# Defining indices of ecosystem variability using biological samples of fish communities: a generalization of empirical orthogonal functions 

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Keywords: Empirical orthogonal function; sea-ice extent; eastern Bering Sea; bottom trawl; vector autoregressive spatio-temporal model


#### Abstract

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Multivariate data reduction techniques are widely used to describe modes of variability in atmospheric and oceanographic conditions for the world's oceans. Dominant modes of 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 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 compared with biological conditions to describe or predict physical drivers of ecological 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 of biological response associated with each index. We therefore develop a generalization of EOF analysis that can be applied directly to multispecies biological samples using a multivariate spatio-temporal model. These biologically derived indices can then be compared with relevant 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 North Pacific Gyre Oscillation. We then identify three axes of variability in the eastern Bering 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 second represents a multi-decadal trend in distribution for most species; for example, showing an 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 stanzas from 1999-2017 and is highly correlated (0.87) with the extent of the cold bottom temperatures in this region and associated impacts on Alaska pollock and Pacific cod. These


axes represent ecological dynamics for adult fishes and therefore integrate the impact of bottomup and top-down processes, and they also confirm the importance of cold-pool extent for fish distribution in the Bering Sea while visualizing its varied impact on individual species.

Moreover, this spatio-temporal approach allows oceanographers to define annual indices representing modes of variability in diverse biological communities from widely available fieldsampling data.

## Introduction

Oceanographers are confronted with a tremendous challenge in summarizing multivariate physical and biological processes into a tractable number of dominant patterns, which can then be readily communicated to other fields (e.g., fisheries scientists), used as covariates in climate models, or used in many other ways. One approach to this challenge is to identify dominant axes 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. Familiar examples include the Pacific Decadal Oscillation (PDO; Mantua et al., 1997) and North 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 al., 2010), and also tend to capture variability in a suite of cross-correlated, ecologically important processes (Stenseth et al., 2003), making them valuable tools for summarizing climate effects on marine biota. For example, the importance of the PDO was established via 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 concentrations in southern California (NPGO; Di Lorenzo et al., 2008). These ocean climate 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 broadly use the term "EOF analysis" to refer to the algorithm generating both the temporal index 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 the temporal index).

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 of data reduction techniques applied to biological data involve either the aggregation across space and time so to obtain regional averages of multi-species biomass (e.g., PCA applied to annual indices of species biomass, Planque and Arneberg (2018)), or aggregation across species 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 the scope of ecological inference, sacrificing either small-scale spatiotemporal dynamics or the 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 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 represented using a normal distribution; (2) biological samples often arise from spatially unbalanced sampling, which complicates the creation of spatial maps in unsampled areas or when fitting to spatially unbalanced sampling programs; and (3) biological sampling is often available for multiple species, and therefore EOF analysis must be generalized for multivariate data sets, where each variable has both temporal and spatial expression.

Despite the profusion of research using EOFs to characterize physical oceanographic conditions, it is not always clear how to relate these indices to biological processes. This commonly involves a multi-step workflow, where physical variability is first summarized with EOF analysis, and the resulting index is then used as a covariate in a separate statistical model linking physical oceanography to biological or social outcomes. Unfortunately, many published relationships between physical conditions and biological responses in ocean environments either
break down over time or have poor skill when forecasting (Myers, 1998; Thorson, 2019a). We note two ways that this multi-step workflow may contribute to poor out-of-sample predictive skill for statistical physics-biology relationships. First, climate indices derived from EOF analysis summarize physical dynamics across a large spatial domain. These regional indices are then correlated with local conditions, and biological responses are then correlated with local 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 change over time, thus changing their correlation with local biological responses (Litzow et al., 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 secondmost, etc. However, these definitions are fixed in time while different axes of physical variation 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 the age-structure of a population (Hsieh et al., 2006).

Given these difficulties when using oceanographic indices of physical habitat to describe biological responses, we see a useful role for indices estimated directly from multivariate 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; Latimer et al., 2009; Ovaskainen et al., 2017). For example, spatial factor analysis (Thorson et 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 arise from both bottom-up and top-down processes. We propose that these approaches would be 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 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 then demonstrate that this approach generalizes conventional EOF analysis by replicating 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 the eastern Bering Sea. Many studies in this system have linked the spatial distribution of fishes 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 be independent of cold-pool extent for some species (Mueter and Litzow, 2008). Similarly, our method estimates an index of ecosystem variability that is highly correlated with the spatial extent of cold near-bottom waters (termed the "cold pool"), and therefore corroborates the 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.

