variation in biological trait distributions within 319 in each of three surveys were the same. To quantify how much of the variance was accounted for by 320 each biological trait and trait modalities (Table 2) on each axis, correlation ratios and eigenvalues were 321 calculated for each biological trait. Correlation ratios represent percentage of variance accounted for by 322 a given axis. The eigenvalues can be considered the amount of variance as part of the total variance (i.e. 323 total inertia) accounted for by each axis.

324	Statistical analysis and graphical results were carried out in the statistical program R (R
325	Development Core Team 2014), version 3.1.2, with R packages ade4 (Chessel and Dufour, 2004; Dray
326	and Dufour, 2007a, b) and Vegan (Oksanen et al., 2015). All maps were created in ESRI ArcGIS version
327	10.3.
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329	3. Results
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331	3.1. Diversity and dominant taxa
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333	The number of taxa in the Chukchi Sea survey ranged from 26-57 per haul with a total of 215
334	taxa identified, of those 151 were identified to species. The number of taxa ranged from 27-58 per haul
335	in the Beaufort Sea LN survey with a total of 132 taxa. The number of taxa ranged from 35-50 per haul
336	in the Beaufort Sea UN survey with a total of 109 taxa. In total, 17 taxa made up 90% of the total
337	biomass in the Chukchi Sea survey, 10 taxa in the Beaufort Sea LN survey, and 11 taxa in the Beaufort
338	Sea UN survey (Table 1). Within the species contributing the top 90% wet weight biomass, the
339	northeastern Chukchi Sea survey and the western Beaufort Sea surveys (LN and UN combined) shared at
340	least six benthic invertebrate species.
341	The CPUE estimates in the Chukchi Sea hauls were highest at the western-most portion of the
342	study area (Fig. 1). The dominant invertebrates in the Chukchi Sea hauls from nearshore to the canyon
343	were Gorgonocephalus spp. (basket star), and other ophiuroids, and four species of large gastropods
344	(Fig. 3) (Table 1). Of the other biomass-dominant taxa, the sea star, Solaster dawsoni arcticus, occurred
345	in all the Chukchi Sea hauls (Fig. 3) while the mud star Ctenodiscus crispatus, only occurred on the outer
346	shelf in the north part of the Chukchi study area (Fig. 3). The sea cucumber, <i>Psolus peronii</i> , was only
347	present in the western portion of the Chukchi study area (Fig. 3).

348 In order to spatially note the taxonomic composition for all three surveys, several taxa were 349 lumped into a single group and were mapped using pie charts (Fig. 3). The Beaufort Sea LN hauls 350 occurred predominantly on the Beaufort slope and in Barrow Canyon (>200 m) and were dominated by 351 brittle stars (mostly Ophiura sarsii), snow crab (Chionoecetes opilio) and gastropods (Fig. 3). Total CPUE 352 estimates were consistently high in the mouth of Barrow Canyon and along the western Beaufort Sea 353 shelf break in water deeper than 300 m (Figs. 2 and 3). Overall, the CPUE estimates for the Beaufort Sea 354 LN hauls were two to three times greater than the estimates for both the Beaufort Sea UN and Chukchi 355 Sea hauls. This discrepancy is in part related to the smaller mesh size (Fig. 1) used for the Beaufort Sea 356 LN hauls; of 12 hauls, only two (12 and 13) had CPUE estimates considerably lower than all other hauls. 357 The Beaufort Sea UN hauls occurred mostly on the shelf and were dominated by a sea cucumber 358 (Psolus peronii) and hermit crab (Pagurus trigonocheirus) in the two western hauls, next to Barrow 359 Canyon (Fig. 3) (Table 1). The catch composition transitioned to hauls dominated by the mud star 360 (Ctenodiscus crispatus) and the same hermit crab (Pagurus trigonocheirus) in the central part of the 361 study area to hauls dominated by the sea peach (Halocynthia aurantium) and two genera of 362 demosponges, Polymastia and Vulcanella, in the east part of the study area (Fig. 3) (Table 1). The CPUE 363 estimates varied across all hauls, but was, with the exception of haul 17, lower than the western- and 364 southernmost Chukchi Sea hauls, and there was no quantitative pattern relating to depth or location 365 (Fig. 1).

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## 367 3.2. Environmental variables and epifaunal biomass

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The bottom temperature ranges observed in the Chukchi Sea survey during the sampling period ranged from -1.72 to 2.68°C, the bottom salinity ranged from 30.93 to 33.28, and bottom depths ranged from 26 to 155 m. Bottom hardness measurements at or near the bottom trawl stations ranged from

3.5 (least hard) to 5.9 (most hard) within the Chukchi Sea study area. During the Beaufort Sea LN
survey, the bottom temperatures ranged from -1.70 to 1.94°C, the bottom salinity ranged from 30.94 to
34.85, and bottom depths ranged from 47 to 445 m. During the Beaufort Sea UN survey, the bottom
temperatures ranged from -1.26 to 2.19°C, the bottom salinity ranged from 30.69 to 34.11, and bottom
depths ranged from 40 to 187 m.

377 The three environmental variables in the CCA accounted for 17% of the variability in taxon 378 biomass (Fig. 4a) (Table 3) on the first two axes, and the only significant term from the ANOVA was 379 bottom hardness on the CCA1 axis (p<0.05) (Table 3). All the hauls with a positive CCA1 axis score (Fig. 380 4a) corresponded to those hauls located either in or in close proximity to Barrow Canyon; this aligned 381 with both an increase in bottom hardness and current flow (Fig.1). The hauls with both a negative CCA1 382 and CCA2 score were those hauls located on the upper Chukchi shelf in the northern most stations (Figs. 383 1 and 4a, hauls 11, 12, 13, and 16). These hauls were also located close to 180° degrees from the 384 direction of bottom hardness, which indicates a decrease in bottom hardness in the location of these 385 hauls (Fig. 4a).

