1	Defining indices of ecosystem variability using biological samples of fish
2	communities: a generalization of empirical orthogonal functions
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18 Abstract:

Multivariate data reduction techniques are widely used to describe modes of variability in 19 atmospheric and oceanographic conditions for the world's oceans. Dominant modes of 20 21 variability such as the Pacific Decadal Oscillation (PDO) are typically defined as a statistical summary of physical measurements, and include both principle components representing modes 22 23 of variability over time, and an empirical orthogonal function (EOF) giving the spatial pattern associated with a positive or negative phase for each mode. Typically, these indices are 24 compared with biological conditions to describe or predict physical drivers of ecological 25 26 dynamics. In some circumstances, however, it may instead be useful to apply EOF analysis directly to biological measurements, estimating indices of biological variability as well as maps 27 of biological response associated with each index. We therefore develop a generalization of 28 EOF analysis that can be applied directly to multispecies biological samples using a multivariate 29 spatio-temporal model. These biologically derived indices can then be compared with relevant 30 31 indices derived from physical data, or used as covariates in spatially-varying coefficient models. We first show that a spatio-temporal model can replicate previous EOF estimates of the PDO and 32 North Pacific Gyre Oscillation. We then identify three axes of variability in the eastern Bering 33 34 Sea using biomass-sampling data for fourteen bottom-associated fishes and decapod crustaceans from 1982-2017. The first axis represents habitat preferences that are stable over time, and the 35 36 second represents a multi-decadal trend in distribution for most species; for example, showing an 37 increasing density for Alaska skate and arrowtooth flounder in the middle and inner domain. Finally, the third axis shows high interannual variability from 1982-1998 switching to multiyear 38 39 stanzas from 1999-2017 and is highly correlated (0.87) with the extent of the cold bottom 40 temperatures in this region and associated impacts on Alaska pollock and Pacific cod. These

41 axes represent ecological dynamics for adult fishes and therefore integrate the impact of bottom42 up and top-down processes, and they also confirm the importance of cold-pool extent for fish
43 distribution in the Bering Sea while visualizing its varied impact on individual species.
44 Moreover, this spatio-temporal approach allows oceanographers to define annual indices
45 representing modes of variability in diverse biological communities from widely available field46 sampling data.

48 Introduction

Oceanographers are confronted with a tremendous challenge in summarizing multivariate 49 physical and biological processes into a tractable number of dominant patterns, which can then 50 be readily communicated to other fields (e.g., fisheries scientists), used as covariates in climate 51 models, or used in many other ways. One approach to this challenge is to identify dominant axes 52 53 of variability for a given process, and then to represent the process as an index that varies over time, combined with a loadings map expressing the spatial pattern associated with the index. 54 Familiar examples include the Pacific Decadal Oscillation (PDO; Mantua et al., 1997) and North 55 56 Pacific Gyre Oscillation (NPGO; Di Lorenzo et al., 2008). These indices capture patterns of basin-scale variability that play leading roles in the dynamics of regional ecosystems (Schwing et 57 al., 2010), and also tend to capture variability in a suite of cross-correlated, ecologically 58 important processes (Stenseth et al., 2003), making them valuable tools for summarizing climate 59 effects on marine biota. For example, the importance of the PDO was established via 60 61 comparison with records of salmon returns in the Pacific Ocean (Mantua et al., 1997), while the importance of the NPGO was demonstrated via its correlation with nutrients and chlorophyll 62 concentrations in southern California (NPGO; Di Lorenzo et al., 2008). These ocean climate 63 64 indices are typically referred to as principal components (PCs) for the temporal index and empirical orthogonal functions (EOFs) for the loadings (e.g., Trenberth et al., 2014). Here we 65 broadly use the term "EOF analysis" to refer to the algorithm generating both the temporal index 66 67 and the spatial map associated with it, while recognizing that different authors use different algorithms for individual steps in the EOF analysis (e.g., PCA vs. factor analysis for generating 68 69 the temporal index).

70 There have been several applications of EOF analysis to biological spatio-temporal data in the last decade (e.g., Morfin et al., 2012; Marshall et al., 2016). However, previous applications 71 of data reduction techniques applied to biological data involve either the aggregation across 72 space and time so to obtain regional averages of multi-species biomass (e.g., PCA applied to 73 74 annual indices of species biomass, Planque and Arneberg (2018)), or aggregation across species 75 so to obtain single-species measurement of biomass across multiple sampling sites (e.g., EOF analysis of spatio-temporal data from single species). These data manipulations inevitably reduce 76 the scope of ecological inference, sacrificing either small-scale spatiotemporal dynamics or the 77 78 ability to generalize the effect of climate across multiple taxa (Puerta et al., 2019). In particular, we identify three technical hurdles when applying EOF analysis directly to 79 80 biological sampling data: (1) biomass-sampling data has many zeros with few extreme values (a highly skewed "dust bunny distribution" sensu McCune and Root (2015)) which is poorly 81 represented using a normal distribution; (2) biological samples often arise from spatially 82 unbalanced sampling, which complicates the creation of spatial maps in unsampled areas or 83 when fitting to spatially unbalanced sampling programs; and (3) biological sampling is often 84 available for multiple species, and therefore EOF analysis must be generalized for multivariate 85 86 data sets, where each variable has both temporal and spatial expression. Despite the profusion of research using EOFs to characterize physical oceanographic 87 88 conditions, it is not always clear how to relate these indices to biological processes. This 89 commonly involves a multi-step workflow, where physical variability is first summarized with

90 EOF analysis, and the resulting index is then used as a covariate in a separate statistical model

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92 relationships between physical conditions and biological responses in ocean environments either

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linking physical oceanography to biological or social outcomes. Unfortunately, many published

break down over time or have poor skill when forecasting (Myers, 1998; Thorson, 2019a). We 93 note two ways that this multi-step workflow may contribute to poor out-of-sample predictive 94 skill for statistical physics-biology relationships. First, climate indices derived from EOF 95 analysis summarize physical dynamics across a large spatial domain. These regional indices are 96 then correlated with local conditions, and biological responses are then correlated with local 97 98 conditions for each individual population (Newman et al., 2016; Stenseth et al., 2003; Wills et al., 2018). However, the correlation between regional indices and local physical conditions can 99 change over time, thus changing their correlation with local biological responses (Litzow et al., 100 101 2018; Newman et al., 2016). Second, the solution of the EOF analysis is defined such that the first axis explains the most variance in the underlying data, the second axis explains the second-102 most, etc. However, these definitions are fixed in time while different axes of physical variation 103 104 may have a larger impact on biological during some conditions than others, e.g., physical variation associated with juvenile production may be more important after fishing has reduced 105 106 the age-structure of a population (Hsieh et al., 2006).

