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30 sympatric range. Here, we present evidence against trophic equivalence of these sibling species 31 based on flounder size and space-stratified analyses of stomach contents data from 5,724 32 flounders sampled from the eastern Bering Sea (EBS) during summer 2007-2016. We found 33 interspecific trophic niche separation occurred between flounders ≤39 cm fork length, before 34 convergence at larger size classes. Our findings are consistent with the hypothesis that foraging 35 efficiency differences arise due to divergence in gill raker counts. Arrowtooth Flounder have 36 more gill rakers than Kamchatka Flounder and consistently consumed more zooplankton than 37 Kamchatka Flounder, while Kamchatka Flounder typically consumed more benthic fishes. While 38 a benthivory-zooplanktivory axis of resource partitioning is common among sympatric trophic 39 polymorphs in freshwater ecosystems, our findings are novel for marine sibling species and 40 flatfishes (Pleuronectiformes). Abundance estimates for both Arrowtooth Flounder and 41 Kamchatka Flounder have substantially increased in Alaska in recent years, and we suggest 42 trophic niche separation alleviates interspecific competition, which may buffer sympatric 43 carrying capacities for Arrowtooth Flounder and Kamchatka Flounder.

44 45

#### 46 INTRODUCTION

47 Arrowtooth Flounder, *Atheresthes stomias* and Kamchatka Flounder, *Atheresthes*  48 *evermanni* are large-mouth flatfishes, sympatric in the Bering Sea, Aleutian Islands and 49 southwestern Gulf of Alaska. Morphological differences between species are subtle, such that 50 field identification confidence during stock assessment surveys conducted by the National 51 Marine Fisheries Service was only moderate through 2008 (Stevenson and Hoff 2009). The 52 primary morphological differences are the position of the left eye, which is on the dorsal margin 53 of the body in Arrowtooth Flounder and below the dorsal margin in Kamchatka Flounder, and 54 higher gill raker counts on each of the first three gill arches in Arrowtooth Flounder 55 (Wilimovsky et al. 1967; Yang 1988). Gill rakers of both species are lathe-shaped and of 56 moderate length (Yang 1991), but Arrowtooth Flounder have a combined 28-37 gill rakers on the 57 first three gill arches (mean: 31.4), while Kamchatka Flounder have 19-29 gill rakers (mean: 58 24.8; Yang 1988; M.S. Yang, unpublished data). 59 Interspecific morphological differences suggest foraging capabilities are likely to differ For external to the state of the first means the state of the means and convergence and represent and convergence in gill raker counts. Arrowtooth Flounder the means of the mean through the state and consistent with the hy

61 mutually beneficial ecological advantage where they co-occur. Phenotypic divergence of feeding 62 structures is common among sympatric sibling species and conspecific polymorphs (Smith and 63 Skúalson 1996; Ward et al. 2006), and can facilitate alleviation of resource competition through 64 resource specialization (Swanson et al. 2003). A common pattern of niche separation occurs 65 between sympatric benthic specialists and zooplankton specialists, where zooplankton specialists 66 have higher gill raker counts (Wimberger 1994; Smith and Skúalson 1996) that increase capture 67 efficiency for small-bodied zooplankton prey (MacNeill and Brand 1990; Sanderson et al. 1991; 68 Link and Hoff 1998; Roesch et al. 2013). Thus, Arrowtooth Flounder may be more efficient 69 zooplankton consumers than Kamchatka Flounder. However, the only prior comparative diet 70 study between species concluded Arrowtooth Flounder and Kamchatka Flounder are "trophically 71 equivalent," and that interspecific competition was unlikely to be important due to a high 72 abundance of prey resources in the eastern Bering Sea (Yang and Livingston 1986). 73 In this paper, we re-examine the trophic equivalency of Arrowtooth Flounder and 74 Kamchatka Flounder in the eastern Bering Sea (EBS) using a larger data set than was available 75 for the earlier comparative diet study. Yang and Livingston's (1986) study design may not have 76 allowed detection of interspecific trophic niche differences because their available sample size 77 was relatively small, covered a limited spatial extent, and was collected during a two-week 78 period in a single year. In addition, the importance of resource competition between *Atheresthes* 79 has likely increased since Yang and Livingston's (1986) study because aggregate Arrowtooth 80 Flounder and Kamchatka Flounder biomass has increased four-fold within their sympatric range 81 (Spies et al. 2016; Wilderbuer et al. 2016). Based on spatial biomass trends, Zador et al. (2011) 82 suggested that Arrowtooth Flounder may be near carrying capacity in some parts of the EBS. 83 Under these conditions, resource competition may have increased, and the expression of 84 morphology-based resource partitioning, or trophic niche separation, between Arrowtooth 85 Flounder and Kamchatka Flounder may be more pronounced (Svanbäck and Bolnick 2007). As 86 such, we address three main questions: 87 1. Do the trophic niches of Arrowtooth Flounder and Kamchatka Flounder differ within 88 their sympatric range? 89 2. If so, does trophic niche separation persist spatially and across predator size classes? 90 3. Are patterns of trophic niche partitioning consistent with expectation based on 91 interspect of the specialists and zooplankton specialists whose higher gill reker counts (Wimberger 1994; Smith and Skúalson efficiency for small-bodied zooplankton prey (MacNeill and Brand 168 Link and Hoff 1998; Roes

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93 METHODS

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### 95 *Study area and sample collection*

96 The eastern Bering Sea (EBS) is a large coastal ecosystem with a broad (~500 km) 97 continental shelf that gradually increases in depth to  $\sim$  200 m, before an abrupt depth increase 98 along the continental slope into the deep Aleutian Basin (Fig. 1). Strong seasonal dynamics and 99 inter-annual environmental variability influence EBS community spatial structure.