386 In the Beaufort Sea LN hauls four environmental variables were used in the CCA and accounted 387 for 46% of the variability in taxon biomass (Fig. 4b) (Table 3) on the first two axes. The ANOVA permutations resulted in three significant terms (p<0.05); bottom temperature, bottom depth, and 388 389 Mean Year Speed (Table 3). The arrows for bottom depth and Mean Year Speed are at a 180° angle 390 indicating these two variables are inversely related; at deeper hauls, current flow is slower (Fig. 4b). 391 Hauls 3, 11 and 12 (Fig. 4b) were at the shallowest stations but had the highest current flows whereas all 392 the hauls clustered in the upper right quadrant were the deepest stations and had the lowest current 393 flows.

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In the Beaufort Sea UN hauls, 48% of the variability on taxon biomass in the CCA was accounted for by bottom depth, North Velocity, and Mean Year Speed Var (Fig. 4c). The significant variables from the ANOVA were bottom depth and North Velocity (Table 3).

397

398 3.3. Biological Traits Analysis

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400 The FCA accounted for 58% of the variance in the distribution of traits for the Chukchi Sea hauls 401 (Figs. 5a-b) (Table 4). Hauls with a positive axis 1 score were dominated by large, predatory carnivores 402 that sexually reproduce and shed eggs (Fig. 5b). Flat bodied, predatory organisms with an endoskeleton 403 also dominated these hauls and similar to the results of the taxon biomass and environmental variables, 404 these traits occurred where there was an increase in hard bottom (Fig. 5a, black circle). Hauls 11-13 and 405 16 had a positive axis 2 score and grouped together; these hauls were dominated by small opportunistic 406 scavengers with a hard exoskeleton (Figs. 5a-b). These four hauls were located on the Chukchi shelf, 407 north of Barrow Canyon (Fig. 1).

408 The FCA accounted for 73% of the variance in the distribution of traits for the Beaufort Sea LN 409 hauls (Figs. 5c-d) (Table 4). Similar to the results based on taxon abundance and environmental 410 variables, the tight cluster of hauls with a negative axis 1 score were hauls that corresponded to the 411 stations >200 m in the Beaufort Sea survey (Fig. 5c, black circle). These hauls were dominated by small, 412 flat bodied deposit feeders, and predators with a hard exoskeleton, reflecting the dominance of brittle 413 stars and snow crab in these hauls (Figs. 5c-d). Those hauls with a positive axis 1 score were dominated 414 by soft, medium/large bodied, filter/suspension feeders (Fig. 5d). These stations were positioned from the slope onto the Beaufort shelf and, in general, were in water <200 m (Figs. 5c-d). 415

The FCA accounted for 79% of the variance in the distribution of traits for the Beaufort Sea UN
hauls (Fig. 5e-f) (Table 4). Unlike the clustering of hauls in the Chukchi and Beaufort LN surveys, the

418 Beaufort Sea UN survey showed little clustering among the hauls based on traits. This likely reflects the 419 higher degree of taxonomic diversity where many of the species do not overlap in trait similarities 420 compared to the Chukchi and Beaufort LN surveys (Fig. 5e). Traits with a positive axis 1 and axis 2 score 421 were dominated by soft body/protected, erect, and medium size traits and were associated with hauls 422 23 and 24 (axis 2) on the southern slope/shelf of Barrow Canyon, which also coincided with a 423 predominance of sea cucumbers in these two hauls (Fig 3c). Hauls that had a negative axis 2 score (17, 424 20, 21) occurred on the Beaufort Sea shelf in the east part of the Beaufort study area, and were 425 dominated by large, filter/suspension feeders that sexually reproduce and shed larvae (Figs. 5e-f). Flat 426 bodied deposit feeders dominated hauls 18 and 26 and were located on the Beaufort Sea shelf (Figs. 5e-427 f).

428 The axis score coordinates for each trait from the FCA analysis were plotted by survey to 429 visualize which trait scores accounted for the most variance in the distribution of hauls (i.e. points 430 located far from the zero coordinate for both axes). For example, the trait modality "small" for both the 431 Chukchi and Beaufort Sea UN surveys showed coordinate scores distant from zero on both axes, which 432 means this trait contributed to the observed variance in the pattern of haul distribution (Fig. 6). In 433 contrast, the variation in the distribution of the trait modality "large" showed little to no variation in trait distribution for all three surveys (i.e. all three points are located at or near the zero coordinate) 434 435 (Fig. 6). Biological traits and their subsequent modalities, that showed little to no variation in their 436 distribution across hauls, included Degree of Attachment, Mobility, Propagule Dispersal, and Larval 437 Dispersal (Fig. 6). The trait modality "deposit feed" showed large variation in the distribution of the trait 438 both within a survey (e.g. the Chukchi Sea, asterisk) and between surveys (e.g. the three points are 439 widely distributed); however, there was little variation in the distribution both within and between 440 surveys for the other trait modalities within Feeding Mechanism (e.g. "filter/sus", "opp/scavenger") (Fig. 441 6).

There was no significant difference in the variation of biological trait distribution among the
three surveys (Chukchi Sea, Beaufort Sea LN, Beaufort Sea UN) based on the results of the Kruskal-Wallis
test for significance on the first axis of the FCA ordination (p=0.90).