Given these difficulties when using oceanographic indices of physical habitat to describe 107 biological responses, we see a useful role for indices estimated directly from multivariate 108 109 biological variables. In particular, recent improvements in statistical computation and spatial statistics have allowed the growth of multivariate spatio-temporal models (Clark et al., 2014; 110 111 Latimer et al., 2009; Ovaskainen et al., 2017). For example, spatial factor analysis (Thorson et 112 al., 2015b) could be used to generalize EOF analysis using biomass samples for multiple species simultaneously, and therefore could represent indices of variation in the ecological dynamics that 113 114 arise from both bottom-up and top-down processes. We propose that these approaches would be 115 useful for compressing one or more variables to a single (or few) easily visualized time series.

Indices derived from biological sampling data could then be correlated with similar indices
estimated from physical data to explore physics-biology linkages, as a descriptive summary of
ecological dynamics (e.g., McClatchie et al., 2018), or used directly as covariates in other
biological models (e.g., O'Leary et al., 2018).

In this study, we develop a method to estimate dominant modes of variability (i.e., one or 120 121 more indices, each associated with a map showing spatial patterns in the positive phase of a given index) that can be applied to noisy, multispecies field samples of biological variables. We 122 then demonstrate that this approach generalizes conventional EOF analysis by replicating 123 124 estimates of PDO and the NPGO using North Pacific sea surface temperature fields. Finally, we demonstrate the approach using data for fourteen bottom-associated fish and decapod species in 125 the eastern Bering Sea. Many studies in this system have linked the spatial distribution of fishes 126 127 to the location and spatial extent of cold near-bottom temperatures (e.g., Wyllie-Echeverria and Wooster, 1998; Baker and Hollowed, 2014), although long-term trends in distribution appear to 128 be independent of cold-pool extent for some species (Mueter and Litzow, 2008). Similarly, our 129 method estimates an index of ecosystem variability that is highly correlated with the spatial 130 extent of cold near-bottom waters (termed the "cold pool"), and therefore corroborates the 131 132 important role and estimates the spatially varying effect of coupled winter ice cover and summer bottom temperature in that region over the past 36 years. 133

134 Methods

135 *A brief history of Empirical Orthogonal Function (EOF) analysis in physical oceanography*

136 Climatologists, meteorologists, and oceanographers have used empirical orthogonal functions

to characterize dominant modes of variability in physical ocean conditions for over fifty years.

138 Grimmer (1963) applied factor analysis to sea surface temperature anomalies in the North

Atlantic to show that 80% of variance can be explained by a small number of latent variables. 139 Kidson (1975a) introduced a similar method using principal components analysis (PCA) applied 140 to monthly average precipitation and sea surface pressure to similarly show that a reduced set of 141 variables could explain >75% of the original variance. These studies established the common 142 practice wherein a multivariate statistical techniques (PCA or factor analysis) is applied to 143 144 spatially replicated measurements of a physical variable at multiple times (e.g., years or months), where a small number of dominant axes can be used to explain the large portion of variance in 145 the original process. 146

147 Empirical orthogonal functions have subsequently been used to define or analyze several of the most widely-known global and regional processes in oceanography. Kidson (1975b) used 148 149 EOF to measure the Southern Oscillation, a pattern in equatorial surface pressure and precipitation that was originally described by Walker (1924). More recently, Mantua et al. 150 (1997) used EOF to define the PDO as the dominant mode of variability in sea surface 151 temperature (SST) in the North Pacific, and the second mode was further explored and 152 interpreted by Bond et al. (2003) and subsequently called the "Victoria mode" (Ding et al., 153 2015). Similarly, Di Lorenzo et al. (2008) defined the NPGO as the second mode of variability 154 155 for sea surface height (SSH) anomalies, and the NPGO is correlated with the Victoria mode via the high correlation between SST and SSH. 156

The PDO and NPGO have seen broad use in biological oceanography due to the correlation between these physical indices and biological production that underlies changes in fishery productivity (Di Lorenzo et al., 2008; Mantua et al., 1997). For example, the importance of the PDO was originally demonstrated based on cycles in salmon productivity between Alaska and Oregon/Washington stocks (Mantua et al., 1997). However, the correlation between the PDO

and salmon productivity has declined since the description of the PDO, apparently due to the
 changing associations between the PDO and localized physical processes that underlie salmon
 productivity for individual salmon stocks (Litzow et al., 2018).

165 Finally, we note ongoing research that defines EOFs in the context of a statistical model that

separately estimates measurement errors from physical variation. For example, Grimmer (1963)

used factor analysis to separate measurement and process errors, and Ghil et al. (1981)

subsequently defined a multivariate Kalman filter that explicitly models covariation in physical

169 dynamics. This Kalman-filter interpretation of EOF was subsequently extended by Wikle and

170 Cressie (1999), who introduced a descriptive spatial process wherein the physical process is

171 more similar at nearby than at distant locations. Subsequently, these "spatio-temporal" models

have rapidly developed due to improvements in statistical and computational techniques (e.g.,

173 Lindgren et al., 2011), and now are also widely used in ecology and fisheries science

174 (Ovaskainen et al., 2017; Thorson, 2019b). To our knowledge, however, this statistical

generalization of EOF analysis to generate the index and the spatial map simultaneously has not

been applied to multi-guild biological data (e.g., abundance or biomass for species in a

177 community) while accounting for variable dynamics across both space and time.

