1 Extracting socio-spatial networks from photo-ID data using multilevel

2 multinomial models

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22 Abstract:

23 Photo identification of individuals within a population is a common data source that is 24 becoming more common given technological advances and the use of computer vision and 25 machine learning to re-identify individuals. These data are collected through hand-held 26 cameras, drones, and camera traps, and often come with biases in terms of sampling effort 27 and distribution. In spite of these biases, a common goal of collecting these datasets is to 28 better understand the habitat use pattern of individuals and populations. Here, we examine the 29 potential for multilevel multinomial models to generate socio-spatial networks that capture 30 the similarities in individual users across the spatial distribution of a species. We use this 31 approach with 18 years of photo-ID data to better understand population structuring of beluga 32 whales in the St. Lawrence River. We show using permuted and simulated data that this 33 approach can identify community network structures within populations in a way that 34 accounts for biases in collections methods. Applying this method to the entire 18 years 35 dataset for SLE beluga, we found three spatially distinct communities. These results suggest 36 that within the population's summer range individuals are moving within restricted areas (i.e., 37 home ranges), and have implications for the estimated impacts of localized anthropogenic 38 stressors, such as chemical pollution or acoustic disturbances on animal populations. We 39 conclude that multilevel multinomial models can be effective at estimating socio-spatial 40 networks that describe community structuring within wildlife populations. 41

42 Keywords: Multinomial Model, Beluga, Photo ID, Socio-Spatial Network, Bayesian
43 Network, Community Detection

44 **1. Introduction**

45	An understanding of the spatial and temporal distribution of a species of concern is of
46	central importance to conservation and management (Evans & Hammond, 2004).
47	Increasingly, photo and video are being used to monitor individuals within populations
48	(hereafter photo-ID data), providing a view of within population social mixing and habitat
49	use (Koivuniemi et al., 2016). The increased use of machine learning to identify individuals
50	from these data streams has greatly facilitated the use of these photo-ID data (Schneider et
51	al., 2019). These individual identifications have facilitated the use of novel statistical and
52	computational methods to quantify within population structures, such as social network
53	analysis (Perryman et al., 2019; Schilds et al., 2019).
54	It is often the case, however, that efforts when collecting photo-ID are not evenly
55	distributed. This differentiation in effort can heavily bias estimates of both habitat usage and
56	population distribution estimates (Hupman et al., 2018). Here we propose the novel use of
57	multilevel multinomial models to account for these biases and to estimate socio-spatial
58	structures within populations.
59	The existence of social structuring within populations, such as communities, can have
60	important ecological and management implications. If a population as a whole can be
61	considered as highly mixed, i.e., with individuals showing no strong patterns of home range
62	use or sub-structuring within the large population, then all individuals are equally likely to
63	feel the impacts of local changes in the environment. In contrast, if the population cannot be
64	considered to be highly mixed, and shows strong sub-structuring and site-fidelity patterns
65	within the larger population, local stressors might have a disproportionate impact on
66	subsections of the population. For example, if noise pollution increased in only one sector, in
67	a highly mixed population all individuals would be lightly impacted, but in a structured
68	population a subset of the population would be highly impacted. These differences in spatial

69 structuring of populations can lead to biased estimation of the likelihood and magnitude of 70 impacts from local stressors both at the individual and population levels (DeFur et al., 2007). 71 The multilevel multinomial modeling approach does not have a large body of 72 literature to draw on for use with photo-ID data. However, it does have some unique 73 advantages (Koster & McElreath, 2017). For instance, if sampling effort is biased in different 74 regions, the mean probability of being seen within highly sampled regions will be biased 75 upwards. By taking advantage, however, of the multilevel structure of the model it is possible 76 to extract individual deviations in the probability of being seen within a particular region. 77 Decisively, these individual level deviations from the mean probability are not biased by 78 changes in the sampling effort. That is the mean probability will increase with sampling 79 effort, but the relative difference between individuals within the sector will not. High users of 80 a particular region of a habitat will consistently be higher compared to low users of that 81 habitat, and this difference between high and low users will not be biased by sampling effort. 82 Furthermore, by comparing the individual differences between regions it is possible to see if 83 the high/low users of one region are similarly the high/low users of another region. The 84 similarity, or dissimilarity, between regions can then provide information about which 85 regions share similar user profiles. We suggest that by using the correlations between these 86 individual level deviations in high/low users between regions it is possible to generate socio-87 spatial networks and identify social structuring within the population. In particular it can help 88 to identify spatial communities, i.e., a set of regions that share similar usage patterns and that 89 differ from other regions.

To evaluate the use of multilevel multinomial models to identify socio-spatial
structuring within a population, we make use of a long term photo-ID dataset of beluga
whales in the St. Lawrence Estuary, Canada. This population has undergone a drastic
decrease from around 10,000 in the late 1800s to less than 1,000 today, and is currently