## Methods

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

Climatologists, meteorologists, and oceanographers have used empirical orthogonal functions to characterize dominant modes of variability in physical ocean conditions for over fifty years. 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. Kidson (1975a) introduced a similar method using principal components analysis (PCA) applied to monthly average precipitation and sea surface pressure to similarly show that a reduced set of variables could explain $>75 \%$ of the original variance. These studies established the common practice wherein a multivariate statistical techniques (PCA or factor analysis) is applied to 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 the original process.

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 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. (1997) used EOF to define the PDO as the dominant mode of variability in sea surface temperature (SST) in the North Pacific, and the second mode was further explored and interpreted by Bond et al. (2003) and subsequently called the "Victoria mode" (Ding et al., 2015). Similarly, Di Lorenzo et al. (2008) defined the NPGO as the second mode of variability for sea surface height (SSH) anomalies, and the NPGO is correlated with the Victoria mode via the high correlation between SST and SSH.

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).

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 dynamics. This Kalman-filter interpretation of EOF was subsequently extended by Wikle and Cressie (1999), who introduced a descriptive spatial process wherein the physical process is 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., Lindgren et al., 2011), and now are also widely used in ecology and fisheries science (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 community) while accounting for variable dynamics across both space and time.

## General approach

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

1. a spatial map $X_{f, c}(s)$ representing whether a given location $s$ has a positive or negative value (as well as the magnitude of that value) during a "positive phase" of the index; and
2. a time-series $\lambda_{f}(t)$ indicating whether a given time $t$ has a positive phase $\left(\lambda_{f}(t)>0\right)$ or negative phase $\left(\lambda_{f}(t)<0\right)$, as well as whether a given year has a weak magnitude $\left(\lambda_{f}(t)\right.$
within the $25 \%$ and $75 \%$ quantiles for $\lambda_{f}$ ) or a strong magnitude ( $\lambda_{f}(t)$ outside the $25 \%$ and 75\% quantiles).

Importantly, the map $X_{f, c}(s)$ associated with each time series $\lambda_{f}(t)$ differs for each category $c$ of $n_{c}$ modeled categories, and this allows our time series to represent multivariate data, for 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 $\tilde{\mathcal{Y}}_{c, t}(s)$ that includes the net effect of all estimated oceanographic indices. Each predictor variable is a linear combination of $n_{f}$ oceanographic indices:

$$
\begin{equation*}
g(\underbrace{\tilde{\mathcal{Y}}_{c, t}(s)}_{\text {predictor variable }})=\underbrace{\beta_{c, t}}_{\text {intercepts }}+\underbrace{\sum_{f=1}^{n_{f}} \overbrace{\lambda_{f}(t)}^{\text {Index phase }} \overbrace{X_{f, c}(s)}^{\text {Index map }}}_{\text {Net effect of indices }} \tag{1}
\end{equation*}
$$

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:

$$
\begin{equation*}
\mathcal{Y}_{c, t}(s) \sim h\left(\widetilde{\mathcal{Y}_{c, t}}(s), \ldots\right) \tag{2}
\end{equation*}
$$

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.

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

We first seek to show that this approach can generalize the EOF analysis that is widely used in physical oceanography to generate indices. Published studies have implemented EOF using
either principle components analysis (PCA) or factor analysis (FA), and both involve a matrix $Y_{t}(s)$ of measurements of a physical variable at each location and time. Anomalies from longterm 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 are used to calculate the sample covariance $V_{t, t}$ for anomalies between any pair of times. The analyst then applies an eigendecomposition (for PCA) or minimization algorithm (FA) to identify a set of $n_{f}$ orthogonal axes of covariation, ranked from most important to least important, such that the first few axes explain the majority of covariation. These axes are then treated as the index $\lambda_{f}(t)$, and the spatial map $x_{f}(s)$ associated with each index is calculated as either the correlation or regression of $\lambda_{f}(t)$ and $y_{f}(s, t)$.

We replicate a univariate EOF analysis by simplifying the general model in four ways:

1. Eliminate notation for multiple categories such that it is applied to a single response;
2. Use an identity link function;
3. Use a normal distribution for observations; and
4. Replace functions with matrices, i.e., $X_{f, c}(s)$ with a matrix $\mathbf{X}$ representing the predicted value of each oceanographic variable $x^{*}\left(s_{i}, f\right)$ at the location for each sample $i$ for each index, and $\tilde{\mathcal{Y}}_{c, t}(s)$ with $\tilde{y}_{i}$ representing the prediction for that sample $i$.