178 *General approach*

We develop a method to estimate one or more ecosystem indices based on biomass-samplingdata for multiple species. Each index *f* includes two components:

181 1. a spatial map $\mathcal{X}_{f,c}(s)$ representing whether a given location *s* has a positive or negative value

182 (as well as the magnitude of that value) during a "positive phase" of the index; and

183 2. a time-series $\lambda_f(t)$ indicating whether a given time t has a positive phase ($\lambda_f(t) > 0$) or

184 negative phase $(\lambda_f(t) < 0)$, as well as whether a given year has a weak magnitude $(\lambda_f(t)$

185 within the 25% and 75% quantiles for λ_f) or a strong magnitude ($\lambda_f(t)$ outside the 25% and 186 75% quantiles).

187 Importantly, the map $\mathcal{X}_{f,c}(s)$ associated with each time series $\lambda_f(t)$ differs for each category *c*

188 of n_c modeled categories, and this allows our time series to represent multivariate data, for

189 example, multispecies biomass samples from bottom trawl surveys. In the following, we focus

on annual variation (i.e., *t* indexes different years), but the process could instead represent other
time intervals (weekly, monthly, etc.).

To estimate these ecosystem indices, we begin by defining a predictor variable $\tilde{Y}_{c,t}(s)$ that includes the net effect of all estimated oceanographic indices. Each predictor variable is a linear combination of n_f oceanographic indices:

$$g\left(\underbrace{\tilde{\mathcal{Y}}_{c,t}(s)}_{predictor variable}\right) = \underbrace{\beta_{c,t}}_{intercepts} + \underbrace{\sum_{f=1}^{n_f} \underbrace{Index \, phase \, Index \, map}_{\mathcal{X}_f(t)} \qquad 1$$

where $g(\tilde{y})$ is a link function transforming the linear predictor to response \tilde{y} and $\beta_{c,t}$ are intercepts that vary among categories and times. This predictor variable is then estimated by minimizing the difference between it and measurements $\mathcal{Y}_{c,t}(s)$ of each response variable:

$$\mathcal{Y}_{c,t}(s) \sim h\left(\widetilde{\mathcal{Y}_{c,t}}(s), \dots\right)$$

where *h* is a probability distribution function for measurements $\mathcal{Y}_{c,t}(s)$ given their predicted values $\widetilde{\mathcal{Y}_{c,t}}(s)$. For a discussion of how this general model relates to previously developed spatio-temporal models, please see Appendix 1.

201 Demonstrating the similarity to Empirical Orthogonal Function (EOF) analysis

202 We first seek to show that this approach can generalize the EOF analysis that is widely used in

203 physical oceanography to generate indices. Published studies have implemented EOF using

204	either principle components analysis (PCA) or factor analysis (FA), and both involve a matrix
205	$Y_t(s)$ of measurements of a physical variable at each location and time. Anomalies from long-
206	term climate are then typically calculated, $Y_t^*(s) = Y_t(s) - \frac{1}{n_t} \sum_{t=1}^{n_t} Y_t(s)$, and these anomalies
207	are used to calculate the sample covariance $V_{t,t}$ for anomalies between any pair of times. The
208	analyst then applies an eigendecomposition (for PCA) or minimization algorithm (FA) to
209	identify a set of n_f orthogonal axes of covariation, ranked from most important to least
210	important, such that the first few axes explain the majority of covariation. These axes are then
211	treated as the index $\lambda_f(t)$, and the spatial map $x_f(s)$ associated with each index is calculated as
212	either the correlation or regression of $\lambda_f(t)$ and $y_f(s, t)$.
213	We replicate a univariate EOF analysis by simplifying the general model in four ways:
214	1. Eliminate notation for multiple categories such that it is applied to a single response;
215	2. Use an identity link function;
216	3. Use a normal distribution for observations; and
217	4. Replace functions with matrices, i.e., $\mathcal{X}_{f,c}(s)$ with a matrix X representing the predicted
218	value of each oceanographic variable $x^*(s_i, f)$ at the location for each sample <i>i</i> for each
219	index, and $\tilde{\mathcal{Y}}_{c,t}(s)$ with \tilde{y}_i representing the prediction for that sample <i>i</i> .
220	These three changes result in the following model:
	$\tilde{y}_i = \beta(t_i) + \sum_{f=1}^{n_f} \lambda_f(t_i) x^*(s_i, f) $ 3a

$$y_i \sim Normal(\tilde{y}_i, \sigma^2)$$
 3b

where σ^2 is the variance of measurement errors, which is minimized by explaining variation in the response y(s, t) to estimated indices, and $x^*(s_i, f)$ is calculated from a predictive-process model of spatial variation. We note that we do not center the data y(s, t) prior to analysis using 224 this generalized model, so the first axis in this generalized model corresponds to persistent spatial differences that would otherwise be eliminated by centering the data. This is different 225 from conventional EOF analysis (which does center the data to calculate anomalies prior to 226 analysis), and this difference means that the 2nd axis from the generalized model is similar to the 227 1^{st} axis from conventional EOF, the 3^{rd} axis is similar to the 2^{nd} from conventional EOF, etc. 228 However, centering the data does not extend to other common forms of data analysis (e.g., 229 generalized linear models) so not centering the data is important for subsequent generalizations. 230 Also differing from conventional EOF, we estimate a separate intercept β_t for every year, so that 231 this term captures interannual variability in y(s, t) among years (i.e., the increasing trend in 232 surface temperatures due to climate forcing). We estimate a separate intercept for each year to 233 234 match model specification for the multi-species extension to biological sampling data, as justified below, and future applications could easily specify a model where this intercept is 235 236 constant across years (to be more similar with conventional EOF analysis). We then compare 237 results with public values for the PDO (obtained Nov. 6, 2018 from http://research.jisao.washington.edu/pdo/PDO.latest.txt) and NPGO (obtained Nov. 6, 2018 from 238 239 http://www.o3d.org/npgo/). 240 Extension to multi-species biomass-sampling data 241 We also seek to show how this approach can include biomass-sampling data for multiple species,

such as are widely available worldwide from resource surveys of fish stocks. For multivariate
biomass-sampling data we define a more complicated sampling process, which involves several
modifications to the general model:

1. Use multiple categories, each with an independent intercept for each species and year;

246 2. Use a Poisson-link delta-model (Thorson 2017) where the linear predictor is associated with 247 numbers density n, where encounter probability p_i for sample i is derived as a 248 complementary log-log link from numbers density $p_i = 1 - \exp(-n_i)$ and expected biomass 249 when encountered r_i is defined such that it is proportional to numbers density, $r_i =$ 250 $\frac{n_i}{p_i}w(c_i, t_i)$, where $w(c_i, t_i)$ is the biomass per individual, which is estimated separately for 251 each category and year; and

- 252 3. Minimize the negative log-likelihood for encounters and sampled biomass for each species.
- 253 These changes result in the following model:

$$\log(n_i) = \beta(c_i, t_i) + \sum_{f=1}^{n_f} \lambda_f(t_i) x^*(s_i, c_i, f)$$

$$\Pr(B = b(s_i, c_i, t_i)) = \begin{cases} 1 - p_i & \text{if } B = 0\\ Lognormal(\log(r_i), \sigma_t^2) & \text{if } B > 0 \end{cases}$$
4a
4b

Where $\beta(c, t)$ is again an intercept for every category and time, Eq. 4b represents a delta-model 254 distribution where $1 - p_i$ is the probability mass associated with B = 0, Lognormal(a, b) is a 255 lognormal probability density function with logmean a and log-variance b, and σ_t^2 is the residual 256 estimated log-variance in positive catch rates. We specify a separate intercept for every category 257 and time because, in our experience, total abundance for marine fishes often varies substantially 258 259 among years due endogenous biological processes (e.g., variable production of juveniles; 260 Thorson et al. (2015a)) and the annually varying intercept "controls" for this variation such that 261 remaining model explain shifts in distribution rather than total abundance. This model includes a log-link (Eq. 4a) such that variation a 0.01 increase in $\lambda_f(t_i)x^*(s_i, c_i, f)$ corresponds to an 262 approximately 1% increase in expected biomass. This relative scale implies that all coefficients 263 264 representing spatial and temporal variation (i.e., the right-hand side of Eq. 4a) are dimensionless.

265 In the introduction, we noted three difficulties with applying EOF to biological sampling data: (1) many zeros and a skewed distribution; (2) spatially unbalanced sampling; and (3) 266 multiple response variables. These difficulties are addressed by (1) applying a delta-model with 267 a linear predictor that is shared among years and species; (2) projecting the linear predictor from 268 269 $n_{\rm s}$ knots to any possible location within the modeled spatial domain, including locations with missing or spatially misaligned data; and (3) associating each mode of variability $\lambda_f(t)$ with a 270 map representing the impact $x^*(s, c, f)$ of a positive phase for each of n_c modeled variables at 271 any location s within a defined spatial domain. In particular, accounting for spatially misaligned 272 273 data allows EOF to be applied to multiple sampling programs, operating at different locations or over different spatial domains, although we do not explore the idea further here. 274

275 *Parameter estimation*

276 Parameters for both configurations of the general model can be estimated using a publicly

277 available R package VAST for vector autoregressive spatio-temporal models (VAST; Thorson

and Barnett, 2017), using release number 3.1.0 (https://github.com/James-Thorson-

279 <u>NOAA/VAST</u>). This package estimates spatial variables x(s, c, f) for n_s "knots" as random

280 effects following a Gaussian Markov random field, uses a predictive process formulation to

interpolate the value of $x^*(s_i, c_i, f)$ for the location s_i and category c_i of sample *i* given

282 x(s, c, f), and models the correlation in x(s, c, f) between any two knots using a stochastic

partial differential equation (SPDE) approach (Lindgren et al., 2011) which approximates a

284 Matérn correlation function (see Appendix 2 for details). Fixed effects are then estimated using

- 285 maximum likelihood (ML) techniques while approximating the marginal likelihood using the
- Laplace approximation (Skaug and Fournier, 2006), and efficiently identifying the ML estimates
- using automatic differentiation (Fournier et al., 2012) as implemented using the TMB package in

288 R (Kristensen et al., 2016). Further details regarding VAST can be found elsewhere (Thorson
289 2019).

290	Factor-analysis models generally require some constraints on loadings matrix $\lambda_f(t)$ to ensure
291	that the model is identifiable. We follow previous practice in fixing $\lambda_f(t) = 0$ for all $f > t$
292	(Thorson et al., 2015b; Zuur et al., 2003), but then rotate results to ensure that they are
293	interpretable similarly to principle components analysis (PCA). In particular, we define a
294	rotation matrix R such that AR has columns identical to the eigenvectors of $\Lambda^t \Lambda$, and then define
295	AR as the climate indices and $\mathbf{RX}(c)$ as the map for each category <i>c</i> . We specifically use a
296	"PCA rotation" (Thorson, 2019b; Thorson et al., 2016a), which maximizes the variance for each
297	axis in sequential order, and this differs from varimax rotation which would instead associate
298	each mode of variability with a minimal subset of species.
299	Validating EOF application using sea surface temperature
300	Having defined a statistical model that generalizes a conventional EOF analysis, we seek to
301	validate that it yields estimates that are similar to an EOF analysis. We therefore download
302	monthly average sea surface temperature for every 2° by 2° grid cell within 20° - 60° N and 132° -
303	250°W for every month from Jan. 1950 through Jan. 2018 from the NOAA Extended
304	Reconstructed Sea Surface Temperature (v.5) product (Huang et al., 2017). We analyze these
305	data with Eq. 3a-3b using three indices, and hypothesize that the 1 st index will represent spatial
306	differences in average temperature, the 2 nd will be correlated with the Pacific Decadal Oscillation
307	(PDO), and the 3 rd will be correlated with the North Pacific Gyre Oscillation (NPGO). However,
308	we note that small differences with the PDO will likely arise because we are analyzing only
309	January SST, while the conventional PDO index is calculated from monthly SST measurements.
310	We also note that differences are likely to be greater for the NPGO because the conventional

NPGO is calculated from sea surface height (SSH), and while SST and SSH are often correlated
they are not guaranteed to be perfectly correlated.