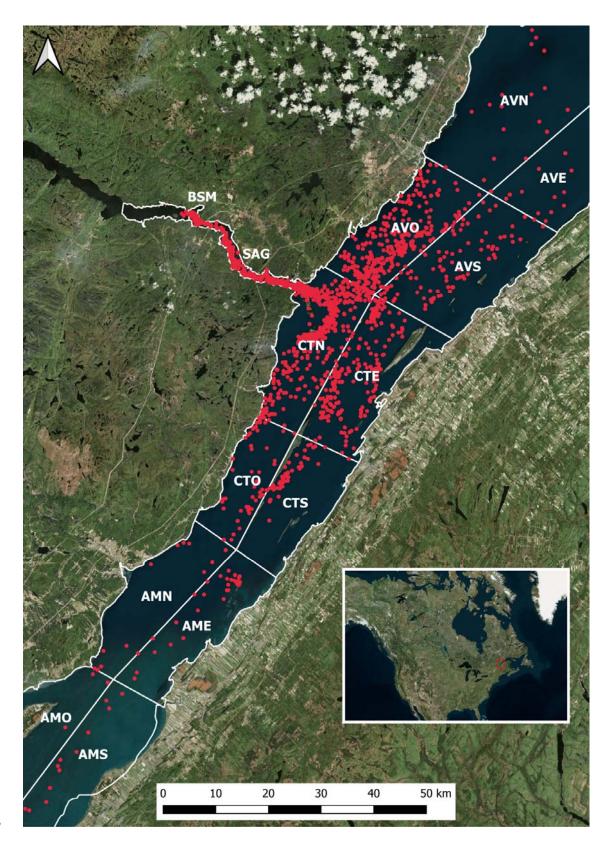
94	considered as endangered in Canada according to the Species at Risk Act (COSEWIC 2014;
95	Fisheries and Oceans Canada, 2012; Mosnier et al., 2015). The population is part of a larger
96	study on the mitigation of noise pollution due to marine traffic (Chion et al., 2017; Lesage et
97	al., 2014; McQuinn et al., 2011; Parrott et al., 2011).
98	In this paper we first evaluate the performance of the multilevel multinomial models
99	using simulated data, testing if the method correctly estimates no community structuring
00	when none is present, and identifies the correct structure when it is present. In both cases we
101	use the 18 year beluga photo-ID dataset, randomly permuting uniquely identified individuals
102	to generate unstructured datasets, and randomly placing individuals within pre-specified
103	communities to generate structured datasets. We then apply the method to the observed data
104	and quantify community structures within the population's summer range in the St. Lawrence
105	Estuary. Finally, we discuss some potential extensions to the multilevel multinomial
106	modeling approach.
107	

1082. Material and Methods

109 2.1 Data

110 Photo-ID data were collected using a handheld camera onboard a boat that was able to 111 navigate near to beluga, hereafter referred to as an encounter. Once near beluga a photo ID 112 protocol was then followed to generate photographs used to attempt to identify individuals. 113 Due to the logistical difficulty of covering a large body of water, the sampling effort was 114 unequally distributed across 14 sectors within the St. Lawrence Estuary (Fig. 1). The photo-115 ID dataset used in this study was collected from 1989-2007 and are part of an ongoing 116 project. The data is stored in a database that facilitates the identification and association 117 between photos to help track the individual identification process. This resulted in a dataset 118 of 7,525 individual encounters where the individual was successfully identified by photo

- sampling (hereafter referred to as a photo-ID), and where a GPS point was taken and the
- 120 sector recorded. This resulted in 821 unique individuals being successfully identified, with a
- 121 mean number of photo-IDs per individual of 9 (min = 1, max = 90) (Fig. 1).



124 Figure 1: Spatial distribution of photo-ID data (red points) within the St. Lawrence Estuary,

- 125 Quebec, Canada (red square in the inset map). The 14 sectors are outlined and labeled in
- 126 white, and covers the summer habitat of this beluga population.
- 127

128 2.2 Statistical Analysis

129 Our aim was to use photo-ID data to estimate the probability of seeing an individual 130 in each delineated sector of the St. Lawrence Estuary, and to use these individual 131 probabilities to estimate socio-spatial structures within the beluga population (Fig. 1). To 132 accomplish this aim we used a multilevel multinomial model, where the dependent variable 133 was the number of times an individual was captured photographically (i.e., photo-identified) 134 in each sector. The use of a multilevel model structure allows for the estimation of both the 135 mean probability of photo-identifying an individual in each sector, and the individual level 136 differences in this probability by using individual ID as a random intercept. If we take, as an 137 example, a case where there is only two sectors, then the log-odds of finding individual *i* in a 138 sector other than the reference sector can be modeled using a multilevel multinomial 139 following (Koster & McElreath, 2017) as:

$$log\left(\frac{p_{1,i}}{p_{r,i}}\right) = \mu_1 + v_{1i}$$
$$log\left(\frac{p_{2,i}}{p_{r,i}}\right) = \mu_2 + v_{2i}$$

140 Where $p_{1,i}$ is the probability of seeing beluga *i* in sector 1, whereas $p_{r,i}$ is the 141 probability of seeing beluga *i* in the reference sector. The μ_1 and μ_2 are the intercepts, i.e., the 142 mean probability of seeing a beluga in sectors 1 and 2. This mean probability represents 143 preference/avoidance of the specified sector, however, it is very likely to be biased due to 144 variation in sampling effort. Finally, the v_{1i} and v_{2i} are the estimated individual differences 145 (i.e., random intercepts) from the mean probability of capture in sectors 1 and 2, respectively.

146 These individual differences from the mean probability capture if an individual beluga is a 147 high/low user of that sector, and is not biased by variation in sampling. It is then possible to 148 model the covariance of the individual differences between two sectors using a multivariate 149 normal distribution:

$$\begin{array}{l} \boldsymbol{v}_{1i}\\ \boldsymbol{v}_{2i} \end{array} \sim Multi \ Normal(0, \boldsymbol{\Omega}_{v}) \\ \\ \boldsymbol{\Omega}_{v} = \begin{bmatrix} \sigma_{1,1} & \sigma_{1,2}\\ \sigma_{2,1} & \sigma_{2,2} \end{bmatrix} \end{array}$$