These three changes result in the following model:

$$
\begin{gather*}
\tilde{y}_{i}=\beta\left(t_{i}\right)+\sum_{f=1}^{n_{f}} \lambda_{f}\left(t_{i}\right) x^{*}\left(s_{i}, f\right) \\
y_{i} \sim \operatorname{Normal}\left(\tilde{y}_{i}, \sigma^{2}\right)
\end{gather*}
$$

where $\sigma^{2}$ is the variance of measurement errors, which is minimized by explaining variation in the response $y(s, t)$ to estimated indices, and $x^{*}\left(s_{i}, f\right)$ 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
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 from conventional EOF analysis (which does center the data to calculate anomalies prior to analysis), and this difference means that the $2^{\text {nd }}$ axis from the generalized model is similar to the $1^{\text {st }}$ axis from conventional EOF, the $3^{\text {rd }}$ axis is similar to the $2^{\text {nd }}$ from conventional EOF, etc. However, centering the data does not extend to other common forms of data analysis (e.g., generalized linear models) so not centering the data is important for subsequent generalizations. Also differing from conventional EOF, we estimate a separate intercept $\beta_{t}$ for every year, so that this term captures interannual variability in $y(s, t)$ among years (i.e., the increasing trend in surface temperatures due to climate forcing). We estimate a separate intercept for each year to 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 constant across years (to be more similar with conventional EOF analysis). We then compare 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 http://www.o3d.org/npgo/).

## Extension to multi-species biomass-sampling data

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;
2. Use a Poisson-link delta-model (Thorson 2017) where the linear predictor is associated with numbers density $n$, where encounter probability $p_{i}$ for sample $i$ is derived as a complementary log-log link from numbers density $p_{i}=1-\exp \left(-n_{i}\right)$ and expected biomass when encountered $r_{i}$ is defined such that it is proportional to numbers density, $r_{i}=$ $\frac{n_{i}}{p_{i}} w\left(c_{i}, t_{i}\right)$, where $w\left(c_{i}, t_{i}\right)$ is the biomass per individual, which is estimated separately for each category and year; and
3. Minimize the negative log-likelihood for encounters and sampled biomass for each species. These changes result in the following model:

$$
\begin{gather*}
\log \left(n_{i}\right)=\beta\left(c_{i}, t_{i}\right)+\sum_{f=1}^{n_{f}} \lambda_{f}\left(t_{i}\right) x^{*}\left(s_{i}, c_{i}, f\right) \\
\operatorname{Pr}\left(B=b\left(s_{i}, c_{i}, t_{i}\right)\right)=\left\{\begin{array}{cl}
1-p_{i} & \text { if } B=0 \\
\operatorname{Lognormal}\left(\log \left(r_{i}\right), \sigma_{t}^{2}\right) & \text { if } B>0
\end{array}\right.
\end{gather*}
$$

4a

Where $\beta(c, t)$ is again an intercept for every category and time, Eq. 4b represents a delta-model distribution where $1-p_{i}$ is the probability mass associated with $B=0, \operatorname{Lognormal}(a, b)$ is a lognormal probability density function with logmean $a$ and log-variance $b$, and $\sigma_{t}^{2}$ is the residual estimated log-variance in positive catch rates. We specify a separate intercept for every category and time because, in our experience, total abundance for marine fishes often varies substantially among years due endogenous biological processes (e.g., variable production of juveniles; Thorson et al. (2015a)) and the annually varying intercept "controls" for this variation such that 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 $\lambda_{f}\left(t_{i}\right) x^{*}\left(s_{i}, c_{i}, f\right)$ corresponds to an approximately $1 \%$ increase in expected biomass. This relative scale implies that all coefficients representing spatial and temporal variation (i.e., the right-hand side of Eq. 4a) are dimensionless.

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) multiple response variables. These difficulties are addressed by (1) applying a delta-model with a linear predictor that is shared among years and species; (2) projecting the linear predictor from $n_{s}$ 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 map representing the impact $x^{*}(s, c, f)$ of a positive phase for each of $n_{c}$ modeled variables at any location $s$ within a defined spatial domain. In particular, accounting for spatially misaligned 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.