100 Diet data used in this study were collected during summer (June-August) bottom-trawl 101 surveys conducted by NOAA's Alaska Fisheries Science Center from 2007 through 2016. 102 Bottom-trawl surveys of the EBS shelf were conducted annually (e.g., see Conner et al. 2017), 103 and bottom-trawl surveys of the EBS slope were conducted biennially, except 2014 (Hoff 2016). 104 Survey designs and sampling gear differed for the EBS shelf and EBS slope due to differences in 105 stock assessment needs and substrate types. The EBS shelf survey sampled across a regularly-106 spaced grid (20 nmi x 20 nmi) of 376 stations, at ~15-200 m bottom depth, with higher density 107 sampling in areas of historically high crab abundance around St. Matthew, St. Paul and St. 108 George Islands (Lauth and Nichol 2013). EBS slope survey samples were randomly stratified by 109 sub-areas and depth (200-1200 m), and ~200 stations were sampled per survey (Hoff 2013). 110 Stomachs were collected from specimens of *Atheresthes* at a subset of survey stations 111 following a size-stratified sampling scheme (Livingston et al. 2017). Up to 15 stomachs were 112 collected per station. Fish were examined for evidence of regurgitation (i.e., prey in the mouth, 113 flaccid stomach) or net-feeding (i.e., undigested prey in the mouth), and whole stomachs were 196<br>
122 Ecosystem Modelling Sea (EBS) is a large coastal ecosystem with a broad (~500 km)<br>
197 continental shadelling traceases in depth to ~200 an, before an abropt depth increase<br>
198 along the continental slope into t

114 sampled from fish not displaying evidence of regurgitation or net-feeding. Predator species, fork 115 length (cm), sex and collection location were recorded for each sample. Stomachs were fixed in

116 neutral-buffered 10% formalin, then transferred to 70% ethanol for storage prior to stomach

117 contents analysis. Stomach samples used for this study were collected during 2007-2016 EBS

118 shelf surveys, and 2008, 2010, 2012 and 2016 EBS slope surveys. An average of 518 non-empty

119 *Atheresthes* stomachs were sampled per EBS shelf survey, and an average of 137 non-empty

120 stomachs were sampled per EBS slope survey (Table 1).

121 Stomach contents analysis was performed by analysts in the Resource Ecology and

123 Center (AFSC; Seattle, WA) following standardized laboratory protocols (Livingston et al. 2017).

124 Briefly, prey items were identified to the lowest practical taxon, fish and crab prey were

125 enumerated, and prey weights were recorded to the nearest milligram.

126

127 *Data analysis* 

128 Multivariate analyses were used to compare trophic niche variation between species, over 129 flounder size and space, based on broad prey categories. Prey categories were assigned based on 130 *a priori* knowledge of important prey categories identified in previous diet studies of Arrowtooth 131 Flounder and Kamchatka Flounder: benthic fishes, assorted pelagic fishes and squids, gadids 132 (Family Gadidae, mostly Walleye Pollock), shrimp (Suborders Dendrobranchiata and 133 Pleocyemata), unidentified fishes, non-shrimp benthic invertebrates, zooplankton (mostly Order 134 Euphausiacea), and "other." Stomachs containing fisheries discards (offal) were omitted prior to 135 analysis (1.6% of stomachs). Flounder groups were aggregated by species, size and spatial area. 136 Flounders were divided into five size classes based on fork length (≤19 cm, 20-29 cm, 30-39 cm, 137 40-49 cm, and  $\geq 50$  cm) to allow interspecific comparisons between flounders expected to have 138 similar stomach capacity and diet composition. Size classes were informed by the four size 139 classes (≤200 mm, 201-300 mm, 301-400 mm, >400 mm) used by Yang and Livingston (1986), 140 but were adjusted based on size class designations used in NOAA's standardized stomach 141 sampling protocol (Livingston et al. 2017), along with the availability of more samples from 142 larger-sized flounders for our analysis. Spatial areas were designated using five EBS shelf strata 143 (A-E) roughly delineating areas with distinct oceanographic features and community 144 composition (Stauffer 2004), and a 200-600 m bottom depth stratum along the EBS slope 145 (stratum F; Fig. 1). Flounder groups from areas where *Atheresthes* were scarce, sympatry was 146 weak, or sample sizes were small (< 10 non-empty stomachs) were excluded from analysis. 147 Overall, 10,110 specimens were assigned among 56 groups, of which 5,762 specimens from 55 148 groups had non-empty stomachs (Table 2). Adequate sample sizes for analyses were available 149 for 46 groups containing 5,724 non-empty stomachs (Table 2). Proportional wet weights of prey 150 were calculated for the groups of non-empty flounder stomachs (*%Wi,k,l,s*): alysis<br>
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\%W_{i,k,l,s} = \frac{\sum_{j=1}^{n_j} W_{i,j,k,l,s}}{\sum_{j=1}^{n_j} \sum_{i=1}^{n_i} W_{i,j,k,l,s}}
$$

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151 where *W* is wet weight of prey category *i*, from the stomach of flounder *j*, flounder size class *l*,

flounder species *s*, and spatial stratum *k*;  $n_i$  is the number of prey categories, and  $n_j$  is the number

153 of non-empty stomachs collected for the flounder group. A Bray-Curtis diet dissimilarity matrix

154 (Bray and Curtis 1957) was calculated from the flounder group proportional diet data for

155 multivariate statistical analysis.