445 To further characterize the variation in the distribution of biological traits, we mapped biological 446 traits and the corresponding trait modalities with correlation ratios greater than 0.20 on axis 1 (Body 447 Design, Body Form, General Prey Type, and Feeding Mechanism) (Fig. 7a-d). (Table 4). Correlation ratio 448 scores greater than 0.20 can be considered those traits with the most variable distribution within a 449 survey. The Chukchi and Beaufort Sea LN surveys each had three traits that met this criteria, followed 450 by one trait in the Beaufort Sea UN survey that met this criteria (Fig. 7a-d). The biological traits with 451 ratios >0.2 showed the highest variation in their distribution within each survey (Fig. 7a-d) (Table 4). 452 There were three biological traits that explained the greatest amount of variation in their distribution on 453 axis 1 and that was Feeding Mechanism (Fig. 7c) in the Beaufort Sea LN and UN surveys and both Body 454 Design and General Prey Type (Figs. 7a and 7d) in the Chukchi and Beaufort Sea LN surveys (Table 4). 455 In the Chukchi Sea survey, the biological trait Body Design had the highest correlation ratio for axis 1 456 explaining 23% of the variability among the traits on axis 1, followed by Body Form and General Prey 457 Type (Table 4). In that, the "soft/protected" (Fig. 7a), "erect" (Fig. 7b), and "herbivore" (Fig. 7d) trait 458 modalities occurred in their highest proportion in the western part of the survey area and, in general, 459 these trait modalities did not occur on the southern slope of Barrow Canyon. Opposite this result, the 460 southern slope of Barrow Canyon in the Chukchi Sea survey was dominated by "endoskeleton/hard 461 shell" (Fig. 7a), "flat/round" (Fig. 7b), "carnivore/omnivore" (Fig. 7d) trait modalities. In the Beaufort Sea LN survey, Body Design and subsequent modalities also had the highest correlation ratio on axis 1, 462 likely due to the trait modality "soft/protected" only occurring at depths <200 m, on the Beaufort Sea 463 464 Shelf (Fig. 7a). Alternatively, the deep slope stations in the Beaufort Sea LN survey showed an absence 465 of the "soft/protected" (Fig. 7a), "erect" (Fig. 7b), "filter/suspension" (Fig. 7c), and "herbivore" (Fig. 7d)

466 trait modalities. The trait modality "deposit feeder" (Fig. 7c) that only occurred in the northeast portion 467 of the Chukchi Sea survey area, was the largest proportion of the Beaufort Sea LN hauls in the deep 468 slope stations. The trait modality "deposit feeders" was also not common on the Beaufort Sea shelf 469 (mostly UN) hauls, which were mostly dominated by "filter/suspension" feeders (Fig. 7c). In the 470 Beaufort Sea UN survey, the biological trait Feeding Mechanism was the only trait >0.2 with a 471 correlation ratio explaining 23% of the variability on axis 1 (Table 4), likely due to the increased 472 occurrence of the trait modality "deposit feeders" around the 50 m depth contour (Fig. 7c, 3 hauls). 473 We selected the two biological traits with ratios less than 0.05 on axis 1 (Propagule Dispersal 474 and Larval Development) for all three surveys to further illustrate patterns in trait distribution between 475 regions (Figs. 7e-f) (Table 4). These two biological traits and subsequent trait modalities showed the 476 least variation in their distribution within each survey. Even though the distribution of trait modalities 477 within a survey was homogenous for these low correlation ratios, the proportion of trait modalities 478 between surveys was quite variable (Figs 7e-f). Specifically, the biological trait Larval Development had 479 a higher proportion of "planktotrophic" larvae (long pelagic larval durations) in the Beaufort Sea LN 480 survey than the Chukchi Survey based on the pie charts (Fig. 7e). In contrast, the Chukchi survey had a 481 higher proportion of "direct" development (low dispersal potential) and "lecithotrophic" larvae (yolk 482 sack attached as source of nutrition; shorter dispersal range than "planktotrophic") throughout the 483 survey area compared to lower proportions in waters > 200 m in the Beaufort Sea LN survey (Fig. 7e). In 484 general, the trait modality "direct" was homogenously distributed and in every haul except one in the 485 Chukchi Sea survey, compared to the other two surveys where the trait is more patchily distributed (e.g. 486 a few hauls in the Beaufort Sea LN and UN surveys have high proportions). The trait modality 487 "substrate" occurs in greater proportions in the Chukchi Sea survey compared to the proportions this 488 trait occurs in both the Beaufort Sea LN and UN surveys (Fig. 7f). In general, the trait modality "pelagic"

was dominate in most of the Chukchi survey hauls and dominated all hauls in the Beaufort Sea surveys(Fig. 7f).

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492 **4. Discussion** 

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494 4.1. Environmental variables and epibenthic invertebrate biomass

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496 The epibenthos in the Arctic's Barrow Canyon are both taxonomically and functionally diverse 497 reflecting the high levels of production and the confluence of three water masses into the canyon. The 498 environmental variables used in this study accounted for a moderate portion of the variance (<47%) in 499 benthic community structure and these results generally aligned with recent studies in the northeast 500 Chukchi and Beaufort seas. In this study, depth was an important factor in the deeper Beaufort Sea 501 stations (Beaufort UN) in that some of the largest biomass estimates from the Beaufort survey occurred 502 at the deepest depths and this result was not entirely due to gear differences (Rand and Logerwell, 503 2011). Ravelo et al. (2015) also showed depth was an important factor in structuring the benthic 504 invertebrate community of the Beaufort Sea shelf and the northeastern Chukchi Sea. In addition, 505 previous studies also showed correlative relationships between water depth and temperature and the 506 structuring of benthic communities (Blanchard et al., 2013; Ravelo et al., 2014). 507 Although bottom hardness measurements were not available for the Beaufort Sea in our study, 508 sediment characteristics are also important factors in structuring the epibenthic communities on the 509 Beaufort Sea shelf (Ravelo et al., 2015) and were influential in the northeast Chukchi Sea, both in our 510 study and Ravelo et al. (2014). In this region, it is well documented that variation in current velocities 511 can act to deposit or carry finer sediments downstream (Darby et al., 2009); in this study, we noted that