313 Estimating an ecosystem index from bottom-trawl data in the eastern Bering Sea

We also seek to demonstrate the benefits of estimating oceanographic indices directly from multispecies biomass-sampling data. To do so, we download bottom trawl sampling data from the eastern Bering Sea bottom trawl survey (Lauth and Conner, 2016) for fourteen commonly occurring bottom-associated fish and crab species. We then fit Eq. 4a-4b to these data while again estimating three oceanographic indices. We hypothesize that the 1st will represent speciesspecific habitat preferences that are stable over time, and do not have strong *a priori* hypotheses regarding the remaining two indices.

321 **Results**

322 Validating EOF application using sea surface temperature

Visualization of the three dominant indices fitted to January average sea surface temperature 323 (Fig. 1) confirms that the spatio-temporal model uses the 1st axis to represent persistent spatial 324 differences in temperature, and the 2nd axis accurately reproduces the annual PDO index with a 325 correlation of 0.92 with the PDO calculated as the EOF of these data. The spatial map associated 326 327 with the PDO index confirms that years with a positive PDO phase have warmer temperatures inshore off the coast of North America and cooler temperatures in the central North Pacific. 328 Similarly, the 3^{rd} index is correlated with the NPGO, although the correlation (0.52) is 329 substantially weaker than for the PDO. The map associated with this index is orthogonal to the 330 PDO map, and shows that a positive phase for the NPGO is associated with decreased water 331 332 temperatures in the California Current from California southward, and elevated temperatures in the northern Bering Sea. The maps for PDO and NPGO are correlated with published estimates 333

and, as expected, the annually varying intercepts ($\beta(t)$ in Eq. 3a) show a gradual increase over time representing climate-forcing of surface temperatures in the Pacific Ocean.

336 Estimating an ecosystem index from bottom-trawl data in the eastern Bering Sea

337 Visualizing the three indices estimated from bottom-trawl survey data for fourteen bottom-

associated species in the eastern Bering Sea (Fig. 2A) shows that these multispecies indices

partition temporal variability into stable (1^{st} index), multi-decadal trend (2^{nd} index), or

340 interannual variability (3rd index). As expected, stable habitat preferences (the 1st axis) explains

the majority (87%) of total (spatial and spatio-temporal) covariation, while long-term shifts (2^{nd})

axis) explains more (9%) than the axis of interannual variability (3^{rd} axis; 4%). For comparison,

we also calculate the proportion of temporal covariation explained by each factor; this confirms
that the first axis explains only 10% of temporal while axis 2 and 3 both explain considerably
more (63% and 28%, respectively).

Visualizing a "habitat preference" index for each species (Fig. 2B, left column) shows 346 347 expected relationships where, for example, snow crab has a more northward distribution than Tanner crab, arrowtooth flounder has elevated density in the outer domain in the southwestern 348 349 edge of the survey, and Pacific cod has relatively weak preferences relative to other species. 350 Variation among years in the magnitude of this index (i.e. a higher value in 1995 than 1998) presumably indicates years where species' distribution is more (1995) or less (1998) in-line with 351 352 their long-term habitat preferences. The "multi-decadal trends" index shows a decrease over the 353 37 years analyzed here, and its spatial map shows, for example, that it is capturing the long-term 354 decrease in density for Alaska skate and arrowtooth in the outer domain relative to the middle 355 domain, as well as a long-term decrease in Tanner crab near Bristol Bay (Fig. 2B middle 356 column). Finally, the "interannual variability" index shows 1-3 year periods of positive or

357 negative stanzas from 1982-1998, then changing to 6-8 year stanzas from 1999-2017. A negative phase for this index (Fig. 2B right column) is associated e.g., with increased density of 358 Alaska pollock and Pacific cod on the northern boundary of the eastern Bering Sea survey and 359 decreased density of Tanner crab in this same area (see Appendix 3, Fig. S1 for patterns for other 360 species). These three indices collectively capture several well-documented trends in the eastern 361 362 Bering Sea, for example, the long-term shift in distribution for arrowtooth from the outer to middle domain (Fig. 3 bottom row). In particular, these indices capture increased density for 363 Alaska pollock and Tanner crab in the northern portion of the survey in 2017 relative to 1982-364 365 1996. Similar patterns are also captured when fitting a model that includes independent spatial and spatio-temporal variation for every species (Appendix 3, Fig. S2), so we conclude that these 366 patterns are not artefacts of the EOF model structure but instead are a low-rank representation of 367 patterns found in the bottom-trawl survey data. Finally, we conclude by noting that that the 368 "interannual variability" index is highly correlated (0.74) with the area of the eastern Bering Sea 369 with summer bottom temperatures less than 2° C (Fig. 4), despite the model not fitting to any 370 physical data, where the small standard errors for this estimated index (black whiskers in Fig. 4) 371 suggest that the pattern is generated by signal within available data rather than statistical error 372 373 introduced by noisy sampling data.

374 **Discussion**

In this paper, we have shown that a multivariate spatio-temporal model can be used to apply empirical orthogonal function analysis to multispecies samples of population densities. EOF analysis is widely used to summarize physical processes (e.g., sea surface temperature) but is less-often used to summarize biological processes due to the noisy, zero-inflated, and multivariate nature of common biological sampling. Multispecies marine resource surveys are

routinely collected in standardized designs in many marine ecosystems worldwide, including
Europe, North America, East Asia, Southern Africa, and Oceana, and the dominant axes of
variability for ecosystem dynamics in these systems can now be summarized using the publicly
available software package, *VAST*.