156 Hierarchical clustering and ordination were used for multivariate statistical analysis of 157 diet data. Hierarchical clustering was conducted to determine which flounder groups had 158 greatest diet similarity and to evaluate which group dimensions (size, stratum, or flounder 159 species) influenced diet similarity. Ordination was performed to determine the relative influence 160 of prey categories on differences between flounder groups.

161 Similar-sized flounders have more similar diets than different-sized flounders, and a 162 strong ontogenetic shift from shrimp prey to fish prey has been identified for *Atheresthes* in the 163 EBS (Yang and Livingston 1986). Since the purpose of this analysis was to identify descriptive 164 differences in flounder group association, rather than synoptic patterns of ontogenetic diet shift, 165 we used space-dilating hierarchical clustering algorithms (flexible beta with β parameter varying 166 from -0.5 to -1.0, and complete linkage clustering) to increase contrast within size classes. 167 Space-dilating algorithms may increase the likelihood of producing small or orphan clusters, but 168 they can also preserve ecologically meaningful differences by requiring a more stringent 169 criterion for assigning cluster membership (Legendre and Legendre 2012). We used cophenetic 170 correlation to compare clustering algorithm performance. An appropriate number of clusters was 171 determined from examination of a scree plot of cluster dissimilarity attained across a range of 172 clusters, and through index-based evaluation of relevant clusters using R package 'NbClust' 173 (Charrad et al. 2014). 1815 multivariate saturateal analysis.<br>
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1927 dietatar-Hierarchical clustering was conducted to determin

174 After relevant clusters were identified, an ANOSIM test was used to test statistical 175 significance of global clusters. ANOSIM is a non-parametric rank-based permutation test which 176 compares within-cluster similarity to between cluster similarity (Clarke 1993). The ANOSIM *R* 177 statistic ranges from -1 to 1 and indicates the level of similarity between clusters, where 1 178 indicates all flounder groups within a cluster are more similar to each other than to groups from 179 other clusters, while an R of 0 indicates no difference between clusters. Significant differences 180 between clusters were also tested using pairwise ANOSIM tests. Global and pairwise ANOSIM

182 SIMPER analysis (Clarke 1993) was then used to determine which prey categories were most 183 different between clusters.

184 Non-metric multidimensional scaling (NMDS) ordination (Kruskal 1964) was performed 185 on the diet dissimilarity matrix to determine which prey categories were associated with flounder 186 group dissimilarity. NMDS was chosen over other ordination methods because predator group 187 diet data were not multivariate normally distributed, and relative differences between groups 188 were more important than absolute differences. Random starting placements (up to 100) were 189 used to find a convergent ordination solution, and an appropriate number of dimensions for the 190 ordination (*k*) was determined by examining stress values on a scree plots for 1-8 *k*-dimensions. 191 Statistical significance of the final ordination was tested using a random permutation test (999 192 permutations) and goodness-of-fit was evaluated based on the correlation between the observed 193 dissimilarity and ordination distance on a Shephard diagram. Prey loadings in ordination space 194 were calculated using linear correlation analysis, and statistical significance of prey category 195 loadings were tested using a random permutation test. Differences between flounder groups were 196 evaluated from visual inspection of NMDS plots. If NMDS plots indicated a consistent diet 197 difference between interspecific flounder pairs sharing size class and stratum, a two-tailed sign 198 test was conducted to test the null hypothesis of interspecific trophic equivalence (Zar 1999). 199 Under the null hypothesis, a flounder group would be equally likely to have a higher or lower 200 proportion of a prey type as its interspecific counterpart from the same size class and stratum. 218 and the most instantant which we adequate sample sizes were available for the sample sizes were available sizes were available (210 non-empty sizes were available sizes were available differences between groups used t

201 Proportional wet weight of prey provides a relative measure of prey contribution to diet, 202 but does not reflect absolute consumption of a prey type. To ensure that results of multivariate 203 analyses were not simply a reflection of interspecific biases in overall prey consumption, we 204 compared stomach fullness indices and prey frequency of occurrence (*%F*) among non-empty 205 stomachs to make interspecific comparisons of consumption. Fullness indices followed Lilly 206 (1991) except in lieu of cubed length  $(L^3)$ , we used estimated flounder biomass calculated from 207 length-weight regressions (*Mj)*, available from NOAA/AFSC's Grounndish Trophic Interactions 208 Database (Livingston et al. 2017). Total fullness index (*TFI*) was calculated as:  $TFI_j =$ 209  $\sum_{i=1}^{Q} W_{ij} M_j^{-1} \cdot 10^4$ , where  $W_{i,j}$  is the wet weight of prey type *i* in flounder stomach *j*,  $M_j$  is 210 estimated biomass of the flounder j, and *Q* is the number of prey categories. *PFI* was calculated 211 as:  $PFI_{i,j} = W_{i,j}M_j^{-1} \cdot 10^4$ . Prey types for *PFI* calculations were the same as for multivariate

213 significant interspecific differences between fullness indices for size class-stratum flounder pairs

214 using the two-tailed Mann-Whitney test. We used the non-parametric Mann-Whitney test

215 because fullness index distributions were non-normal and *PFI* had a high proportion of zeroes.