512 there was a decrease in bottom hardness on the north part of the Chukchi shelf, suggesting this part of

513 the study area was comprised of finer sediments. This was corroborated by Grebmeier and Cooper 514 (2014) who reported that sediment becomes finer from the west side of Barrow Canyon, onto the 515 offshore Chukchi (Fig. 8). One of the variables we examined that represented current velocity in our 516 study was "speed (velocity) on the day of sampling" and the results were narrowly insignificant in the 517 Chukchi Study area (p=0.059). In this study area of the Chukchi Sea, current velocity does coincide with 518 the Pacific-origin WW (PWW) that flows from the northeastern shelf of the Chukchi Sea but then shifts 519 from the west side of Barrow Canyon to the eastern side as it heads northward into the Pacific Arctic 520 Basin (Gong and Pickart, 2015). This water mass flow erodes the fine sediment especially in the central 521 Canyon and leaves coarser sediments (Pisareva et al., 2015).

522 Our analysis for the Chukchi Sea shows that the faunal distribution patterns are in part related 523 to these sediment and associated current velocity patterns, but the moderate to low correlation factors 524 suggest that other factors, such as competitive interactions, may also influence faunal distribution 525 patterns as examples illustrate here. The coarser sediment and higher flow is associated with taxa that 526 are larger bodied such as the basket star, Gorgonocephalus spp., gastropods, and sea stars (Asteroidea). 527 The basket star, Gorgonocephalus spp., is known to inhabit areas of high current since they filter 528 organisms from the water column (Patent, 1970). In the Chukchi Sea, Gorgonocephalus spp., was 529 almost exclusively located in this higher current flow (i.e. bottom hardness and Mean Year Speed were 530 correlated). The dominance of predatory sea stars within this PWW and coarser sediment coincides with 531 both the absence of snow crabs, possibly due to competitive interactions, and a marked increase in 532 nutrient rich waters transported from the Pacific into Barrow Canyon (Grebmeier et al., 2006). Another example of faunal changes with sediment and current velocity patterns is the presence of the filter-533 534 feeding sea cucumber, Psolus peronii. This species of sea cucumber occurred in high abundances (Figs. 1 535 and 3) in areas of higher current velocity and increased bottom hardness (Fig. 2). The predatory lyre 536 crab, Hyas coarctatus, also occurs primarily west and on the Chukchi Sea Shelf and does not co-occur

537 with the basket star Gorgonocephalus spp.; possibly, this may indicate avoidance of sea stars and 538 gastropods due to competition and/or a preference for finer sediment and lower current velocities. 539 Only four hauls captured the deposit feeding mud star, Ctenodiscus crispatus, in the northeast corner of 540 the Chukchi Sea study area. This result may indicate that this portion of the Chukchi Sea study area may 541 have the finest sediment and lowest current velocities as supported in Figs. 2 and 8. The brittle star 542 Ophiura sarsii occurred in very low numbers in the northeastern Chukchi Sea survey even though it was 543 dominant in several other studies in the Chukchi Sea (Frost and Lowry, 1983; Ambrose et al., 2001; 544 Feder et al., 2005; Bluhm et al., 2009; Ravelo et al., 2014). This does not appear to be an artifact of the 545 sampling method because this species was captured with the same gear in the Beaufort Sea UN hauls. 546 Possibly, this may be due to competitive interactions between these large, predatory taxa in the central 547 part of Barrow Canyon.

548 We observed another taxonomic transition between the Barrow Canyon slope in the Beaufort 549 Sea and the Beaufort Sea Shelf. There is higher benthic biomass on the Beaufort Sea slope and a shift in 550 dominant taxa between the Beaufort slope (brittle stars and snow crab) and the Beaufort Shelf (sea 551 cucumbers, mussels, mud stars, etc.). This taxonomic transition coincides with a transition from the 552 PWW and Chukchi summer water (CSW) entering Barrow Canyon in the Chukchi Sea to both the 553 Beaufort shelfbreak jet <200 m, and the warmer more saline Atlantic water that dominates depths >200 554 m along the Beaufort Sea slope (Pickart, 2004; Weingartner et al., 2005b; Gong and Pickart, 2015). 555 Unlike the Chukchi Sea portion of Barrow Canyon, the brittle star Ophiura sarsii, and the snow crab, 556 Chionoecetes opilio, dominate these deeper depths (>200 m) that exit Barrow Canyon and lead into the 557 deep Arctic basin. Our results also illustrated an inverse relationship between depth and current velocity 558 in that these deeper basin stations also experience, on average, lower currents than those stations on 559 the Beaufort slope and shelf. Although we do not know the exact mechanism influencing these 560 distributions, they are consistent with previous findings by Frost and Lowry (1983) and Ravelo et al.

561 (2015), and we hypothesize that *O. sarsii* and snow crabs prefer the warmer, slower, and saltier Atlantic
562 influenced water.

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4.2. Patterns in biological and functional traits

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566 Our two surveys in the Chukchi and Beaufort seas encompass most of Barrow Canyon, sharing 567 some of the hydrographic characteristics, but they only share a portion of the biomass-dominant taxa. 568 Because of this result, we asked if these two study areas were, in fact, functionally different or shared 569 similarities. Although the regional comparison is somewhat biased by the different mesh size in the 570 Beaufort Sea LN survey, we were able to shed light on the variability of the biological traits both within 571 and adjacent to Barrow Canyon. The fact that there was not a biological trait or trait modality exclusive 572 to either system infers that these two systems share similarities in their biological processes which may 573 be surprising given the differences in taxonomy between the two study areas. This is one of the 574 fundamental benefits to a biological traits analysis; even though the taxonomic composition and 575 distribution is different, the underlying functional processes are similar.