150 This multivariate normal distribution has a mean of 0 and a covariance matrix Ω_{ν} . 151 Here the diagonal entries in the covariance matrix (e.g., $\sigma_{1,1}$) represent the magnitude of 152 individual differences within a sector, i.e., are their high and low users in a sector or are all individuals equally likely to be seen? The off-diagonal entries (e.g., $\sigma_{2,1}$) are the covariance 153 154 estimates between sectors, i.e., do sectors share similar high and low users? By estimating the 155 correlation of individual differences between sectors, this multilevel modeling approach 156 quantifies how much information individual differences in one sector can provide about 157 another sector. Positive correlations suggest that the high/low users in one sector are similarly 158 high/low users in another sector, while negative correlations suggest high/low users in one 159 sector are the low/high users in another sector. 160 This model can be fit using brms in the R environment using a multivariate syntax: 161 $bf(y | trials(n) \sim 1 + (1|q|ID)) + multinomial()$. Here, y is a set of column vectors where each 162 column is a sector and each row is an individual. The values in this column vector indicate 163 how many times each individual was seen in each sector. The n is the total number of times 164 an individual was captured, and q is an arbitrary character choice that allows correlations 165 between the estimates of random intercepts for each sector (Bürkner, 2017). 166

167 2.2.1 Addition of a common reference sector

168	Tests using simulated data suggested that adding a preset reference category (i.e., a
169	fifteenth sector used as a reference sector) to the data was required to estimate the correlation
170	between sectors (Fig. S1). To create this reference category, we tally up the observations for
171	each individual and add that number of observations to the new reference sector. This
172	essentially sets the probability of a capture in the reference sector to 0.5 for all individuals,
173	i.e., equal to the probability of being captured outside of this reference sector. This ensures
174	that all individuals have the same baseline probability in the reference category, and as the
175	parameters in the multinomial model measure deviations away from the reference sector, we
176	gain better estimates of the relative deviations between individuals (Fig. S1).
177	

178 2.2.2 Dealing with biases in photo-ID datasets

179 This multilevel multinomial approach accounts for repeated sampling of individuals, 180 and provides an estimate of whether some individuals are found more or less often than the 181 mean probability of captures in each sector. We are particularly interested in the estimates of 182 individual differences from the mean probability of capture (i.e., the random intercepts) as 183 these estimates are not impacted by bias in sampling effort among sectors. This is not the 184 case for estimates of the mean probability of capture for each sector, which are expected to 185 increase in highly sampled sectors. For example, the Saguenay River is over-sampled 186 compared to the other sectors (SAG in Fig. 1), increasing the mean probability of capturing 187 individuals in that sector. However, over-sampling should not affect the individual 188 differences in the probability of being captured, i.e., all individuals' chances of being captured 189 go up or down equally.

Similarly, potential biases due to ease of recognition, e.g., some beluga or age classes
might have more distinctive markings, are minimized using a multilevel multinomial
approach where the differences in the probability of being seen between sectors is the main

193 focus. For example, if juveniles are less likely to be successfully identified by photo, then 194 they might have a reduced number of photo-IDs compared to other age classes, but the 195 difference in distribution of these fewer photo-IDs across sectors will not be impacted. For 196 example, if both an adult and juvenile spend twice as much time in the SAG sector compared 197 to all other sectors, you might expect a photo-ID distribution (Sag:not-sag) of 10:5 and 2:1, 198 respectively. In both cases the probability of being captured in the SAG sector is twice that of 199 the remaining sector. Due to the adaptive partial pooling properties of multilevel models, 200 individuals with few photo-IDs will, however, be less likely to show differences to the mean 201 probability, i.e., they contain less information. This means that if an age class has very little 202 chance of being identified by photo-ID, they will likely contribute less to the estimated socio-203 spatial structures estimated by the multilevel multinomial approach.

204 By using a multilevel modeling approach we also reduce the chance of false positives 205 when making comparisons between many different individuals in many different sectors (i.e., 206 problem of multiple comparisons). For example, if we were to estimate the differences in the 207 probability of each sector separately for each individual, the risk of false positives would be 208 increased. By using a multilevel approach to estimating the differences we can make effective 209 use of partial pooling of information to reduce extreme values, especially where the number 210 of photo IDs is not equal between individuals. Furthermore, by running this in a Bayesian 211 framework we are able to place priors on the individual differences within sectors that start 212 the model assuming that there are no differences between individuals in their use of each 213 sector, e.g., student_t(3,0,1).

214

215 2.3 Social Network Analysis

Social networks are often used when visualizing and quantifying social structureswithin populations, with individuals often represented as nodes and their interactions as edges

218 between these nodes. In our case, we use sectors as nodes, and the similarities in user profiles 219 between sectors as edges. The correlations between sectors estimated from the multilevel 220 multinomial model can be used to create a network where the posterior predictions of each 221 correlation parameter corresponds to an edge weight in the network. In this way, each edge 222 has a posterior distribution and can be used to create many networks from which a 223 distribution of network metrics can be generated, e.g., the distribution of node strength values 224 can be calculated for each sector. The advantage of having distributions of network measures 225 is that the measures can be readily compared, e.g., does one sector have a higher node 226 strength than another? It is also possible to use the distribution of edge weights, and a chosen 227 threshold (e.g., 95% credible interval), to highlight only the edges where the sign of the 228 correlation is known with a particular range of certainty. In this paper, we used this latter 229 approach to generate a signed network (i.e., a network with positive and negative edges) and 230 use a simple signed-edge rule to define communities: where a distinct community is a set of 231 nodes that share positive edges but no negative edges. We also made use of signed 232 blockmodeling, an algorithm that can also be used to identify blocks of nodes that maximize 233 within block positive edges and minimize within block negative edges (Doreian & Mrvar, 234 2015). While the signed-edge rule generally provides relatively intuitive results with simple 235 networks, using the signed blockmodeling is likely to be particularly advantageous when 236 dealing with large networks.