216

217 RESULTS

218 Prey types consumed by Arrowtooth Flounder and Kamchatka Flounder were 219 qualitatively similar and four hierarchical clusters were relevant for describing predator diets 220 (Fig. 2). Complete linkage clustering (Sørensen 1948) provided the best hierarchical clustering 221 performance (cophenetic correlation  $r^2 = 0.89$ ). Cluster W included mostly predators in 20-29 cm 236 Gadids, shrimp, zooplankton and benthic fishes contributed most  $(\geq 70\%)$  to between 237 cluster dissimilarity, but ranked contribution to differences varied among cluster pairs (Table 3). 238 A global ANOSIM test revealed significant differences across clusters  $(R = 0.96, P = 0.001)$ , and 239 five out of six pairwise ANOSIM tests identified significant and substantial diet differences 240 between clusters  $(0.91 \le R \le 1.00$ ; Table 3). Clusters Y and Z were not significantly different, 241 although an ecological difference was not convincingly rejected due to small cluster membership and a low p-value  $(P = 0.09)$ . 222 and 30-39 cm size classes, but also included Stratum C Arrowtooth Flounder ≤19 cm and 223 Stratum F Kamchatka Flounder  $\geq 50$  cm. Cluster X had the largest group membership, including 224 all large size classes (40-49 cm,  $\geq$  50 cm) except Cluster F Kamchatka Flounder  $\geq$  50 cm. Six 225 small and medium size classes were also in Cluster X. Cluster Y only contained the smallest size 226 class ( $\leq$ 19 cm), and included both species. Cluster Z contained only smaller size classes ( $\leq$ 19 cm, 227 20-29 cm) of Kamchatka Flounder. All stratum B size class pairs showed concordant cluster 228 membership, indicating greater interspecific diet overlap than in other strata for which at least 229 three size classes were represented. Among small and medium size classes  $(\leq 19 \text{ cm}, 20\text{-}29 \text{ cm},$ 230 30-39 cm), the only interspecific predator group pair sharing cluster membership, aside from 231 stratum B, were 30-39 cm size classes in stratum D. Overall, interspecific pairs of small and 232 medium size class predator groups were in different clusters 63.6% (7/11) of the time. Within 233 Cluster W, there was a considerable distance between a branch where seven out of eight leaves 234 were medium size class Arrowtooth Flounder, and a monospecific Kamchatka Flounder branch. 235 This difference occurred at a dissimilarity of 0.56, slightly below the phenon line at 0.6. 217 RESULTS<br>
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243 Important prey categories in the best NMDS ordination were consistent with the prey 244 categories that contributed most to hierarchical cluster dissimilarity (Table 3; Fig. 3). NMDS 245 diagnostics indicated a two-dimensional  $(k = 2)$  ordination was appropriate, and the ordination 246 resulted in a stress value of 0.09 (Fig. 4). More of the variation was explained by the ordination 247 than expected by random chance  $(P < 0.001)$  and fit to the data was good (non-metric  $r^2 = 0.99$ ). 248 Benthic fishes ( $P < 0.001$ ,  $r^2 = 0.37$ ), gadids ( $P < 0.001$ ,  $r^2 = 0.91$ ), shrimp ( $P < 0.001$ ,  $r^2 = 0.71$ ) 249 and zooplankton ( $P < 0.001$ ,  $r^2 = 0.50$ ) categories had highly significant vector loadings. Benthic 250 invertebrates ( $P = 0.01$ ,  $r^2 = 0.06$ ), unidentified fishes ( $P = 0.04$ ,  $r^2 = 0.02$ ), and other ( $P = 0.01$ , 251  $r^2 = 0.05$ ) prey categories had statistically significant vector loadings but, compared to the highly 252 significant prey vectors, did not account for much of the variation in ordination space (low  $r^2$ -253 values) so were omitted from the NMDS plot (Fig. 3). Miscellaneous pelagic fishes and squids 254 were not significant  $(P = 0.31, r^2 = 0)$ .