576 Of the ten biological traits examined, however, four traits and their respective trait modalities 577 were considered variable enough within their respective study area to further examine patterns. The 578 four biological traits, Body Design, Body Form, Feeding Mechanism, and General Prey Type share some 579 characteristics. For example, the trait modality "soft/protected" often coincides with the trait 580 modalities "erect", and "filter/suspension" feeding. Similar to the connection in water masses and 581 taxonomic distribution, the variability we observed in trait distribution can in part be attributed to 582 habitat heterogeneity and transitions (e.g. sediment, hydrography) that occur in this region. The 583 Chukchi Sea survey area contained a high prominence of filter/suspension feeders in line with the 584 convergence of the PWW and increased current velocities as this water mass accelerates north into

585 Barrow Canyon. High current velocities tend to contain high loads of suspended particles that serve as 586 food for these filter feeders. These filter/suspension feeders maintain a presence in Barrow Canyon as it 587 enters into the Beaufort and extends onto the shelf with the PWW transitioning into the Beaufort 588 shelfbreak jet. The filter/suspension feeders do not extend to the Barrow Canyon slope or waters >200 589 m where deposit feeders dominate. This is likely related to the lower suspended particle content in 590 slowing waters and the fine-grained, less eroded sediment that is more suitable for deposit feeders. 591 This supports the taxonomic distribution results and again highlights an important transition zone from 592 the fast flowing Beaufort Sea shelf environment (<200 m) dominated by filter/suspension traits to one 593 dominated by the deposit feeding trait within the deeper Atlantic water. These relationships between 594 feeding mechanism and flow velocity match results of a study by Pisareva et al. (2015) that used the 595 feeding modes of benthic fauna and sediment characteristics as indicators of hydrographic flow on 596 varying time scales. They also concluded that, in general, benthic suspension feeders were associated 597 with regions of stronger flow and deposit feeders with regions with weaker flow (Pisareva et al., 2015), 598 and that these faunal patterns were reflective of the long-term flow velocities. Further, Pisareva et al. 599 (2015) found that there was a higher proportion of suspension feeding fauna in the central Barrow 600 Canyon and a higher proportion of deposit feeding taxa outside the Canyon based on flow speed and 601 sediment grain size. This shift in both taxa and traits from the head of Barrow Canyon in the Chukchi 602 Sea to the Beaufort Sea shelf occurs within an oceanographically complex region. The hydrography at 603 the head of Barrow Canyon in the Chukchi Sea hosts both vertical and horizontal transition zones that 604 include lateral fronts, vertical stratifications and the influence of swift flows interacting with steep 605 bottom topography gradients (Fig. 2).

Some biological traits had little variability in their distribution within a survey but differed
between surveys. For example, those traits representing reproduction modes and early life history
stages (Propagule Dispersal and Larval Development) showed little to no variation within a survey. It is

609 possible that these traits are less influenced by currents, water mass properties, and sediment 610 composition than morphological or feeding traits. In contrast to this result, between-survey differences 611 included a high proportion of "direct" development in the Chukchi Sea survey that was observed to a 612 lesser degree in the Beaufort Sea. This trait may contribute to some of the differences we observed in 613 the taxonomy between the Chukchi and Beaufort parts of Barrow Canyon. The retention of "direct" 614 developers (low dispersal) at the head of Barrow Canyon may cause these taxa to remain in the Chukchi 615 Sea and not be advected into the Beaufort Sea. Whether the prominence of taxa with this Larval 616 Development trait at the head of Barrow Canyon are fostered by conditions such as nutrients and 617 upwelling, or whether these differences are driven by competitive interactions within the community 618 itself, is difficult to determine. Although we do not quantitatively compare the absolute biomass of these traits between surveys (e.g. there is "more" of a trait in a survey), the fact that some of these trait 619 620 modalities are evenly distributed within a survey is interesting. Life history strategies in marine benthic 621 invertebrates, in particular reproduction, are especially complex processes with multiple tradeoffs that 622 can translate into differences, for example, in fecundity among closely related species inhabiting 623 different habitats (Llodra, 2002). This suggests the possibility that the processes affecting those taxa at 624 the head of Barrow Canyon are not the same as those in the Beaufort Sea side of Barrow Canyon in 625 depths less than 200 m, even though the water mass transport, bathymetry, and current velocities 626 appear to be similar and given these sites are within only 60 km of each other.

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### 628 4.3. Study limitations and recommendations for future studies

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The limitations of our functional traits analysis is that in the Arctic, life history and other
information on benthic invertebrates is particularly sparse. In the absence of life history traits for many
of these Arctic species, our approach was, therefore, to apply known trait modalities from those species

633 within the same genus or family. Given the lack of species-specific knowledge, we used traits that 634 focused on the general functions of an organism, like Feeding Mechanism, rather than those traits 635 describing complex processes like growth rates. We recommend that future studies on impacts of 636 climate change and other human impacts also include those traits characteristic of Arctic fauna such as 637 (slow) growth rates, (high) longevity and (late) age at maturity. Also, the focus of this study was on the 638 top 90% by biomass of the epibenthic invertebrates; this eliminated 80-90% of the taxa that were 639 captured in the surveys, shifting the focus to organisms that make up the largest biomass and for which 640 information on biological traits could be gathered. The weakness in focusing on only a relatively small 641 number of taxa, in this case <20 per survey area, is that information was lost on taxa or species that may 642 have been common to all stations but comprised a very small fraction of the total biomass at that 643 station and across the survey area. These taxa or species could be filling a small, but important, 644 ecological niche in the overall function of a local community. The BTA approach could also be refined by 645 using fuzzy instead of binary coding for the traits since many species exhibit more than one trait 646 modality within a trait category. For example, most sea stars were placed in the "predator" modality as 647 their primary method of feeding, but they can also be "opportunistic/scavengers" and "deposit 648 feeders"; fuzzy coding would allow better resolution of the complexity of their functional contributions 649 at the community level.