237

238 2.4 Testing data

To assess the accuracy of the multilevel multinomial modelling approach, we generated test datasets from the observed photo-ID data. We ensured that the test datasets contained the same number of unique individuals, distribution of sightings (i.e., some individuals are seen more than others), and overall number of photo-IDs as the observed dataset. We, however,

243	varied the spatial mixing of the test datasets. To test if the proposed method correctly
244	detected no pattern when none existed, we created a completely random test dataset by
245	permuting the sector associated with each photo-ID in the observed dataset. The expected
246	result was to find no correlations between sectors given that the sectors for each photo-ID had
247	been randomly permuted. To then test whether the proposed method could also correctly
248	identify patterns when a known pattern existed, we generated a structured test dataset by
249	randomly assigning each individual to four equally populated communities with the
250	following and hypothetical home range of adjacent sectors: community 1-BSM, SAG, CTN,
251	community 2- CTN, CTO, AMN, community 3- AVO, AVS, AVN, and community 4-AME,
252	CTS, CTE. Following this, we altered the sector of where the individual photo-IDs were
253	taken so as to fall within sectors associated with an individual's community, i.e., one of their
254	home range sectors. We did this by choosing a sector for each photo-ID based on the
255	individual's assigned community 80% of the time; a random sector was chosen for the other
256	20% of the time, introducing noise in the assignment of sectors. We then tested whether the
257	model correctly identified the correlations between sectors that defined the home range of
258	each of the communities.
259	
260	3. Results
261	3.1 Testing data
262	When the multinomial multilevel model was fit to the data with sectors randomly

263 permuted between all photo-IDs, the model found no evidence for positive/negative

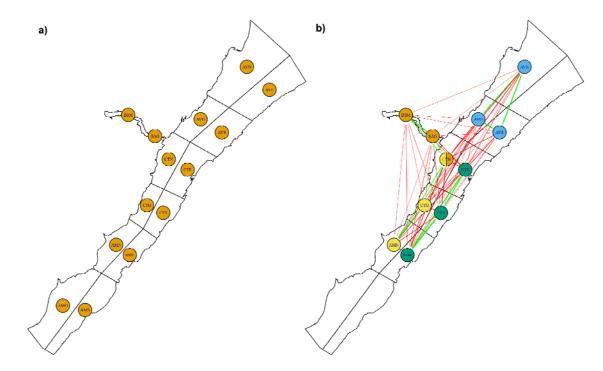
264 correlations between sectors (Fig. 2a). Similarly, when we simulated data with some known

structure, i.e., when we artificially created spatially distinct communities, we found that the

266 model accurately estimated the correlations between sectors that defined these artificial

communities (Fig. 2b). The simple signed-edge rule and blockmodelling algorithm applied to

- the simulated datasets both revealed the four artificially generated communities, though the
- blockmodeling algorithm had difficulty with the multi-membership node as it could not
- assign a node to two blocks (i.e., the CTN node that was shared between communities 1 and
- 271 2).
- 272





274 Figure 2: Similarity and dissimilarity between sectors in the simulated datasets: a) randomly 275 permuted data, where there are no spatial communities, and b) structured data, where there 276 are four distinct communities. In b) the simulated communities are represented by color codes 277 for each of their sectors (Note: CTN is part of the orange and yellow communities). The 278 green edges between two sectors signify that the sectors share high/low users, while red 279 edges signify that they have dissimilar high/low users. The lack of an edge signifies that the 280 high/low users of one sector does not provide information about the high/low users of other 281 sectors. Nodes represent sectors, and are coloured based on the communities imposed when 282 simulating the data.

284 *3.2 Observed data*

285	The results from our multilevel multinomial model found that within sectors there
286	were consistent individual differences in how likely it was to see individual beluga (Table 1).
287	That is, within sectors, there were some beluga that used the sector heavily, while others did
288	not. The model also found that between sectors these individual differences were correlated
289	(Table S1). These correlations quantify the magnitude of similarity/dissimilarity between
290	sectors in terms of which beluga are using those sectors heavily or rarely. If we take two
291	sectors as examples, e.g., the SAG and CTE sectors, representing, respectively, a tributary to
292	the St. Lawrence Estuary and a sector on the opposite side close to the South shore of the
293	Estuary, and we look at the top 10 estimated high users (i.e., relatively high probability of
294	being found there) within the SAG, we find that they are found to be low users in the CTE
295	sector (see blue dots in Fig 3 a) and b)).

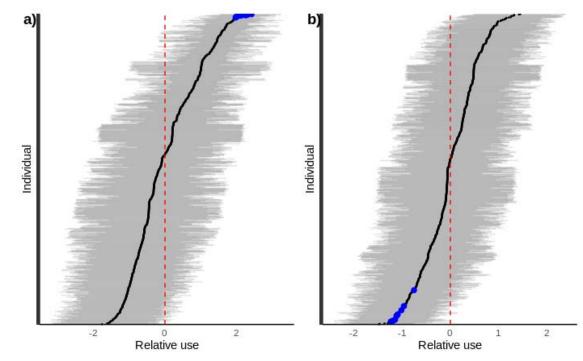




Figure 3: Estimate of the relative use (i.e., deviation from mean use) for each individual
within the SAG (a) and CTE sectors (b) of the St. Lawrence beluga summer habitat. The
values are deviations from the mean probability of observing individuals within a sector and

are on the logit scale. The red dashed line represents the mean use, black points represent the estimated deviation from the mean, while the horizontal grey lines represent the 95% credible interval. To highlight how correlations are estimated between sectors, the estimated top 10 users of the SAG sector are represented by blue dots (panel a), and those same individuals are also highlighted in blue in the CTE sector (panel b).