255 Shrimp and gadid vectors were divergent and were associated with an ontogenetic shift in 256 diet (Fig. 3). Smaller *Atheresthes* were more strongly associated with the shrimp vector, while 257 larger *Atheresthes* had a higher proportion of gadids in their diet. Zooplankton and benthic fish 258 prey categories were divergent and were associated with trophic niche separation between 259 Arrowtooth Flounder and Kamchatka Flounder. Among small and medium flounder size classes 260 (≤19 cm, 20-29 cm, 30-39 cm) there was a consistent within-stratum difference between 11 size 261 class pairs of Arrowtooth Flounder and Kamchatka Flounder. Arrowtooth flounder were more 262 strongly associated with zooplankton prey, and Kamchatka Flounder were more strongly 263 associated with benthic fish prey, suggesting interspecific trophic niche separation occurred 264 along a benthic fish-zooplankton prey gradient. The same benthic fish-zooplankton arrangement 265 also occurred in some larger size classes (40-49 cm in strata B, C, E and A;  $\geq$ 50 cm in strata A, B 266 and F), but not for 40-49 cm size classes in stratum D and E. Stratum F Kamchatka Flounder  $\geq 50$ 267 cm were dissimilar from other  $\geq$ 50 cm groups and were associated with the benthic fishes vector. 268 Arrowtooth Flounder groups consumed higher proportions of zooplankton than 269 Kamchatka Flounder in 100% of pairwise comparisons (sign-test; *P* < 0.001, *n* = 19). Stratum A 270 pairs were tied and omitted from sign-tests because neither group consumed zooplankton. 271 Kamchatka Flounder were more likely to have a higher proportion of benthic fishes in the diet 272 than Arrowtooth Flounder (sign test; *P* < 0.05, *n* = 21). Kamchatka Flounder groups had a higher 2747 than expected by readom chance  $(P < 0.001)$  and fit to the data was good (non-metric  $r^2 = 0.99$ ).<br>
248 Benthic fishes  $(P^2 < 0.001, r^2 = 0.37)$ , gadids  $(P < 0.001, r^2 = 0.91)$ , shrimp  $(P < 0.001, r^2 = 0.71)$  and zoopfundit

274 comparisons. There were no evident size or stratum trend for the 23.8% (5/21) of pairs where 275 Arrowtooth Flounder consumed more benthic fishes.

276 Interspecific differences in *PFIi* and *%F<sup>i</sup>* were generally consistent with patterns of 277 trophic niche separation observed in multivariate analyses. Significant differences (*P* < 0.05) in 278 *PFI<sup>i</sup>* were detected in 31 out of 154 flounder group comparisons (Fig. 4; Table S.1). Mean 279 *PFIzooplankton* and *%Fzooplankton* were higher for Arrowtooth Flounder than Kamchatka Flounder for 280 100% (19/19) of pairwise comparisons, and there were 11 statistically significant differences in 281 *PFIzooplankton*. Mean *PFI*benthic\_fishes was higher for Kamchatka Flounder in 76.2% (16/21) of group 282 comparisons, there were seven statistically significant differences in *PFI*<sub>benthic</sub> fishes, 283 and *%Fbenthic\_fishes* was higher in Kamchatka Flounder for 76.2% (16/21) of comparisons. 284 Significant differences in *PFI* were also identified for shrimp, Gadidae, benthic invertebrates, 285 miscellaneous pelagic, and unidentified fishes. Kamchatka Flounder ≤39 cm had higher mean 286 *PFIshrimp* and *%Fshrimp* in 90.9% (10/11) of group comparisons, among which five statistically 287 significant differences in *PFIshrimp* were observed. *PFIshrimp* was significantly higher for 40-49 288 cm Arrowtooth Flounder in stratum A, but shrimp were only a minor contributor to total 289 consumption. *PFIGadidae* was higher for Arrowtooth Flounder in one comparison, and higher for 290 Kamchatka Flounder in two comparisons. *PFIbenthic.inverts* was higher for 30-39 cm Kamchatka 291 Flounder in stratum E, but contributed very little to overall consumption. *PFImisc\_pelagic* was 292 significantly higher for ≥50 cm Kamchatka Flounder in stratum F. *PFI*<sub>fish\_unid</sub> was higher for 20-293 29 cm Arrowtooth Flounder in stratum E, and higher for  $\geq$ 50 cm Kamchatka Flounder in stratum 294 F. Despite the preponderance of significant differences in *PFI*, significant interspecific 295 differences in *TFI* only occurred in stratum B, for 30-39 cm, 40-49 cm and ≥50 cm size classes 296 (Table S.2). 274 FIF), were distanced in 31 out of 154 flounder group comparisons (Fig. 4; Table S.1). Mean<br>
279 FIF), were distanced in 31 out of 154 flounder group comparisons (Fig. 4; Table S.1). Mean<br>
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- 297
- 298 DISCUSSION

299 Our findings indicate trophic niche separation occurs between Arrowtooth Flounder and 300 Kamchatka Flounder at smaller sizes (≤39 cm) until trophic niches converge ontogenetically. 301 Trophic niche partitioning is consistent with hypothesized differences in foraging efficiency due 302 to gill raker functional morphology. Arrowtooth Flounder examined for this study consumed 303 more zooplankton than Kamchatka Flounder, and small-sized to medium-sized Kamchatka