We have also demonstrated our approach using two case studies: temperature in the North 384 385 Pacific and biomass-samples for bottom-associated fishes and decapods in the eastern Bering Sea. In both examples, the dominant axis represents persistent spatial variation in a given 386 physical or biological variable, and this axis explains 87% (for biological) and 99% (for 387 388 physical) variance. This portion of variance is not typically quantified during EOF analysis, which is often applied to anomalies and therefore long-term patterns are eliminated prior to 389 analysis. We retain this portion of variance (rather than centering data prior to analysis) so that 390 the same algorithm can be applied to biological sampling data that are best described using a 391 delta-model or other "dust-bunny" distributions (McCune and Root, 2015; Thorson, 2018). The 392 2nd axis then represents longer-term spatial trends in the biological case-study, in this case 393 represented in part by the shift in distribution of several flatfishes from offshore to onshore areas. 394 Additional trends may be driven by east-west movements (e.g., great sculpin). The distribution 395 396 shift for arrowtooth flounder has been previously documented (Spencer et al., 2016; Thorson et al., 2016b), and has been attributed to both the extent of the cold pool (Kotwicki and Lauth, 397 398 2013) and an interaction between the bottom temperature and population abundance (Ciannelli et al., 2012). The 3rd axis, by contrast, corresponds to the widely documented role of sea-ice and 399 cold-pool extent in the Bering Sea (Hunt et al., 2011; Stabeno et al., 2012). It is comforting that 400 401 an EOF analysis of biological variables is able to recover this well-documented, bottom-up 402 driver of ecological dynamics. However, the fact that the "cold pool" axis explains less variance

than the "multi-decadal trends" axis underlines the importance of future studies to identifyingbottom-up or top-down drivers of long-term spatial redistribution of flatfishes in this region.

We also note a paradox regarding the 3rd axis in the eastern Bering Sea. This axis has an 405 index that closely corresponds to the extent of the cold pool, and fluctuations in cold-pool extent 406 are primarily driven by changes in bottom temperature in the northern portion of the EBS 407 408 (Appendix 3, Fig. S3) although some cold years (e.g., 1999, 2006-2010, 2012) have large enough cold pool extent that they then decrease bottom temperatures in the southern EBS. However, the 409 species-specific maps associated with the 3rd axis of variability resemble the map of bottom 410 411 temperatures (Appendix 3, Fig. S3) for some species but not others. In particular, the spatial areas associated with changing bottom temperatures overlap with the 3rd axis maps for several 412 species (e.g., Pacific cod. walleye pollock, Tanner crab) but not others (e.g., yellowfin sole, 413 Alaska plaice, great sculpin). These latter species appear to show changes in spatial distribution 414 between years with large and small cold-pool extent, but their distribution does not necessarily 415 expand as a function of local bottom temperatures. This supports previous research showing that 416 local bottom temperature in isolation is not sufficient to explain or predict future distribution for 417 many EBS species (Litzow, 2017; Thorson et al., 2017b; Thorson, 2019a). In fact, it suggests 418 419 that distribution for these species may be driven by a mechanism that is correlated with fluctuations in cold-pool extent, but not necessarily bottom temperature itself. This could 420 421 include changes in the timing of the plankton bloom as driven by timing of ice melt and 422 indirectly related to bottom temperatures (Hunt et al., 2011; Sigler et al., 2014), the timing of ontogenic movement from nearshore to offshore habitats (Nichol et al., 2019) or the opening of 423 424 thermal gateways across the middle shelf which allow species located in the outer shelf to move 425 toward the inner shelf (Ciannelli and Bailey, 2005). These and other potential mechanisms could

426 result in a spatial distribution that differs between warm and cold years (and hence is correlated with the cold-pool extent), but where distribution in those years is not necessarily correlated with 427 local bottom temperatures. Alternatively, spatio-temporal dynamics that are in phase with the 428 429 extent of the cold pool index may also be driven by a change in catchability of the bottom trawl sampling gear. The bottom trawl only samples the lower few meters of the water column during 430 431 a fixed summer season, and catchability may vary among years either due to variation in gear performance or spatial availability (von Szalay and Somerton, 2005; Nichol et al., 2019). These 432 many processes could result in species catch rates that covary with sea-ice extent but not with 433 434 localized temperatures, necessitating a flexible approach to modelling region-scale drivers of distribution for these species. Relatedly, these species vary widely in generation time and life-435 history strategy, such that it is surprising that the majority show a strong response to cold-pool 436 extent (i.e., the 3rd factor). Despite these differences in life-history, our results suggest that these 437 various bottom-associated species all shift their spatial distribution in response to behavioral 438 439 cues, prey availability, or other mechanism that are strongly correlated with cold-pool extent. A common approach for examining the effect of large-scale climate fluctuations on the 440 biological productivity of marine organisms is to reduce complex physical dynamics to one or 441 442 more indices and then correlate these indices with measures of biological production (Stenseth et al., 2003). This approach has contributed to characterizing patterns of climate-productivity 443 444 covariation in the ocean, but it is challenged by the evolving nature of these relationships over 445 time and space (Litzow et al., 2018). Analysts could also use the statistical generalization of EOF analysis to analyze the link between biology and physics in two new ways. First, as shown 446 447 here, estimated modes of biological variation can be compared with relevant measurements of 448 physical variables across scales, to provide some interpretation of what physical processes are