305

306 The use of a multilevel model also allowed us to estimate the magnitude of individual

307 differences in each sector, i.e., the extent to which there are high/low beluga users in a sector.

308 Our model found that the CTN sector showed very little individual differences in use (Table

309 1, i.e., low "sd" value) compared to other sectors, suggesting very little differences in high

and low users of that sector. While the BSM sector showed large individual differences, with

311 some very high/low users of that sector (Table 1).

312

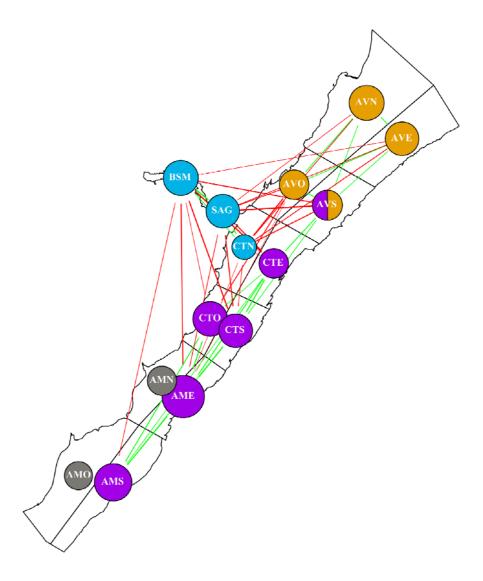
313 Table 1: Parameter estimates from the multilevel multinomial model predicting the 314 probability of capturing a photo-ID by sector. Estimated magnitudes of individual differences 315 (sd) are presented for each sector. Higher 'sd' estimates indicate more individual differences 316 in individual use of that sector, whereas lower estimates indicate individuals are using the 317 sector at very similar levels. To facilitate interpretation we have ordered the table by lowest 318 to highest estimates of individual differences. As the number of parameters in the model is 319 large, the overall mean by sector, and estimated correlations between individual differences, 320 are presented in the supplementary section (Table S1).

Parameter	Estimate	SD	l-95% CI	u-95% CI
sd(mu_CTN)	0.33	0.03	0.27	0.39
sd(mu_AMO)	0.65	0.38	0.05	1.41
sd(mu_AMN)	0.76	0.29	0.14	1.31

sd(mu_AVO)	0.78	0.05	0.68	0.88
sd(mu_CTE)	0.78	0.05	0.68	0.89
sd(mu_AVS)	0.83	0.07	0.70	0.97
sd(mu_AVE)	1.16	0.20	0.78	1.57
sd(muy_SAG)	1.16	0.08	1.01	1.31
sd(mu_CTS)	1.18	0.11	0.97	1.40
sd(mu_CTO)	1.27	0.14	1.00	1.55
sd(mu_AVN)	1.30	0.17	0.98	1.65
sd(mu_BSM)	1.30	0.11	1.09	1.52
sd(mu_AMS)	1.53	0.22	1.10	1.98
sd(mu_AME)	2.00	0.19	1.64	2.38

³²¹

322 Using the between sector correlations to generate a signed network overlaid on top of 323 the sectors in the St. Lawrence, suggests, for example, that individuals that are seen in the 324 SAG sector a lot, are also seen in the BSM and CTN sectors a lot, but are seen very little in 325 the CTE and CTS sectors (Fig. 4). Applying the simple-signed rule and the blockmodeling 326 algorithm to delineate communities, both find that there are three distinct communities (Fig. 327 4). Though, in the case of AVS the simple sign-rule suggested multi-membership, while the 328 blockmodeling algorithm found AVS to be part of the cluster containing (AVO, AVN, AVE) 329 or that the two clusters (orange and purple in fig. 4) merged into one depending on the choice 330 of weighting parameter (i.e., emphasizing positive or negative edges).



332 Figure 4: Similarity and dissimilarity between sectors in the beluga whale population of the 333 St. Lawrence. The green edges between two sectors signify that the sectors share high/low 334 users, while red edges signify that they have dissimilar high/low users. The lack of an edge 335 signifies that the high/low users of one sector does not provide information about the 336 high/low users of other sectors. Nodes represent sectors, and are coloured based on shared 337 communities: i.e., shared green edges, and no red edges. Node sizes represent the magnitudes 338 of individual differences in use within the sector, i.e., larger nodes suggest larger differences 339 between high and low users.

340

341 **4. Discussion**

Here we've shown that using photo-ID data with a multilevel multinomial model it is possible to estimate socio-spatial networks, identifying spatial communities while controlling for sampling biases. Our results suggest that the beluga population shows a non-random social-spatial structuring within the summer range.

346 Within sectors of the St. Lawrence, our model suggests that subsets of belugas are 347 heavily using some sectors, while other sectors show little evidence of differences in use. The 348 magnitude of individual differences in each sector, i.e., how much individuals differ in their 349 probability of being observed in a particular sector, shows that the CTN sector, in particular, 350 has very little in the way of individual differences in the probability of being seen in that 351 sector (Table 1). This result suggests that the CTN sector is used similarly by most 352 individuals, and represents a potential high mixing zone for the population. In contrast, the 353 AME sector shows the highest level of individual differences, suggesting that there are large 354 differences in how beluga are using this sector. These results suggest that the population is 355 not randomly mixing with the St. Lawrence, and that there are belugas that make use of some 356 sectors more than other belugas.