305 Some diet overlap occurred between species of the same size class and the degree of 306 trophic niche separation between species varied among strata. Multiple factors likely contributed 307 to spatial variation in trophic niche separation. Diet overlap is common among allied resource 308 competitors because diet is jointly influenced by foraging capability and prey availability 309 (Amundsen et al. 2004; Chavarie et al. 2016). Greater diet overlap often occurs when a shared 310 prey resource is abundant and available. As resource competition increases, aggregate trophic 311 niche breadth can increase (Svanbäck and Persson 2004) and morphology-based resource 312 partitioning can become more pronounced (Svanbäck and Bolnick 2007). Hypothetically, this 313 would increase trophic niche separation between Arrowtooth Flounder and Kamchatka Flounder. 314 However, causes of spatial variation in niche separation cannot be determined because prey 315 abundances are not known. Thus, in stratum B, the comparatively high trophic niche overlap 316 could be due to either an abundance of preferred resources, or a scarcity of alternate resources. 317 A notable exception to the shrimp-gadid ontogenetic shift was evident for Stratum F 318 Kamchatka Flounder ≥50 cm, which may reflect a non-trophic dimension of ecological niche 319 separation between Arrowtooth Flounder and Kamchatka Flounder. With increasing depth, 320 Kamchatka Flounder abundance increases relative to Arrowtooth Flounder abundance on the 321 continental slope, and Arrowtooth Flounder become scarce on the outer continental slope (600- 322 1200 m; Zimmermann and Goddard 1996). Consequently, Stratum F Kamchatka Flounder ≥50 323 cm were sampled from deeper (mean: 474 m, SD: 72 m) than Stratum F Arrowtooth Flounder 324 ≥50 cm (mean: 405 m, SD: 98 m), so interspecific diet differences may reflect depth-dependent 325 changes in prey availability, including the lower *PFI<sub>Gadidae</sub>* for stratum F Kamchatka Flounder. 326 The primary aim of this analysis was to determine whether trophic niche separation occurs in 327 sympatry, so the outer continental slope was excluded from analysis. However, the Kamchatka 328 Flounder diet on the outer continental slope (600-1200 m) is distinct from strata included in this 329 analysis, and the stratum F diet may resemble a transitional diet between the inner and outer 330 continental slope. On the outer slope, the diet of Kamchatka Flounder  $\geq$  50 cm is dominated by 331 Continental stope, and Arrowtorth Foundate hand the streamed the present of the present of the breadth can increase (Svanbäck and Persson 2004) and morphoraritioning can become more pronounced (Svanbäck and Bolnick 21

332 Interspecific differences in fullness indices and prey frequency of occurrence were 333 consistent with our finding of trophic niche divergence linked to morphology, but also produced 334 two surprising outcomes which may warrant further investigation of interspecific ecological 335 differences. While the large number of comparisons we made increased the potential for Type I

336 errors, repeat differences in prey-specific *PFI* comparisons and highly significant differences in 337 *PFI* for important prey types are noteworthy. Higher *PFIshrimp* and *%Fshrimp* for Kamchatka 338 Flounder ≤39 cm may be due to greater benthivory by Kamchatka Flounder, although we were 339 unable to determine whether differences were due to consumption of greater numbers of the 340 same shrimp taxa, larger shrimp, a broader diversity of shrimp taxa, or a combination of factors. 341 Counting prey shrimp was not a standard protocol for laboratory analysis, the number of 342 available prey size measurements was not adequate to make interspecific comparisons of shrimp 343 size, and an insufficient proportion of shrimp were identified to a taxonomic level suitable for 344 more detailed prey categories. Dominant EBS shrimp taxa include a combination of benthic 345 (families Crangonidae, Hippolytidae), pelagic (family Pasiphaeidae, Sergestidae), and semi-346 pelagic (family Pandalidae) taxa (Wicksten 2012), and both flounders consume shrimp from all 347 major EBS families (Yang and Livingston 1986; Yang 1995; Orlov and Moukhametov 2004). 348 For larger-sized flounders (40-49 cm,  $\geq$ 50 cm) in Stratum B, interspecific differences in 349 *PFIGadidae* were explained by higher *%FGadidae* in non-empty Kamchatka Flounder stomachs, but 350 we were unable to determine why the difference in  $\%F_{\text{Gadiidae}}$  occurred. Potential explanations 351 include interspecific differences in allometric scaling of size-structured interactions, feeding 352 activity level, aspects of the prey search-detection-capture sequence, rates of digestion, feeding 353 chronology, or spatial overlap with prey on a finer spatial scale than we considered in our 340 same shrimp taxa, la<br>
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355 Environmental variation influences community spatial structure in the EBS across 356 seasonal and inter-annual temporal scales (e.g. Mueter and Litzow 2008; Kotwicki and Lauth 357 2013; Barbeaux 2017). During summer months, the spatial distribution of many commercially 358 and ecologically important species is influenced by the cold pool, an area with near bottom water 359 temperatures  $\leq 2^{\circ}$ C which approximately reflects the southern extent of winter sea ice and 360 persists through summer months (Mueter and Litzow 2008). During some years, this limits 361 distributions of Arrowtooth Flounder and Kamchatka Flounder to the outer continental shelf, 362 because they avoid the cold pool (Spencer 2008; Barbeaux and Hollowed 2018). On the EBS 363 shelf (but not the slope), Kamchatka Flounder ≥40 cm tend to occupy slightly colder habitat than 364 Arrowtooth Flounder ≥42 cm (Barbeaux 2017; Barbeaux and Hollowed 2018). Hypothetically, 365 during cold years, this may allow comparatively higher overlap of large Kamchatka Flounder 366 with cold-tolerant Gadid prey (Walleye Pollock), than for large Arrowtooth Flounder. While diet

367 data in the present study included both cold (2007-2013) and warm (2014-2016) years,

368 Kamchatka Flounder sample sizes were not sufficient to compare concurrent interspecific shifts 369 in diet over time.

