**Bottom up and top down effects of temperature on body growth, population size spectra and yield – an application of a multi-species size-spectrum model**

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**Abstract**

Resolving the combined effect of climate warming and exploitation in a food web context is key for predicting future biomass production, size-structure and potential yields of marine fishes. Previous studies based on mechanistic size-based models have found that bottom-up processes are important drivers of size-structure and fisheries yield at the ecosystem level in changing climates. However, we know less about the joint effects bottom up and top down effects (direct of warming on individual-level physiology) on exploited food webs. Using the Baltic Sea as a case study, we assess how various food web metrics are affected by warming and exploitation in a species-resolved size-based food web. We parameterize a dynamic size-spectrum model representing the offshore Baltic Sea food web, and investigate how individual growth rates, size-structure, relative abundances of species and yields are affected by warming under different scenarios of fishing mortality and physiological scaling. We find that the model is able to reproduce realistic biomass densities in the relatively stable time period (1992-2002). Evaluation of steady-state properties at simulated higher temperatures suggests that both warming leads to positive impacts on growth of younger fish within species, but the positive effect levels off and becomes negative for older cod within projected warming scenarios. Any key explanation why? The effect of warming on the abundance size-spectrum, spawning stock biomass and yields of the cod? fishery depend more strongly on the effect of temperature on basal resources than on ??. Results from our model can be used to qualitatively evaluate how warming and exploitation interactively affect the Baltic Sea fish community, which can inform about potential future yields in a multispecies scenario and thus support strategic management decisions.

**Introduction**

Climate change affects aquatic food webs directly by shifting the distribution of species (Pinsky *et al.* 2013), abundance (McCauley *et al.* 2015), body size (Daufresne *et al.* 2009; Baudron *et al.* 2014), and ecosystem function (Pontavice *et al.* n.d.), which can lead to indirect effects through the rewiring of food webs (Frainer *et al.* 2017). Global retrospective analysis of warming and fish population dynamics has revealed that declines in productivity of scientifically assessed fish populations across ecoregions has already declined due to climate change (Free *et al.* 2019). Interestingly, these findings are also matched, both in magnitude and direction, by projections from an ensemble of mechanistic ecosystem models, which predicts further declines in biomass by 2100 especially at higher trophic levels (Lotze *et al.* 2019). Large scale models are crucial for informing global assessments about the implications of warming affecting key ecosystem-level processes (Lotze *et al.* 2019). However, by necessity, these models rely on simplifying assumptions (Fulton *et al.* 2019), which may lead to challenges in translating predictions across scales (Brander *et al.* 2013). Thus, there is a need to increase our understanding of climate change impacts on individuals and how that affects regional dynamics (Blanchard *et al.* 2012).

The primary effect of temperature is on individual physiology, and mechanistic models should therefore be based on physiological principles (Lefevre *et al.* 2018; Neubauer & Andersen 2019). There is however no consensus or a single framework that accurately describes the mechanistic basis for the empirical patters behind the temperature-size rule (Atkinson 1994; Ohlberger 2013; Audzijonyte *et al.* 2018), which predicts faster growth rates but smaller asymptotic body size. Attempts have been made to describe this phenomena from a supply and demand perspective, using a generalization of the von Bertalanffy growth equation (VBGE) (Perrin 1995; Cheung *et al.* 2013). Specifically, it has been suggested that increasing oxygen limitation with warming due to elevated metabolic demands impose growth constraints that lead to smaller asymptotic body masses. This model (and other similar growth models) has been criticized for not accounting for the energetic costs of reproduction (Marshall & White 2019a, b), and for not representing the underlying physiological processes accurately – both in definitions and parameters (Lefevre *et al.* 2018). Importantly, evaluating changes in food web size-structure based on static growth models (as in no food-dependence) alone is not ideal, as it has long been known that the physiological performance, of e.g. growth, depends on an interplay between body size, temperature and food availability (Brett *et al.* 1969). For instance, the optimum temperature for growth declines with reduced food rations (Huey & Kingsolver 2019), which means that the effects of temperature are mediated by changes in resources levels. Linking ecology and physiology in a trait based setting can be a fruitful venue for overcoming too simplistic physiological explanations while maintain a high degree of generality (Neubauer & Andersen 2019). Thus, models should be based on physiological principles (Neubauer & Andersen 2019) and evaluated in an ecological context to account for the food dependence of growth.