357 Between sectors of the St. Lawrence, our results add to the evidence that the beluga 358 population cannot be assumed to be randomly mixing within its summer habitat. Rather, 359 comparing the individual differences in beluga usage patterns within sectors suggests 360 similar/dissimilar user populations across sectors (Fig. 4). By using correlations between 361 sector usage patterns to create a socio-spatial network, and running community detection 362 algorithms, our results found that there are spatially distinct communities that make use of 363 particular regions of the St. Lawrence and the Saguenay River (Fig. 4). We found that the 364 beluga population in the St. Lawrence could be separated into three distinct communities: 1)

The lower St. Lawrence (AVO, AVS, AVE, AVN), the Saguenay River and mouth (BSM,
SAG, CTN), and the upper and eastern portion of the St. Lawrence (CTE, CTS, CTO, AME,
AMS) (Fig. 4).

368 Our findings have direct implications for estimating the impacts of anthropogenic 369 disturbances on this population. As the population shows evidence of spatially restricted 370 habitat use, disturbances to particular regions can have a disproportionate impact on 371 particular segments of the larger population. In particular, the cumulative impacts over time 372 are likely to be greatly increased in some segments while reduced in other segments of the 373 population, altering estimations of the distribution of impacts. If cumulative impacts, such as 374 noise, or environmental contaminants, have a threshold beyond which individual survival is 375 greatly reduced, properly estimating the distribution of cumulative impacts can have large 376 implications for conservation management. Our results add to the current understanding of 377 socio-spatial structuring within this population (Michaud, 1993, 2005), and suggest that more 378 empirical data, e.g., photo-ID data, movement data, aerial surveillance, should be collected to 379 better refine socio-spatial mixing in this population.

380 The modeling approach presented in this paper relies on defined sectors within a 381 particular spatial range, e.g., SAG sector, CTN sector... etc (Fig. 1). In some cases, these 382 delineations can be justified as they identify management zones, but in other cases, the 383 delineation and scale of these sectors can be delineated somewhat arbitrarily. Future work 384 could assess the use of continuous random effects (as opposed to categorical) where 385 individual differences in the probability of being seen could be on a continuous surface. Point 386 estimates of individual differences could then be estimated at any location, and correlations 387 between individual differences obtained between any two points in continuous space. This 388 approach could avoid reliance on user-defined sectors and facilitate a means of looking at the 389 results at different scales (e.g., grids of points at various scales could be used when estimating correlations). Similarly, given that the method requires repeated sampling of individuals to
obtain probability of being seen in any one sector, there is a reliance on longitudinal data. In
well sampled populations, it could be feasible to estimate the change in time of individuals
being seen in particular sectors. Here differences in being seen in any particular area could
be explicitly modeled to capture any temporal changes in community substructures, or
developmental trajectories related to habitat use.

396 In terms of implementing the multilevel multinomial model on other photo-ID

datasets, the use of test datasets should hold a prominent role in the analysis. The use of

398 permutation/randomization methods to both generate structured and unstructured datasets,

399 while maintaining the sample size distribution of the original datasets, can be very valuable in

400 helping to set model priors and to interpret the final model results. The use of permutation

401 approaches is common in social network analysis (Croft et al., 2011; Farine, 2017), and is

402 becoming more common in statistical workflows more generally (Gelman et al., 2013;

403 McElreath, 2020).

404

405 **5.** Conclusions

406 We have introduced the use of multilevel multinomial modeling to estimate socio-407 spatial networks from photo-ID data. We've shown, using testing datasets, that the proposed 408 method is effective at detecting socio-spatial structures. When applied to 18 years of photo-409 ID data from an endangered population of beluga whales in the St. Lawrence, our results 410 suggest strong evidence that the population has three distinct spatial communities. We 411 suggest that multilevel multinomial models can be effective in extracting socio-spatial 412 structuring within animal populations monitored by photo-ID, and can have direct 413 implications for conservation management.

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417	collecting these long term data.
418	
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421	Secrétariat à la stratégie maritime du Québec.
422	
423	8. Data accessibility
424	The permuted and simulated photo ID datasets are available on github
425	(github.com/tbonne/photoID_multinomial), along with code used in the analysis.
426	
427	9. Author Contributions
428	RM collected the data; TRB conceived the analytical methodology and performed the
429	analysis; all authors contributed critically to the drafts and gave final approval for
430	publication.
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497 Supplementary material:

498

499 *Use of a preset reference sector:*

We use the simulated dataset with structure, presented in the main text, to fit a multilevel multinomial model with and without a preset reference sector. That is, in the model with the preset reference sector we duplicated each individual's photo-IDs and placed them in the reference sector. This results in a probability of 0.5 for being seen in the reference sector for all individuals. The results suggest that using a preset reference sector where all individuals have the same probability of being seen is required to estimate correlations between sectors and produces appropriate socio-spatial networks (Fig S1).

507

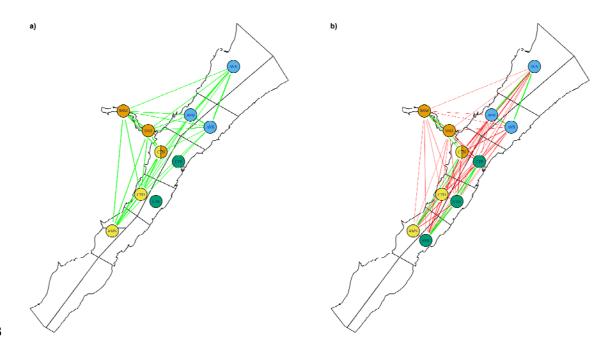


Fig S1: The estimate socio-spatial networks for a multilevel multinormial model a) without a preset reference sector, i.e., AME is used as the reference, and b) with a preset reference sector. The green edges between two sectors signify that the sectors share high/low users, while red edges signify that they have dissimilar high/low users. The lack of an edge signifies that the high/low users of one sector does not provide information about the high/low users of

514 other sectors. Nodes represent sectors, and are coloured based on the communities imposed

- 515 when simulating the data.
- 516
- 517 Full model table:
- 518

519	Table S1: Parameter estimates from the multilevel multinomial model predicting probability
520	of capturing a photo-ID by sector. Estimated mean probability (logit scale), individual
521	differences in mean probability ('sd'), and correlation of individual differences between the
522	different sectors are presented along with an estimate of their 95% credible intervals. Positive
523	correlations suggest that the high/low users in one sector are similarly high/low users in
524	another sector, while negative correlations suggest high/low users in one sector are the
525	low/high users in another sector.