370 It is unclear how Arrowtooth Flounder diet shifts relative to Kamchatka Flounder diet as 371 the environment and prey availability change. Discrete sampling of Arrowtooth Flounder and 372 Kamchatka Flounder would improve understanding of how environmental variability influences 373 diet composition. In the Gulf of Alaska, zooplankton contribution to Arrowtooth Flounder diet is 374 highest during spring, when zooplankton are most abundant (Knoth and Foy 2008). No 375 analogous seasonal comparison of Kamchatka Flounder diet has been conducted. Stable isotope 376 analysis would extend the temporal range of summer sampling, improve understanding of 377 benthic and pelagic trophic pathway duration, and provide a relative measure of trophic position 378 between species. In the Gulf of Alaska, Arrowtooth Flounder trophic position, inferred from bulk  $379$   $\partial^{15}$  N ratios was lower during years when euphausiids were unusually abundant in stomach 380 samples (Marsh et al. 2015).

381 Support for the hypothesized association between morphology and diet is ample. 382 Sympatric benthivore and planktivore polymorphs have been identified in gasterosteids (McPhail 383 1984; Taylor and McPhail 2000), osmerids (Taylor and Bentzen 1993), salmonids (Foote et al. 384 1999; Guiguer et al. 2002; Chavarie et al. 2016), and coregonids (Amundsen et al. 2004; Østbye 385 et al. 2005; Gowell et al. 2012). Gill raker functional morphology has repeatedly been linked to 386 adaptive resource partitioning between benthivore and zooplanktivore polymorphs in freshwater 387 ecosystems (Schluter and McPhail 1992; Wimberger 1994; Smith and Skúalson 1996). In marine 388 systems, gill raker functional morphology has been linked to prey size divergence between 389 sibling planktivores (e.g. Castillo-Rivera et al. 1996), but had not previously been linked to 390 benthivory-zooplanktivory resource partitioning. Regardless of prey type consumed, foraging by 391 both species of *Atheresthes* likely occurs near-bottom, in contrast with common patterns of 392 habitat partitioning between demersal foraging benthivore morphs and pelagic foraging 393 zooplanktivore morphs in freshwater systems. 397 due environment and prey availability change. Discrete sampling of Arrowtooth Flounder and<br>372 Kamchatka-Floonder would improve understanting of how environmental variability influences<br>373 disc composition: In the Gu

394 A morphological mechanism for trophic niche separation implies niche specialization 395 should persist across the ranges of Arrowtooth Flounder and Kamchatka Flounder. Throughout 396 the range of Arrowtooth Flounder, euphausiids (zooplankton) are frequent prey and often

398 1969; Rose 1980; Yang and Livingston 1986; Yang 1995; Buckley et al. 1999; Knoth and Foy

399 2008). By contrast, euphausiids are uncommon or unimportant in Kamchatka Flounder diet

400 (Yang and Livingston 1986; Orlov 1997; Orlov and Moukhametov 2004). While ecosystem

401 dynamics and prey availability certainly vary across the North Pacific, diet differences

402 throughout the species' ranges support the association between gill raker morphology and trophic 403 niche separation.

404 Unlike gill rakers, the morphological difference in left-eye position between the two 405 species does not have a clear association with the observed pattern in benthivory-zooplanktivory 406 resource partitioning. The position of the left eye of Arrowtooth Flounder (more 407 zooplanktivorous), intersecting the dorsal margin (Yang 1988), suggests a field of view that may 408 encompass more of the benthos than that of Kamchatka Flounder (more benthivorous). The eye 409 position in flatfish is less indicative of specific feeding behaviors than other morphological 410 characteristics and specializations (Gibb 1997; Bergstrom and Palmer 2007), so this may be due 411 to random phenotypic divergence. However, we speculate on the possibility that the marginal 412 position of the left eye in Arrowtooth Flounder may result from parasite mediated selection by 413 the pathologic copepod, *Phrixocephalus cincinnatus* (Wilson, 1908). A high percentage of 414 Arrowtooth Flounder off British Columbia, Canada, become infected with this parasite (Kabata 415 1969; Blaylock et al. 2005). The right eye was more commonly infected than the left eye (the eye 416 nearer the dorsal margin) in Arrowtooth Flounder (Kabata 1969; Blaylock et al. 2005), but 417 infection rates of Pacific Sanddab *Citharichthys sordidus*, having eyes located nearly level on the 418 eyed-side of the head, have similar infection rates of the left and right eyes (Perkins and Gartman 419 1997). Binocular infections by this copepod certainly lead to death (Kabata 1969; Perkins and 420 Gartman 1997) while monocular infections may also result in substantial impairment of the host 421 (Blaylock et al. 2005). However, a captured Pacific Sanddab, was observed to survive the 422 monocular infection and completion of the life cycle (death) of the parasite, and it grew as fast as 423 uninfected Pacific Sanddabs also in captivity (Perkins and Gartman 1997). There are no reports 424 in the literature of a similar pathology for Kamchatka Flounder in the western North Pacific 425 Ocean. If marginal eye position imparts increased host-survival of infections by pathologic 426 copepods at minimal or no associated energetic cost, the trait would be beneficial to a population 427 that suffers high rates of infection and could be maintained in the absence of the parasite (Ebert 428 2005). But the speci<br>paration.<br>Jnlike gill radios not have<br>partitioning<br>ktivorous), in<br>ass more of t<br>in flatfish is<br>ristics and sp<br>m phenotypi<br>of the left e:<br>ologic coper<br>ooth Flounde<br>laylock et al<br>le dorsal ma<br>n rates of Pa<br>le o