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#### 651 4.3 Conclusions and monitoring recommendations

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It is well documented that the effects of climate change are accelerated in the Arctic (Pithan and
Mauritsen, 2014). What remains unclear, is the magnitude in which these rapidly changing conditions
will affect marine life in the coming years. For example, Cross et al. (this issue) has shown that
decreased saturation rates of calcium carbonate (e.g. ocean acidification) from anthropogenic changes

657 in atmospheric carbon dioxide, has fundamentally altered the duration and intensity of the PWW. Both 658 in- and epi- faunal invertebrates are particularly vulnerable to changes in ocean acidification. Since the 659 PWW is the primary source of transport from the Pacific to the Arctic Basin, through Barrow Canyon, a 660 well-documented biological "hotspot" for micro- to macro- fauna (Moore et al., 2014; Moore and 661 Stabeno, 2015; Grebmeier et al., 2015), it is an especially important region of the Pacific Arctic to 662 monitor for the effects of climate change. A traits analysis, along with traditional taxonomic methods, 663 has become a useful tool in ecosystem monitoring and management (Usseglio-Polatera et al., 2000; 664 Doledec et al., 2006; Tomanova et al., 2008). In these studies, researchers used community composition 665 analyses (taxonomic and/or traits) and changes therein as a direct reflection of ecosystem health. 666 Although changes in taxonomic diversity can be indicative of change that is occurring, extending this to 667 analyzing biological traits makes a direct connection between environmental processes and the ecological functioning of the systems organisms. 668

669 This study improved the knowledge of epibenthic communities in the Barrow Canyon area, 670 which has previously not been well resolved nor their connections to the complex regional 671 hydrodynamics. This effort contributes to the baseline data in terms of both species and functional 672 diversity of the epibenthic invertebrate communities within this region. Monitoring the epibenthic 673 invertebrate community at these "hotspots" is in line with the recently established Distributed 674 Biological Observatory (Grebmeier et al. 2010, 2015), the US-wide Marine Biodiversity Observation 675 Network (MBON) including its Arctic component and the Conservation of Arctic Flora and Fauna's 676 Circumpolar Biodiversity Monitoring Program. Benthic taxa can be excellent proxies for change over time because they are relatively long lived at high latitudes and can be indicative of changes in sediment 677 678 characteristics and water mass influences due to warming, acidification, and the loss of sea ice (Kortsch 679 et al., 2012; Grebmeier et al., 2015a; Cross et al., this issue). We recommend biological trait analysis 680 approaches be included in such monitoring efforts.

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# Table 1 The taxa listed comprise 90% of the total biomass of epibenthic invertebrates, used in all

analyses, separated by survey and, within the Beaufort survey, by lined net (LN) or unlined net (UN). The

920 mean is the cube-root transformed CPUE kg/km<sup>2</sup> with the standard deviation kg/km<sup>2</sup>.

Species	Common name (class)	Survey	Moon	Std day
Sheries		Survey	kg/km <sup>2</sup>	kg/km <sup>2</sup>
Berinaius berinaii	sea snail (gastropod)	Chukchi	2.22	1.56
Bryozoa	bryozoa	Chukchi	1.42	2.15
, Crossaster papposus	, sea star (asteroidea)	Chukchi	3.52	1.27
Ctenodiscus crispatus	mud star	Chukchi	1.60	3.25
Eualus sp.	shrimp	Chukchi	0.57	1.82
Gersemia rubiformis	coral	Chukchi	2.26	3.08
Gorgonocephalus spp.	basket star (ophiuroid)	Chukchi	4.79	6.96
Hyas coarctatus	lyre crab (crab)	Chukchi	2.34	2.40
Musculus discors	mussel	Chukchi	1.40	4.21
Neptunea heros	sea snail (gastropod)	Chukchi	3.61	2.65
Neptunea ventricosa	sea snail (gastropod)	Chukchi	1.84	2.05
Pagurus trigonocheirus	hermit crab	Chukchi	2.17	2.96
Psolus peronii	sea cucumber	Chukchi	3.00	6.76
Pyrulofusus deformis	sea snail (gastropod)	Chukchi	2.22	1.85
Solaster dawsoni arcticus	sea star (asteroidea)	Chukchi	4.21	2.98
Strongylocentrotus droebachiensis	sea urchin	Chukchi	1.73	3.25
Unicina crassicornis	sea anenome	Спикспі	1.70	2.77
Actiniaria	sea anenome	Beaufort LN	5.01	7.32
Buccinum polare	sea snail (gastropod)	Beaufort LN	5.86	5.30
Chionoecetes opilio	snow crab (crab)	Beaufort LN	16.16	9.57
Ctenodiscus crispatus	mud star	Beaufort LN	9.32	8.93
Gorgonocephalus spp.	basket star (ophiuroid)	Beaufort LN	2.79	5.61
Musculus sp.	mussel	Beaufort LN	5.97	11.32
Neptunea sp.	sea snail (gastropod)	Beaufort LN	6.89	3.12
*Ophiura sp.	brittle star (ophiuroid)	Beaufort LN	25.55	15.69
Psolus peronii	sea cucumber	Beaufort LN	5.31	6.89
Strongylocentrotus sp.	sea urchin	Beaufort LN	3.48	6.60
Volut middendorfii and	sea snail (gastropod)	Beaufort LN	7.05	4.42
Habevolutopsius				
Chionoecetes opilio	snow crab (crab)	Beaufort UN	2.78	1.77
Ctenodiscus crispatus	mud star	Beaufort UN	3.48	4.21
Halocynthia aurantium	sea peach	Beaufort UN	6.53	5.79
Hyas coarctatus	lyre crab (crab)	Beaufort UN	4.16	2.04
Neptunea heros	sea snail (gastropod)	Beaufort UN	1.44	3.07
Pagurus trigonocheirus	hermit crab	Beaufort UN	4.72	2.06
<i>Polymastia</i> sp.	sponge (demosponge)	Beaufort UN	2.33	4.98
Psolus peronii	sea cucumber	Beaufort UN	4.25	4.65
Pyrulofusus deformis	sea snail (gastropod)	Beaufort UN	2.36	2.58
Stomphia sp.	sea anenome	Beaufort UN	4.31	1.67
Strongylocentrotus sp.	sea urchin	Beaufort UN	3.46	5.45
Vulcanella sp.	sponge (demosponge)	Beaufort UN	3.86	6.79
*95% of the <i>Ophiura</i> sp. was comprised				
of Ophiura sarsi				