Туре	Parameter	Estimate	SD	l-95% CI	u-95% CI
Mean probability (logit scale)					
	mu_AME	-5.85	0.28	-6.43	-5.33
	mu_AMN	-5.60	0.24	-6.15	-5.17
	mu_AMO	-6.68	0.34	-7.41	-6.08
	mu_AMS	-6.33	0.33	-7.00	-5.73
	mu_AVE	-5.47	0.24	-5.99	-5.02
	mu_AVN	-5.06	0.22	-5.52	-4.65
	mu_AVO	-1.82	0.05	-1.93	-1.72
	mu_AVS	-2.90	0.07	-3.04	-2.76
	mu_BSM	-3.78	0.13	-4.04	-3.52
	mu_CTE	-2.20	0.05	-2.30	-2.10
	mu_CTN	-1.24	0.03	-1.30	-1.18
	mu_CTO	-4.58	0.16	-4.92	-4.28

	mu_CTS	-4.09	0.13	-4.35	-3.85
	mu_SAG	-2.78	0.09	-2.96	-2.61
Individual Di	fferences				
	sd(mu_AME)	2.00	0.19	1.64	2.38
	sd(mu_AMN)	0.76	0.29	0.14	1.31
	sd(mu_AMO)	0.65	0.38	0.05	1.41
	sd(mu_AMS)	1.53	0.22	1.10	1.98
	sd(mu_AVE)	1.16	0.20	0.78	1.57
	sd(mu_AVN)	1.30	0.17	0.98	1.65
	sd(mu_AVO)	0.78	0.05	0.68	0.88
	sd(mu_AVS)	0.83	0.07	0.70	0.97
	sd(mu_BSM)	1.30	0.11	1.09	1.52
	sd(mu_CTE)	0.78	0.05	0.68	0.89
	sd(mu_CTN)	0.33	0.03	0.27	0.39
	sd(mu_CTO)	1.27	0.14	1.00	1.55
	sd(mu_CTS)	1.18	0.11	0.97	1.40
	sd(mu_SAG)	1.16	0.08	1.01	1.31
Correlations	Correlations between individual differences				
	cor(mu_AME,mu_AMN)	0.39	0.19	-0.03	0.71
	cor(mu_AME,mu_AMO)	0.27	0.24	-0.26	0.67
	cor(mu_AMN,mu_AMO)	0.14	0.24	-0.36	0.57
	cor(mu_AME,mu_AMS)	0.63	0.11	0.39	0.82
	cor(mu_AMN,mu_AMS)	0.25	0.21	-0.19	0.62
	cor(mu_AMO,mu_AMS)	0.27	0.24	-0.26	0.67
	cor(mu_AME,mu_AVE)	0.18	0.16	-0.14	0.46
	cor(mu_AMN,mu_AVE)	0.15	0.21	-0.28	0.56

cor(mu_AMO,mu_AVE)	0.10	0.23	-0.36	0.52
cor(mu_AMS,mu_AVE)	0.05	0.19	-0.32	0.41
cor(mu_AME,mu_AVN)	-0.04	0.15	-0.33	0.25
cor(mu_AMN,mu_AVN)	-0.02	0.21	-0.42	0.39
cor(mu_AMO,mu_AVN)	0.00	0.22	-0.42	0.44
cor(mu_AMS,mu_AVN)	-0.07	0.18	-0.41	0.28
cor(mu_AVE,mu_AVN)	0.50	0.14	0.19	0.75
cor(mu_AME,mu_AVO)	-0.21	0.10	-0.40	0.00
cor(mu_AMN,mu_AVO)	0.03	0.19	-0.35	0.40
cor(mu_AMO,mu_AVO)	-0.06	0.21	-0.45	0.36
cor(mu_AMS,mu_AVO)	-0.28	0.14	-0.54	0.01
cor(mu_AVE,mu_AVO)	0.48	0.13	0.22	0.71
cor(mu_AVN,mu_AVO)	0.50	0.10	0.28	0.69
cor(mu_AME,mu_AVS)	0.13	0.12	-0.10	0.36
cor(mu_AMN,mu_AVS)	0.21	0.18	-0.16	0.53
cor(mu_AMO,mu_AVS)	0.13	0.21	-0.31	0.50
cor(mu_AMS,mu_AVS)	0.03	0.15	-0.26	0.32
cor(mu_AVE,mu_AVS)	0.40	0.14	0.12	0.66
cor(mu_AVN,mu_AVS)	0.38	0.12	0.14	0.60
cor(mu_AVO,mu_AVS)	0.54	0.08	0.38	0.69
cor(mu_AME,mu_BSM)	-0.57	0.09	-0.75	-0.38
cor(mu_AMN,mu_BSM)	-0.37	0.17	-0.66	0.00
cor(mu_AMO,mu_BSM)	-0.22	0.23	-0.62	0.26
cor(mu_AMS,mu_BSM)	-0.37	0.13	-0.62	-0.10
cor(mu_AVE,mu_BSM)	-0.34	0.13	-0.59	-0.06
cor(mu_AVN,mu_BSM)	-0.22	0.12	-0.44	0.02