429 Evolutionary mechanisms which initially led to trophic niche separation between 430 Arrowtooth Flounder and Kamchatka Flounder are unclear because the evolutionary history of 431 the genus *Atheresthes* is not known. Sympatric speciation caused by resource-driven character 432 displacement may be possible, although an allopatric origin is more likely considering the glacial 433 history of the North Pacific. Glacial isolation during the Pleistocene epoch has been suggested as 434 the reason for east-west genetic population structures in Pacific Cod (Canino et al. 2010) and 435 Pacific Herring (Liu et al. 2011) in the North Pacific. There is a clear genetic difference between 436 Arrowtooth Flounder and Kamchatka Flounder and no evidence of interspecific hybridization 437 (Ranck et al. 1986; De Forest et al. 2014). However, the genetic difference between species is 438 minor, suggesting recent speciation (Ranck et al. 1986). Although genetic studies have been 439 conducted to validate species identities in *Atheresthes*, more thorough investigation of genetic 440 population structure is necessary to elucidate patterns of phylogenetic divergence.

441 Ecological niche width imposes a constraint on population growth (Hutchinson 1957; 442 Schoener 1974), so a broader combined trophic niche for smaller Arrowtooth Flounder and 443 Kamchatka Flounder size classes may facilitate a higher carrying capacity than might be 444 expected under trophic niche equivalence. An important consequence of higher *Atheresthes*  445 carrying capacity is elevated predation pressure, which may dampen recruitment of 446 commercially important Walleye Pollock in the EBS (Mueter et al. 2011; Holsman et al. 2015; 447 Spencer et al. 2016). Predation on Walleye Pollock may be especially impactful during warm 448 years, when higher bottom temperatures allow greater Arrowtooth Flounder and Kamchatka 449 Flounder spatial overlap with juvenile Walleye Pollock (Mueter et al. 2006; Kotwicki and Lauth 450 2013; Spencer et al. 2016; Barbeaux 2017). Warm year frequency is expected to increase due to 451 climate change, and the potential corollary of increased predation on Walleye Pollock is a 452 management concern, so modeling efforts have sought to forecast climate-mediated shifts in 453 Walleye Pollock predation mortality (Mueter et al. 2011; Holsman et al. 2015; Spencer et al. 454 2016). Separately considering demographic changes and diet compositions for Arrowtooth 455 Flounder and Kamchatka Flounder could improve ecosystem (energy flow) models. While 456 abundances of sympatric Arrowtooth Flounder and Kamchatka Flounder increased concurrently 457 in recent decades, ecological niche partitioning, including trophic niche divergence, suggests <sup>443</sup><br>433 bistory of the North Pacific. Glacial isolation during the Pleistocen<br>the reason for east-west genetic population structures in Pacific Co<br><sup>433</sup> history of the North Pacific. Glacial isolation during the Pleistoc

459 Our study contributes to understanding of interspecific differences in the life history and 460 ecology of Arrowtooth Flounder and Kamchatka Flounder in their sympatric range. Although 461 larval traits and spatial-temporal dispersal patterns overlap (De Forest et al. 2014) and both 462 species exhibit a shift towards deeper water with increasing size and age (Zimmermann and 463 Goddard 1996), Arrowtooth Flounder grow faster and mature at a smaller size and younger age 464 than Kamchatka Flounder (Stark 2012). In addition, large Kamchatka Flounder tend to be 465 distributed at greater depths and in warmer oceanic waters than large Arrowtooth Flounder 466 (Zimmermann and Goddard 1996). Through an accumulation of dietary data (Livingston et al. 467 2017), we find evidence against trophic equivalence between species, and identify an 468 interspecific difference in trophic niche consistent with hypothesized variation in foraging 469 efficiency due to divergent gill raker morphology. 489 Garbeaux, S. J., Barbeaux, S. J., Barbeaux, S. J., and A. B. Hollowed. 2018. Concerned a started a strained a star of the Camerana and The Camerana and Warrell and The Commentation and Concerned Manuscript (Stark 2012

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### 708 TABLE CAPTIONS

709 Table 1. Sample sizes stomachs sampled per year, by species and survey. Numerator indicates 710 the number of non-empty stomachs, denominator indicates total number of stomachs, number in 711 parentheses indicates percentage of non-empty stomachs. Dashes (-) indicate no bottom-trawl 712 survey was conducted.

713

714 Table 2. Sample sizes of stomachs for flounder groups, assigned by species, spatial stratum, and 715 size class. Numerator indicates the number of non-empty stomachs, denominator indicates total 716 number of stomachs, number in parentheses indicates percentage of non-empty stomachs.

717 Italicized groups, with sample sizes <10, were excluded from analysis.

718

719 Table 3. Pairwise ANOSIM R statistics (lower diagonal) for the four relevant predator clusters

720 (W, X, Y, Z) identified using complete linkage clustering. Significance levels denoted by:  $P <$ 

721 0.1 (^), *P* < 0.05 (\*), *P* < 0.01 (\*\*), *P* < 0.001 (\*\*\*). SIMPER results (upper diagonal) indicate

722 which prey categories cumulatively contribute  $\geq$  70% of between-cluster dissimilarity, ranked by

723 contribution. Prey categories are: BF – Benthic fish, GA – Gadidae, SH – Shrimp, ZP –

- 724 Zooplankton.
- 725
- 726 TABLES
- 727 Table 1.



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729 Table 2.



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