## Parameter estimation

Parameters for both configurations of the general model can be estimated using a publicly 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-ThorsonNOAA/VAST). This package estimates spatial variables $x(s, c, f)$ for $n_{s}$ "knots" as random effects following a Gaussian Markov random field, uses a predictive process formulation to interpolate the value of $x^{*}\left(s_{i}, c_{i}, f\right)$ for the location $s_{i}$ and category $c_{i}$ of sample $i$ given $x(s, c, f)$, 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 Matérn correlation function (see Appendix 2 for details). Fixed effects are then estimated using 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

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

Factor-analysis models generally require some constraints on loadings matrix $\lambda_{f}(t)$ to ensure that the model is identifiable. We follow previous practice in fixing $\lambda_{f}(t)=0$ for all $f>t$ (Thorson et al., 2015b; Zuur et al., 2003), but then rotate results to ensure that they are interpretable similarly to principle components analysis (PCA). In particular, we define a rotation matrix $\mathbf{R}$ such that $\boldsymbol{\Lambda} \mathbf{R}$ has columns identical to the eigenvectors of $\boldsymbol{\Lambda}^{t} \boldsymbol{\Lambda}$, and then define $\Lambda \mathbf{R}$ as the climate indices and $\mathbf{R X}(c)$ as the map for each category $c$. We specifically use a "PCA rotation" (Thorson, 2019b; Thorson et al., 2016a), which maximizes the variance for each axis in sequential order, and this differs from varimax rotation which would instead associate each mode of variability with a minimal subset of species.

## Validating EOF application using sea surface temperature

Having defined a statistical model that generalizes a conventional EOF analysis, we seek to validate that it yields estimates that are similar to an EOF analysis. We therefore download monthly average sea surface temperature for every $2^{\circ}$ by $2^{\circ}$ grid cell within $20^{\circ}-60^{\circ} \mathrm{N}$ and $132^{\circ}-$ $250^{\circ} \mathrm{W}$ for every month from Jan. 1950 through Jan. 2018 from the NOAA Extended Reconstructed Sea Surface Temperature (v.5) product (Huang et al., 2017). We analyze these data with Eq. $3 \mathrm{a}-3 \mathrm{~b}$ using three indices, and hypothesize that the $1^{\text {st }}$ index will represent spatial differences in average temperature, the $2^{\text {nd }}$ will be correlated with the Pacific Decadal Oscillation (PDO), and the $3^{\text {rd }}$ will be correlated with the North Pacific Gyre Oscillation (NPGO). However, we note that small differences with the PDO will likely arise because we are analyzing only January SST, while the conventional PDO index is calculated from monthly SST measurements. 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.

## 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. $4 \mathrm{a}-4 \mathrm{~b}$ to these data while again estimating three oceanographic indices. We hypothesize that the $1^{\text {st }}$ will represent speciesspecific habitat preferences that are stable over time, and do not have strong a priori hypotheses regarding the remaining two indices.

## Results

## Validating EOF application using sea surface temperature

Visualization of the three dominant indices fitted to January average sea surface temperature (Fig. 1) confirms that the spatio-temporal model uses the $1^{\text {st }}$ axis to represent persistent spatial differences in temperature, and the $2^{\text {nd }}$ axis accurately reproduces the annual PDO index with a correlation of 0.92 with the PDO calculated as the EOF of these data. The spatial map associated 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. Similarly, the $3^{\text {rd }}$ index is correlated with the NPGO, although the correlation (0.52) is substantially weaker than for the PDO. The map associated with this index is orthogonal to the PDO map, and shows that a positive phase for the NPGO is associated with decreased water 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
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. Estimating an ecosystem index from bottom-trawl data in the eastern Bering Sea Visualizing the three indices estimated from bottom-trawl survey data for fourteen bottomassociated species in the eastern Bering Sea (Fig. 2A) shows that these multispecies indices partition temporal variability into stable ( $1^{\text {st }}$ index), multi-decadal trend ( $2^{\text {nd }}$ index), or interannual variability ( $3^{\text {rd }}$ index). As expected, stable habitat preferences (the $1^{\text {st }}$ axis) explains the majority ( $87 \%$ ) of total (spatial and spatio-temporal) covariation, while long-term shifts (2 ${ }^{\text {nd }}$ axis) explains more ( $9 \%$ ) than the axis of interannual variability ( $3^{\text {rd }}$ 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 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 edge of the survey, and Pacific cod has relatively weak preferences relative to other species. 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 their long-term habitat preferences. The "multi-decadal trends" index shows a decrease over the 37 years analyzed here, and its spatial map shows, for example, that it is capturing the long-term decrease in density for Alaska skate and arrowtooth in the outer domain relative to the middle domain, as well as a long-term decrease in Tanner crab near Bristol Bay (Fig. 2B middle column). Finally, the "interannual variability" index shows 1-3 year periods of positive or
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 Alaska pollock and Pacific cod on the northern boundary of the eastern Bering Sea survey and decreased density of Tanner crab in this same area (see Appendix 3, Fig. S1 for patterns for other species). These three indices collectively capture several well-documented trends in the eastern 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 Alaska pollock and Tanner crab in the northern portion of the survey in 2017 relative to 19821996. 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 patterns are not artefacts of the EOF model structure but instead are a low-rank representation of patterns found in the bottom-trawl survey data. Finally, we conclude by noting that that the "interannual variability" index is highly correlated (0.74) with the area of the eastern Bering Sea with summer bottom temperatures less than $2^{\circ} \mathrm{C}$ (Fig. 4), despite the model not fitting to any physical data, where the small standard errors for this estimated index (black whiskers in Fig. 4) suggest that the pattern is generated by signal within available data rather than statistical error introduced by noisy sampling data.