cor(mu_AVO,mu_BSM)	-0.16	0.08	-0.32	0.01
cor(mu_AVS,mu_BSM)	-0.66	0.07	-0.80	-0.51
cor(mu_AME,mu_CTE)	0.54	0.10	0.34	0.72
cor(mu_AMN,mu_CTE)	0.34	0.18	-0.04	0.65
cor(mu_AMO,mu_CTE)	0.21	0.23	-0.29	0.60
cor(mu_AMS,mu_CTE)	0.38	0.14	0.10	0.62
cor(mu_AVE,mu_CTE)	0.07	0.14	-0.21	0.36
cor(mu_AVN,mu_CTE)	-0.11	0.12	-0.35	0.14
cor(mu_AVO,mu_CTE)	-0.10	0.08	-0.26	0.07
cor(mu_AVS,mu_CTE)	0.52	0.08	0.34	0.67
cor(mu_BSM,mu_CTE)	-0.80	0.06	-0.89	-0.68
cor(mu_AME,mu_CTN)	-0.26	0.12	-0.49	-0.03
cor(mu_AMN,mu_CTN)	-0.24	0.19	-0.59	0.16
cor(mu_AMO,mu_CTN)	-0.12	0.21	-0.50	0.32
cor(mu_AMS,mu_CTN)	-0.09	0.15	-0.39	0.21
cor(mu_AVE,mu_CTN)	-0.53	0.13	-0.77	-0.26
cor(mu_AVN,mu_CTN)	-0.44	0.12	-0.67	-0.18
cor(mu_AVO,mu_CTN)	-0.75	0.07	-0.86	-0.60
cor(mu_AVS,mu_CTN)	-0.63	0.09	-0.79	-0.44
cor(mu_BSM,mu_CTN)	0.47	0.09	0.28	0.64
cor(mu_CTE,mu_CTN)	-0.25	0.10	-0.46	-0.05
cor(mu_AME,mu_CTO)	0.63	0.09	0.43	0.79
cor(mu_AMN,mu_CTO)	0.28	0.20	-0.15	0.63
cor(mu_AMO,mu_CTO)	0.16	0.22	-0.30	0.56
cor(mu_AMS,mu_CTO)	0.55	0.13	0.27	0.78
cor(mu_AVE,mu_CTO)	-0.12	0.17	-0.45	0.22

cor(mu_AVN,mu_CTO)	-0.24	0.15	-0.54	0.07
cor(mu_AVO,mu_CTO)	-0.55	0.09	-0.72	-0.36
cor(mu_AVS,mu_CTO)	-0.16	0.12	-0.39	0.07
cor(mu_BSM,mu_CTO)	-0.33	0.11	-0.53	-0.12
cor(mu_CTE,mu_CTO)	0.39	0.11	0.17	0.60
cor(mu_CTN,mu_CTO)	0.19	0.12	-0.06	0.42
cor(mu_AME,mu_CTS)	0.70	0.08	0.53	0.85
cor(mu_AMN,mu_CTS)	0.42	0.19	0.00	0.73
cor(mu_AMO,mu_CTS)	0.21	0.23	-0.27	0.61
cor(mu_AMS,mu_CTS)	0.42	0.14	0.14	0.67
cor(mu_AVE,mu_CTS)	0.16	0.16	-0.15	0.45
cor(mu_AVN,mu_CTS)	0.06	0.14	-0.23	0.33
cor(mu_AVO,mu_CTS)	-0.04	0.10	-0.23	0.15
cor(mu_AVS,mu_CTS)	0.42	0.11	0.20	0.61
cor(mu_BSM,mu_CTS)	-0.72	0.08	-0.85	-0.55
cor(mu_CTE,mu_CTS)	0.64	0.09	0.46	0.80
cor(mu_CTN,mu_CTS)	-0.37	0.11	-0.57	-0.14
cor(mu_CTO,mu_CTS)	0.48	0.11	0.25	0.68
cor(mu_AME,mu_SAG)	-0.43	0.09	-0.61	-0.24
cor(mu_AMN,mu_SAG)	-0.36	0.17	-0.66	0.00
cor(mu_AMO,mu_SAG)	-0.17	0.22	-0.56	0.29
cor(mu_AMS,mu_SAG)	-0.22	0.13	-0.47	0.03
cor(mu_AVE,mu_SAG)	-0.44	0.12	-0.68	-0.19
cor(mu_AVN,mu_SAG)	-0.37	0.10	-0.57	-0.16
cor(mu_AVO,mu_SAG)	-0.48	0.06	-0.60	-0.35
cor(mu_AVS,mu_SAG)	-0.80	0.06	-0.90	-0.68

cor(mu_BSM,mu_SAG)	0.83	0.04	0.74	0.91
cor(mu_CTE,mu_SAG)	-0.68	0.06	-0.79	-0.55
cor(mu_CTN,mu_SAG)	0.67	0.07	0.52	0.80
cor(mu_CTO,mu_SAG)	-0.12	0.11	-0.32	0.09
cor(mu_CTS,mu_SAG)	-0.62	0.08	-0.77	-0.45

526

527

528

529 *Note on interpreting low sd within sectors:*

530 Similar to the CTN sector, the sector AMO is also estimated to have a low magnitude of

531 individual differences. However, it has a large uncertainty in this estimate. This highlights

that it is possible to have little individual differences in a sector due to either: 1) limited data,

resulting in all individuals being pooled to the mean value, and 2) limited data is not a factor,

but individuals are using this sector relatively equally. Care should therefore be taken when

535 interpreting the magnitude of individual differences, nevertheless, the estimated uncertainty

around magnitude estimates is one way to help identify sectors with limited data.