Process-based food web models aim represent the dynamics or structure of ecosystems based on an understanding of the key mechanisms behind the phenomena considered, and models can therefore vary in their performance depending on the process they are tailored for (Spence *et al.* 2018). It is therefore important to both identify the driving mechanism and to capture it in the model (Curtsdotter *et al.* 2019). Traditional Lotka-Volterra-type models describing pairwise species interactions often fail to accurately represent the life history of fish, which in general grow several orders of magnitude in body size and change feeding niches and trophic levels during ontogeny (Werner & Gilliam 1984). Hence, it is often necessary to account for intraspecific variation in body size and allometry to capture the ecological function of a typical fish (Hartvig *et al.* 2011). Not surprisingly there have been pioneering work in fish and fisheries ecology that aim to capture processes such as ontogenetic growth and size-dependent feeding interactions, for example in physiologically structured models (PSPM) (Metz & Diekmann 1986; De Roos & Persson 2001). The development of trait based size spectrum models allowed for more generic formulations of physiologically based life history description and using body weight at maturation as a distinguishing trait (Hartvig *et al.* 2011), which has provided general insights and mechanisms on e.g. trophic cascades (Andersen & Pedersen 2009). More recently, extensions of the size spectrum framework for modelling size-based food webs include more precise characterization of species, e.g. by defining species feeding preference and life history characteristics – so called multi-species size spectrum models (MSSMs) (Blanchard *et al.* 2014; Scott *et al.* 2014). Such species- and trait-based models have led to a closer connection to observational data, by e.g. both informing and validating model predictions. For instance, (Reum *et al.* 2019), by characterizing species predation preference by both species- and body size, they were able to more accurately describe the true feeding links, which in addition weakened the effects of fishing on trophic cascades. Phoebe here! + MSSMs therefore provide a good framework for assessing the impacts of climate on growth by accounting for physiological description of life history and a bioenergetic perspective.

The Baltic Sea

Research questions: How do bottom up and top down (direct effect of warming on individual-level physiology) affect growth rates, population size-structure and fisheries yield of a harvested food web? Here we apply temperature dependence to key physiological rates describing the bioenergetic life history of fish, based on meta-analysis to overcome unclear parameters and estimates. We also include uncertaininty in theses parameters as that is a very common things. Brief summary of paper and findings. Relate to similar/large scale/ensemble models, e.g. 5%

**Materials and Methods**

*Food web*

In this study we develop a Multi-species size spectrum model (MSSM) inspired by a simplified version of the food web in the Baltic proper (ICES sub divisions 25-29+32, Appendix Fig SX). This food web is here characterized by three fish species: cod (*Gadus morhua*), sprat (*Sprattus sprattus*) and herring (*Clupea harengus*), and this choice is based on data availability (they all have analytical stock assessments), they are the most important species commercially in this sub-region and they constitute the main species in terms of biomass. We also model one pelagic and one benthic background resource spectra.

*Size-spectrum model*

Our model is based on source code for the multi-species implementation of size-spectrum models in the ‘R’-package *mizer* (Blanchard *et al.* 2014; Scott *et al.* 2014, 2019; R Core Team 2018), which we extend by adding multiple background resources and temperature-scaling of key physiological processes and background resource levels. All model code (parameterization, calibration and analysis), together with necessary data is available on GitHub (<https://github.com/maxlindmark/mizer-rewiring/tree/rewire-temp/baltic>), and will be deposited on Zenodo (<http://doi.org/>) upon publication. In this section we describe the key elements of the MSSM, which mostly follow the standard multispecies mizer model (Scott *et al.* 2014).

In MSSMs, individuals are characterized by their weight () and species identity (). The core equation is the McKendrik-von Foerster equation, which here describes the change in abundance at size through time from food dependent somatic growth and mortality, based on bioenergetic principles:

where [] is somatic growth (dependent on the availability of background resources or prey) and [] is total mortality. The total mortality, , consists of three components: constant allometric background mortality, , predation mortality (all that is eaten by other species, (Hartvig *et al.* 2011)) and fishing mortality ).