## 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 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 physical or biological variable, and this axis explains $87 \%$ (for biological) and $99 \%$ (for 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 analysis. We retain this portion of variance (rather than centering data prior to analysis) so that the same algorithm can be applied to biological sampling data that are best described using a delta-model or other "dust-bunny" distributions (McCune and Root, 2015; Thorson, 2018). The $2^{\text {nd }}$ axis then represents longer-term spatial trends in the biological case-study, in this case represented in part by the shift in distribution of several flatfishes from offshore to onshore areas. Additional trends may be driven by east-west movements (e.g., great sculpin). The distribution 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, 2013) and an interaction between the bottom temperature and population abundance (Ciannelli et al., 2012). The $3^{\text {rd }}$ axis, by contrast, corresponds to the widely documented role of sea-ice and cold-pool extent in the Bering Sea (Hunt et al., 2011; Stabeno et al., 2012). It is comforting that an EOF analysis of biological variables is able to recover this well-documented, bottom-up 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 identifying bottom-up or top-down drivers of long-term spatial redistribution of flatfishes in this region.

We also note a paradox regarding the $3^{\text {rd }}$ axis in the eastern Bering Sea. This axis has an index that closely corresponds to the extent of the cold pool, and fluctuations in cold-pool extent are primarily driven by changes in bottom temperature in the northern portion of the EBS (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 species-specific maps associated with the $3^{\text {rd }}$ axis of variability resemble the map of bottom temperatures (Appendix 3, Fig. S3) for some species but not others. In particular, the spatial areas associated with changing bottom temperatures overlap with the $3^{\text {rd }}$ axis maps for several species (e.g., Pacific cod. walleye pollock, Tanner crab) but not others (e.g., yellowfin sole, Alaska plaice, great sculpin). These latter species appear to show changes in spatial distribution between years with large and small cold-pool extent, but their distribution does not necessarily expand as a function of local bottom temperatures. This supports previous research showing that local bottom temperature in isolation is not sufficient to explain or predict future distribution for many EBS species (Litzow, 2017; Thorson et al., 2017b; Thorson, 2019a). In fact, it suggests 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 include changes in the timing of the plankton bloom as driven by timing of ice melt and 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 thermal gateways across the middle shelf which allow species located in the outer shelf to move toward the inner shelf (Ciannelli and Bailey, 2005). These and other potential mechanisms could
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 local bottom temperatures. Alternatively, spatio-temporal dynamics that are in phase with the 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 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 many processes could result in species catch rates that covary with sea-ice extent but not with 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 lifehistory strategy, such that it is surprising that the majority show a strong response to cold-pool extent (i.e., the $3^{\text {rd }}$ factor). Despite these differences in life-history, our results suggest that these various bottom-associated species all shift their spatial distribution in response to behavioral 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 biological productivity of marine organisms is to reduce complex physical dynamics to one or 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 covariation in the ocean, but it is challenged by the evolving nature of these relationships over 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 here, estimated modes of biological variation can be compared with relevant measurements of physical variables across scales, to provide some interpretation of what physical processes are
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 biological variation that is not otherwise explained. Importantly, an estimated mode of biological variation may be correlated with a known physical process during one portion of the time-series and not another, and this can be used to indicate when the linkage between these 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 populationdynamics model (e.g., Schirripa et al., 2009). This second approach allows modes of biological variation to be propagated even when the physical mechanisms driving variation are poorly understood, and therefore represents ecosystem drivers without directly modelling them.