# Table 2 The biological traits for those taxa in Table 1 used in the FCA analysis. Within each Biological

Trait are several Trait Modalities; each unique combination of Trait and Modality is assigned a Trait
 Code.

Biological Trait	Trait Modalities	Trait Code
Size	small (<10 g)	S1
	medium (10-50 g)	S2
	large (>50 g)	S3
Body Design	soft	BD1
	soft/protected	BD2
	endoskeleton	BD3
	hard exoskeleton	BD4
	hard shell	BD5
Body Form	erect	BF1
	round	BF2
	flat	BF3
General Prey Type	herbivore	PT1
	omnivore	PT2
	carnivore	PT3
Feeding Mechanism	deposit feeder	FM1
	filter/suspension	FM2
	opportunist/scavenger	FM3
	predator	FM4
Degree of attachment	none	DA1
	semi-permanent	DA2
	permanent	DA3
Mobility	sessile	M1
	motile	M2
Propagule Dispersal	pelagic	PD1
	substrate	PD2
Reproductive Mode	sexual/release eggs	RM1
	sexual/release larvae	RM2
	sexual/brood	RM3
Larval Development	direct	LD1
	planktotrophic	LD2
	lecithotrophic	LD3

Table 3 Results of the CCA analysis on species abundance constrained by environmental variables. The

929 observed variance accounted for by the variables are shown in percent for both axes CCA1 and CCA2.

930 Variables that were used in the CCA are listed under "Variables Full Model" and the corresponding

significance level from the ANOVA test are listed under "Significance (p<0.05)"; significant values are in</li>bold.

Survey	CCA 1	CCA 2	Variables Full Model	Significance (p<0.05)
Chukchi Survey	12%	5%	Bottom Hardness	0.001
			Bottom Depth	0.085
			East Velocity	0.140
Beaufort LN Survey	24%	22%	Bottom Temperature	0.004
			Bottom Depth	0.004
			Mean Year Speed	0.010
			North Velocity	0.279
Beaufort UN Survey	28%	20%	Bottom Depth	0.020
			Mean Year Speed Var	0.168
			North Velocity	0.051

Table 4 Correlation Ratios for the FCA ordination of biological traits. The proportion of variance on each

935 axis that can be explained by the Trait Modalities (Table 2) within each Biological Trait. Ratios greater

than 20% on axis 1 are in bold and less than 5% on axis 1 in italics. Biological Traits that are mapped in

937 Fig. 7 are shown in bold text. The respective eigenvalues for each survey are given for each axis. The

total inertia (in parentheses) represents the total variance accounted for in an ordination. The percent

- 939 of variance that can be explained by each axis is listed by survey and axis.
- 940

Survey	Biological Traits	Axis 1	Axis 2
Chukchi	Size	0.110	0.070
Chukchi	Body Design	0.237	0.030
Chukchi	Body Form	0.218	0.013
Chukchi	General Prey Type	0.218	0.102
Chukchi	Feeding Mechanism	0.143	0.193
Chukchi	Degree of attachment	0.028	0.083
Chukchi	Mobility	0.028	0.083
Chukchi	Propagule Dispersal	0.033	0.010
Chukchi	Reproductive Mode	0.079	0.007
Chukchi	Larval Development	0.002	0.051
Percent of Variance	(Total Inertia)	37%	21% (0.299)
Eigenvalues		0.109	0.064
Beaufort LN	Size	0.082	0.035
Beaufort LN	Body Design	0.225	0.072
Beaufort LN	Body Form	0.196	0.022
Beaufort LN	General Prey Type	0.207	0.054
Beaufort LN	Feeding Mechanism	0.202	0.025
Beaufort LN	Degree of attachment	0.172	0.022
Beaufort LN	Mobility	0.179	0.015
Beaufort LN	Propagule Dispersal	0.000	0.001
Beaufort LN	Reproductive Mode	0.022	0.130
Beaufort LN	Larval Development	0.021	0.178
Percent of Variance	(Total Inertia)	51%	22% (0.253)
Eigenvalues		0.130	0.055
Beaufort UN	Size	0.141	0.107
Beaufort UN	Body Design	0.067	0.119
Beaufort UN	Body Form	0.084	0.054
Beaufort UN	General Prey Type	0.120	0.036
Beaufort UN	Feeding Mechanism	0.235	0.077
Beaufort UN	Degree of attachment	0.151	0.003
Beaufort UN	Mobility	0.151	0.003
Beaufort UN	Propagule Dispersal	0.025	0.019
Beaufort UN	Reproductive Mode	0.047	0.020
Beaufort UN	Larval Development	0.049	0.018
Percent of Variance	(Total Inertia)	55%	24% (0.193)
Eigenvalues		0.107	0.045