The proportion of encountered food that is consumed, which can be allocated to growth and/or reproduction, follows a Holling functional response type II:

where is the *feeding level* (0:1), is the allometric maximum consumption rate and is the encountered food. The amount of encountered food for a predator of body weight is given by the available food in the system, scaled by a size-selection function and the search rate of the predator. Here, available food, , is the integral of all species and background resource

where is the weight of prey, is the preference of species on resource , and indicates species. Note that in contrast to other MSSMs (Blanchard *et al.* 2014; Reum *et al.* 2019) we have a species-preference for the background resources to account for species feeding differently on benthic and pelagic resource, but assume no preference in terms of the species interactions. This is for simplicity, as the body size-only prey selection and encounter captures the important predation interactions in this system and the species largely occupy similar spatial areas in the time period considered here (as inferred from independent stomach data, see model calibration, Appendix S1). The body size-based selection function, , is given by a log-normal selection function (Ursin 1967):

where parameters and are the preferred predator-prey mass ratio and the standard deviation of the log-normal distribution. Together, the size-based selection function and the amount of available prey is multiplied with the allometric function describing the search rate, to get the actual encountered food.

.

Encountered food is assimilated with efficiency and used to cover basal metabolic costs, which scale allometrically as . The remaining energy, , is allocated to growth or reproduction. This allocation to reproduction () increases smoothly from 0 at the weight maturation, , to 1 at the asymptotic weight, , according to the function:

which has the property that the growth rate, i.e.

approximates a von Bertalanffy growth curve when the feeding level is constant (Hartvig *et al.* 2011). Reproduction is given by the total egg-production in numbers, which is the integral of the energy allocated to reproduction multiplied by an efficiency factor () divided by the egg weight, :

The recruitment of offspring to each species related to the total egg production via a Beverton-Holt stock recruit relationship, such that recruitment reaches a maximum as the egg production increases:

where is the maximum recruitment of each species. is here treated as a free parameter that is found in the calibration process by minimizing the residual sum of squares between spawning stock biomass from stock assessment and the MSSM. In doing so, it also ensures that the species coexist (Hartvig *et al.* 2011).

The temporal dynamics of the background resource () spectra (benthic or pelagic) is defined as:

, where is the population regeneration rate, is the population carrying capacity and is predation mortality, and lambda is defined as .

*Temperature dependence*

Describe the projected temperature increase in the Baltic

* Here we need a good justification about why we assume certain ways that temperature will affect processes. Why maximum intake and not search rate? Why mortality? Any references? I know we decided that last year, but we will need to back it up with refs
* Motivation for which rates are temperature dependent and how (MTE and meta-analysis in another thesis chapter)
* This equation describes a general size-and temperature-dependent rate, based on a generalized version of the MTE equation. Here it applies to maximum intake rate, metabolism and mortality.

|  |  |
| --- | --- |
|  | (1) |

* For resource parameters, we only scale the allometric functions describing growth rate and carrying capacity, so no temp-size-interaction here.

|  |  |
| --- | --- |
|  | (2) |

* The random activation energies on both physiological rates and resource parameters allows us to look into the importance of bottom up vs top down effects of warming

Max, sorry, but I am not sure I follow this. I guess more description in the methods is needed. Initially I thought you are using a few values of activation energies, but now it seems you are using lots of random draws. So are you running multiple simulations for a scenario? Are all E values changing independently? Are you exploring each effect separately or are all five E rates sampled randomly and then all applied? How many simulations? This seems like a great approach to assess uncertainty, but I just need a bit more info to follow it

*Model calibration and analyses*

The Baltic Sea offshore food web has experienced dramatic structure changes over the last decades, which makes it difficult to identify a period of relative stability for model calibration. We chose to calibrate our model to the time period 1992-2002, as in (Jacobsen *et al.* 2017), which is a post-regime shift period characterized by high fishing mortality (F) on cod, low cod and herring abundance and high sprat abundance (Gårdmark *et al.* 2015). The cutoff at 2002 also ensures that we do not calibrate the model to years when the physiological state of cod started to drastically deteriorate, i.e. mid 2000’s and onwards when the growth capacity, condition, proportion of large fish in the population and reproductive capacity started to decline (Svedäng & Hornborg 2014; Mion *et al.* 2018; Neuenfeldt *et al.* 2019).