449 correlated with leading biological variation. Linking physics and biology in this way has some benefits for interpreting biological modes of variation, while still visualizing the residual 450 biological variation that is not otherwise explained. Importantly, an estimated mode of 451 biological variation may be correlated with a known physical process during one portion of the 452 time-series and not another, and this can be used to indicate when the linkage between these 453 454 physical and biological processes is nonstationary. Second, biological indices could be extracted and used directly in a secondary modelling framework, e.g., as an annual index in a population-455 dynamics model (e.g., Schirripa et al., 2009). This second approach allows modes of biological 456 457 variation to be propagated even when the physical mechanisms driving variation are poorly understood, and therefore represents ecosystem drivers without directly modelling them. 458 In the application to groundfish data, EOF analysis decomposes biological variation into 459 stable patterns (1st mode), multi-decadal trends (2nd mode) and interannual variability (3rd mode). 460 The scale order of these indices may not always be the same, depending on how variability at 461 each scale contributes to the total variability of the biological data set. Similarly, other data sets 462 may lack either longer-term trends or strong internanual variability. Nevertheless, decomposing 463 biological variability across temporal scales may enable a more mechanistic understanding of the 464 465 effect of climate on biological productivity (Drinkwater et al., 2010; Ottersen et al., 2010). The generalization that we propose here allows applications to data that are not necessarily 466 467 normally distributed and can handle the zero inflation via application of a delta model. The error 468 structure is flexible to allow temporally correlated measurements and non-stationarity (see Thorson (2019b) for a full description of features available in package VAST). The generalization 469 470 presented here can be used to simultaneously detect patterns of covariation across space and time 471 in multispecies assemblages, and therefore overcome these problems. This ability to provide

simultaneous inference across taxa can provide a mechanistic understanding of bottom up and
top down dynamics affecting species abundance/productivity (Cury et al., 2008). We envision
future biological oceanography applications for the analyses of meroplankton (McClatchie et al.,
2018), remote sensing products for characterization of biogeochemical and ecological oceanic
boundaries (Kavanaugh et al., 2014), and zooplankton collections (Colebrook, 1978), in addition
to the application demonstrated using survey data for groundfishes.

We see three primary avenues to extend this work, which we recommend for future research: 478 1. Spatially varying coefficient models to forecast distribution shifts: Our analysis for the 479 480 eastern Bering Sea suggests that many species have a strong but nonlocal response to regional temperatures, as shown by the similarity between the 3rd axis and cold-pool extent. 481 These indices can be included in species distribution modes using a apatially-varying 482 coefficient (Bacheler et al., 2009; Bartolino et al., 2011), and recent research corroborates 483 that a spatially varying response to cold-pool extent improves forecast skill relative to local 484 temperature in isolation (Thorson, In press). We hypothesize that annual indices such as 485 cold-pool extent will be informative about low-frequency variability in species distribution 486 that is currently lacking in long-term forecasts of distribution shift in response to climate. 487 488 2. Joint rank-reduction models of physical drivers and biological responses: An interesting extension of the generalized EOF analysis used here is to include both physical and 489 biological variables, to make inference on the shared covariance among them and the spatial 490 491 patterns in which this covariance is dominant (Brown and Fiechter, 2012). The signal of anthropogenic climate change is emerging from the envelope of natural variability in many 492 493 ecosystems globally (Henson et al., 2017), including the Bering Sea (Walsh et al., 2018). 494 This anthropogenic change makes nonstationary behavior in environmental variables

495 (changing mean or variance) an increasingly important factor in ecosystem dynamics (Burrows et al., 2011). The spatio-temporal generalization of EOF analysis presented here 496 may be expanded to include both physical and biological variables, to make inferences about 497 shared covariance and to directly infer the role of nonstationary physical dynamics in leading 498 ecosystem patterns (Brown and Fiechter, 2012). In particular, the estimated linkage between 499 500 physical and biological variables could be useful to estimate the spatial map of physical conditions that is most strongly associated with variation in a given biological variable. In 501 essence, this could be used to identify the physical index that maximizes predictive power for 502 503 a given species.

3. Model evaluation via null models and simulation testing: Finally, we recommend future 504 research regarding the expected performance of applying EOF to multivariate biological 505 samples. Likely model performance could be explored via simulation-testing, where fits to 506 data from multiple ecosystems could be used to simulation new data (potentially with 507 different sampling designs or sample sizes) and the model could then be re-fitted to explore 508 measure model performance. The performance of multivariate spatio-temporal models when 509 estimating "factor-loadings" (analogous to EOF indices in this analysis) and associated 510 511 spatial maps has been simulation-tested previously (e.g., Thorson et al., 2016a, 2017a, In press), but performance specifically when generalizing EOF analysis remains a useful topic 512 for further testing. Alternatively, researchers could adapt previously developed "null models" 513 514 (e.g., Planque and Arneberg, 2018) to determine whether the statistical approach to EOF is likely to identify erroneous patterns from autocorrelated noise. 515 516 Another issue raised by anthropogenic climate change is the creation of novel patterns of

517 covariance among different ecologically important physical variables, which may produce

518	nonstationary statistical relationships between environmental drivers and ecological responses, or
519	nonstationary relationships among species (Litzow et al., 2018; Williams and Jackson, 2007;
520	Wolkovich et al., 2014). This possibility is not accounted for in the current study, as our
521	approach assumes stationary relationships among species. We therefore recommend future
522	methodological research to incorporate time-evolving community relationships in our modeling
523	approach in order to examine the potential role of nonstationary community relationships under
524	climate change.
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528	available from the Alaska Fisheries Science Center at
529	http://www.afsc.noaa.gov/RACE/groundfish/survey_data/data.htm. The R package VAST is
530	publicly available online at https://github.com/James-Thorson/VAST/, and can be used to
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Fig. 1 – Visualization of three dominant axes of variability for sea surface temperature in the North Pacific (rows), where each axis includes an index (left columns) showing the magnitude and phase (y-axis) for each year (x-axis), as well as a spatial map (right column) showing the variability in temperature expected during a positive phase, where each map has a spatial variance of 1.0. For each axis we list the proportion of variance explained (top-right of each panel) as well as the proportion of temporal variance explained (calculated after subtracting off the mean across years for each index; listed in parentheses). The second axis is highly correlated with the PDO and the third axis is correlated with the NPGO, so in each case we show the estimated index (black line) relative to the published PDO or NPGO index (red line) and list the correlation between estimated and published indices (top-left of

each panel).