In the application to groundfish data, EOF analysis decomposes biological variation into stable patterns ( $1^{\text {st }}$ mode), multi-decadal trends ( $2^{\text {nd }}$ mode) and interannual variability ( $3^{\text {rd }}$ mode). The scale order of these indices may not always be the same, depending on how variability at each scale contributes to the total variability of the biological data set. Similarly, other data sets may lack either longer-term trends or strong internanual variability. Nevertheless, decomposing biological variability across temporal scales may enable a more mechanistic understanding of the 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 normally distributed and can handle the zero inflation via application of a delta model. The error 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 presented here can be used to simultaneously detect patterns of covariation across space and time 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:

1. Spatially varying coefficient models to forecast distribution shifts: Our analysis for the eastern Bering Sea suggests that many species have a strong but nonlocal response to regional temperatures, as shown by the similarity between the $3^{\text {rd }}$ axis and cold-pool extent. These indices can be included in species distribution modes using a apatially-varying coefficient (Bacheler et al., 2009; Bartolino et al., 2011), and recent research corroborates that a spatially varying response to cold-pool extent improves forecast skill relative to local temperature in isolation (Thorson, In press). We hypothesize that annual indices such as cold-pool extent will be informative about low-frequency variability in species distribution that is currently lacking in long-term forecasts of distribution shift in response to climate.
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 biological variables, to make inference on the shared covariance among them and the spatial 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 ecosystems globally (Henson et al., 2017), including the Bering Sea (Walsh et al., 2018). This anthropogenic change makes nonstationary behavior in environmental variables
(changing mean or variance) an increasingly important factor in ecosystem dynamics (Burrows et al., 2011). The spatio-temporal generalization of EOF analysis presented here may be expanded to include both physical and biological variables, to make inferences about shared covariance and to directly infer the role of nonstationary physical dynamics in leading ecosystem patterns (Brown and Fiechter, 2012). In particular, the estimated linkage between 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 essence, this could be used to identify the physical index that maximizes predictive power for a given species.
3. Model evaluation via null models and simulation testing: Finally, we recommend future research regarding the expected performance of applying EOF to multivariate biological samples. Likely model performance could be explored via simulation-testing, where fits to data from multiple ecosystems could be used to simulation new data (potentially with different sampling designs or sample sizes) and the model could then be re-fitted to explore measure model performance. The performance of multivariate spatio-temporal models when estimating "factor-loadings" (analogous to EOF indices in this analysis) and associated 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 for further testing. Alternatively, researchers could adapt previously developed "null models" (e.g., Planque and Arneberg, 2018) to determine whether the statistical approach to EOF is likely to identify erroneous patterns from autocorrelated noise.

Another issue raised by anthropogenic climate change is the creation of novel patterns of covariance among different ecologically important physical variables, which may produce
nonstationary statistical relationships between environmental drivers and ecological responses, or nonstationary relationships among species (Litzow et al., 2018; Williams and Jackson, 2007; Wolkovich et al., 2014). This possibility is not accounted for in the current study, as our approach assumes stationary relationships among species. We therefore recommend future methodological research to incorporate time-evolving community relationships in our modeling approach in order to examine the potential role of nonstationary community relationships under climate change.

## Acknowledgments

NOAA_ERSST_V5 data provided by the NOAA/OAR/ESRL PSD, Boulder, Colorado, USA, from their web site at https://www.esrl.noaa.gov/psd/. Groundfish data used here are publicly available from the Alaska Fisheries Science Center at
http://www.afsc.noaa.gov/RACE/groundfish/survey_data/data.htm. The R package VAST is publicly available online at https://github.com/James-Thorson/VAST/, and can be used to replicate or update this analysis. We thank S. Zador, M. Scheuerell, and two anonymous reviewers for comments on an earlier draft, and the many scientists who contributed to bottom trawl sampling in the eastern and northern Bering Sea shelf.

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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 for fourteen bottom-associated fish and crab species in the Eastern Bering Sea, where each axis 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 phase for seven of the fourteen modeled species (rows) and each of the three indices (columns), where each map has a spatial variance of 1.0. For each axis we list the proportion of variance explained (top of Fig. 2a), as well as the proportion of temporal variance explained (calculated after subtracting off the mean across years for each index; parentheses in Fig. 2a). For spatial maps associated with each axis of variability for the remaining seven species, see Fig. S1.

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