941 Figure captions

Fig. 1 The catch-per-unit-effort (CPUE metric tons (MT) wet weight/km<sup>2</sup>) by hauls in and near Barrow
Canyon. The Chukchi Sea survey was in 2013 (yellow) and the Beaufort Sea LN (green), and UN (orange)
surveys were in 2008; The Beaufort Sea LN refers to hauls sampled with a net liner and UN refers to
those hauls sampled with no net liner. All hauls are labeled in white text for reference. The CPUE scale
is different for the Beaufort Sea LN hauls; the lowest and highest CPUE estimate for the Beaufort UN and
Chukchi hauls also varies. Note: Haul 22, Beaufort Sea UN survey, is masked by haul 10, Beaufort Sea LN
survey due to a lower CPUE estimate.

Fig. 2 Average near bottom current velocities; results were produced by a 3-dimensional ocean and ice
circulation numerical hindcast model, which was integrated over 2005 to 2011 with the Regional Ocean
Modeling System (ROMS) framework. Large arrows were overlaid to show general current trajectories.

952 Fig. 3 Benthic invertebrate species composition for the top 90% by biomass (CPUE kg/km<sup>2</sup>) for the

953 Chukchi Sea, Beaufort Sea LN (pie charts are shown with thicker outline), and Beaufort Sea UN hauls.

954 The Beaufort Sea LN refers to hauls sampled with a net liner and UN refers to those hauls sampled with

955 no net liner. Each pie chart represents one haul. Several taxa were lumped together under a common956 name for better map interpretation (e.g., sea stars).

Fig. 4 Results of the canonical correspondence analysis (CCA) on species biomass (CPUE kg/km<sup>2</sup>) for the
Chukchi Sea (a), Beaufort Sea LN, and Beaufort Sea UN surveys. The Beaufort Sea LN refers to hauls
sampled with a net liner and UN refers to those hauls sampled with no net liner. The hauls (numbered
black dots) and environmental variables (arrows) are shown on each plot. Species associated with the
hauls are in light gray text. Significant variables are in bold text. The label "Bot" refers to "bottom". The
percentages refer to the amount of variance explained by each axis.

Fig. 5 Results of the fuzzy correspondence analysis (FCA) on the biological traits for the Chukchi Sea (a
and b) (asterisk), Beaufort Sea LN (c and d) (black dots), and Beaufort Sea UN (d and e) (triangles)
surveys. The Beaufort Sea LN refers to hauls sampled with a net liner and UN refers to those hauls
sampled with no net liner. Figures 5a, 5c, and 5e are the ordination results by haul and Figures 5b, 5d,
and 5f are the ordination results by traits (in light gray text) and aids in the interpretation of the results
in the first column. Hauls that are grouped can be considered similar in the biological traits they exhibit.
Trait codes are written out in full black text; some trait codes are truncated, refer to Table 2 for full

970 listing of traits. Hauls in the black circle indicated significance with bottom hardness in the Chukchi Sea971 (5a) and hauls >200 m depth in the Beaufort Sea LN (5c).

Fig. 6 Traits score results from the fuzzy correspondence analysis (FCA) on the biological traits for the
Chukchi Sea (asterisk), Beaufort Sea LN hauls (black dots), and Beaufort Sea UN hauls (triangles) surveys.
The Beaufort Sea LN refers to hauls sampled with a net liner and UN refers to those hauls sampled with
no net liner. This figure is and alternative representation of Figures 5b, 5d, and 5f and is a visual
comparison of the variation for each trait, in each survey, with vertical grouping by the Biological Trait
and subsequent Trait Modalities. Note: some trait codes are truncated (e.g., Opp/Scavenger), refer to
Table 2 for full listing of traits.

- 979 Fig. 7 Biological traits with correlations ratios above 0.20 (a-d) and correlation ratios below 0.05 (e and f)
- 980 are shown for hauls in the Chukchi Sea, Beaufort Sea LN (outlined in black), and Beaufort Sea UN
- 981 surveys. The Beaufort Sea LN refers to hauls sampled with a net liner and UN refers to those hauls
- 982 sampled with no net liner. Each pie chart represents one haul. Note: to illustrate all hauls on one map
- some pie charts two hauls from the Beaufort Sea UN survey are slightly displaced from the actual haul
- 984 location but a line connecting the chart to the location is shown.
- 985 Fig. 8 Map of the Chukchi Sea survey area overlaid with the acoustic transects for bottom hardness and
- 986 the PacMARS Surface Sediment Parameters (Grebmeier, J., and L. Cooper. 2016. PacMARS Surface
- 987 Sediment Parameters. Version 2.0. UCAR/NCAR-Earth Observing Laboratory.
- 988 <u>https://doi.org/10.5065/D6416V3G</u>). The measure of bottom hardness is from vessel acoustics and is
- 989 unitless; higher values imply harder bottom. "Modal Phi Size" refers to the sediment grain size mode.
- 990 This figure illustrates the transition from finer to coarser sediment from offshore (lower current velocity)
- 991 into Barrow Canyon and onto the shelf, where current velocities are greater.



Fig. 1



Fig. 2















Fig. 6







Fig. 8