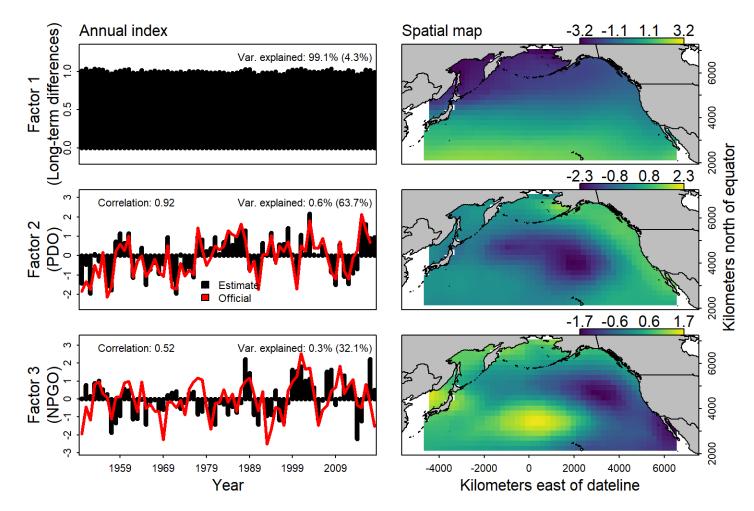
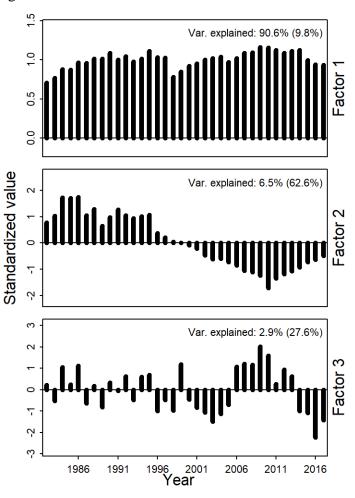


Fig. 2 – Visualization of three dominant axes of variability for bottom trawl sampling of biomass 791 for fourteen bottom-associated fish and crab species in the Eastern Bering Sea, where each axis 792 793 includes an index (Fig. 2a) showing the magnitude and phase (y-axis) for each year (x-axis), as well as a spatial map (Fig. 2b) showing the variability in biomass expected during a positive 794 phase for seven of the fourteen modeled species (rows) and each of the three indices (columns), 795 where each map has a spatial variance of 1.0. For each axis we list the proportion of variance 796 explained (top of Fig. 2a), as well as the proportion of temporal variance explained (calculated 797 after subtracting off the mean across years for each index; parentheses in Fig. 2a). For spatial 798 maps associated with each axis of variability for the remaining seven species, see Fig. S1. 799

800 Fig. 2A



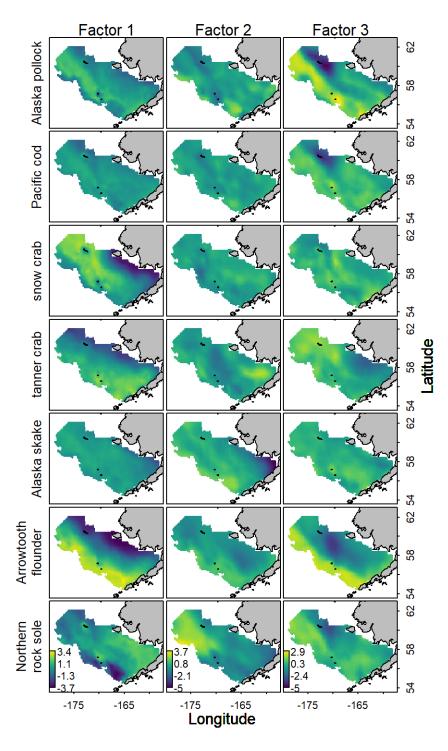


Fig. 3 – Visualization of predicted biomass for three of the fourteen modeled species (rows) in six of the thirty-six modeled years.
 Species are selected from those shown in Fig. 2B, while years are evenly spaced from 1982-2017; for comparison with a model that
 includes independent spatial and spatio-temporal for each species see Appendix 3 Fig. S2.

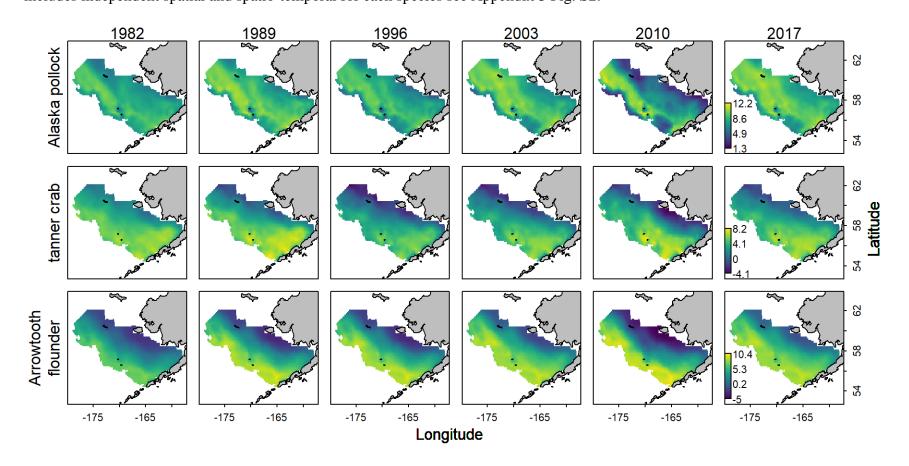


Figure 4 – Comparison of the 3^{rd} estimated index (black bullets) as well ±1.96 times the 810 estimated standard error in each year (black whiskers), as calculated using Monte Carlo 811 simulation of the estimated loadings matrix $\lambda_f(t)$ and the inverse-Hessian matrix of fixed 812 effects, applied to bottom trawl sampling of biomass for fourteen bottom-associated fishes and 813 814 crabs in the Eastern Bering Sea (black line; left y-axis label) and the cold pool area (blue line; 815 right y-axis label). The correlation between estimated and published indices is listed (top-left of the plot). 816