We calibrated the system to the Baltic Sea by finding the vector of -values that minimize the residual sum of square between the natural log of spawning stock biomass (SSB) from stock assessment output and modelled SSB, using the "L-BFGS-B" algorithm in the ‘R’-optimization function ‘*optim’*. Mean SSB and F during the calibration time period (1992-2002) from stock assessment were acquired from ICES Working Group Reports (ICES 2013, 2015), and are in unit 106 kg/area. This area corresponds to ICES subdivisions 25-29+32, which equals approximately 2.49E+11 m2 (estimated using ICES shapefiles in ArcGIS). To express the modelled SSB in the same scale, we first defined it in the units g/m2 and then used the converting factor 240.342. Given that the North Sea is approximately 1E11 m2, we used the and divided by 1E11 as starting values, before calibrating the model. This resulted in a starting value for and of 1 (Blanchard *et al.* 2014). We also ensured that also growth-rates were realistic by comparing them to VBGE curves fitted to trawl survey data, and that the density dependence due to the stock-recruit function was appropriate. This was assessed from the ratio of recruitment before and after stock-recruitment density dependence was applied (Jacobsen *et al.* 2017). Lastly, we compared time series of model output simulated together with historical temperature and fishing effort data with stock assessment output. A detailed description of the step-by-step calibration protocol used in this study, inspired by (Jacobsen *et al.* 2017), can be found in Appendix S1.

Table 1 Parameters of distributions describing activation energies of temperature-dependent rates in the size-spectrum model. See figure 2 for 200 random draws from these distributions which were used as input in projections.

|  |  |  |  |
| --- | --- | --- | --- |
| Symbol | Rate | Distribution | Source |
|  | Metabolism |  | Lindmark et al. (*in prep*) |
|  | Background mortality |  | (Brown *et al.* 2004); Lindmark et al. (*in prep*) |
|  | Maximum consumption |  | Lindmark et al. (*in prep*) |
|  | Background resource carrying capacity |  | (Gilbert *et al.* 2014); (Savage *et al.* 2004) |
|  | Background resource growth rate |  | (Savage *et al.* 2004) |

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Figure 1. Time series of (A) effort by species and (B) temperature from the regional coupled model system RCA4-NEMO using the RCP 8.5 scenario. In panel A, the light grey area corresponds to years with from stock assessment. The dark grey area is the calibration time period. We treat the 60 years prior to the start of the F-time series as burn-in and keep the constant and equal to the first time series (in year 1974). The future projections correspond to the means of the multispecies stock assessments’ estimates and the size-spectrum model. In panel B, temperature is shown as deviations from the average in 1970-1999. Note the different x-axis of the figures.

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Figure 2. 200 random samples from the distributions describing the activation energies of the rates resource carrying capacity and resource growth rates [assumed same for benthic and pelagic background resources], metabolic rate, background mortality rates and maximum consumption rates. See (Table 1). Combinations of these parameters where used in model projections to understand how uncertainty in these parameters affect individual- and population level metrics. The rate-scalars from the activation energies, using Equations 1-2, can be seen in Figure S10.

**Results and discussion**

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*Figure 3. Individual growth trajectories of sprat, herring and cod from model projections. Top row shows size-at-age in 2050 assuming fishing mortality held at mean FMSY levels (Figure 1A) from different stock assessment estimates and this model (Fig. S7) and a temperature increase in the Baltic Sea in line with the projection from the RCA4-NEMO assuming the RCP 8.5 emission scenario (Figure 1B). The dashed line in the top row depicts projections assuming no scaling of physiological rates and are therefore equivalent to constant temperatures. In both the green and pink projections physiological rates are temperature dependent, and in the pink projections also background resources are temperature-dependent via their growth rates and carrying capacities. The shaded area corresponds to minimum and maximum size at age (or relative size at age) that stem from the random parameterization of individual-level scalars. For fit of size-at-age to data, see Figure S4.*

* With the random combinations activation energies that we believe are more of less realistic, and for scenarios with/without effects of temperature on the background resources, body growth rates increase with warming.
* The positive effect of warming diminishes with age.
* For herring and sprat, warming always has a positive effect on growth given the temperature increase we model (which is about +1.5C, see Figure S11 for the scenario here) – even in the most “unfortunate” parameter combination.
* For cod, the “confidence bands” go below 1 in some combinations of parameters, with and without resource temperature-dependence, suggesting warming has a negative effect on size at age even with a 1.5C increase in temperature.
* These results show that the effect of warming on growth rates does not depend that much on how growth of background resources vary with temperature, as the two lines follow each other and largely overlap.
* I also note that the distributions of activation energies are relatively tight, which makes the “confidence bands” pretty tight. This is because it’s not very likely that we draw one really good activation energy for intake and one really bad for metabolic rate at the same time. This is also why I plot the minimum and maximum rather than some quantile.
* (Suggested sensitivity analysis: redo this analysis in a constant temperature and effort setting (not time-varying), and explore extremes of activation energies to get a better feeling for what the effects are. Will also show which the actual activation energies are that give the min, max and mean response that correspond to the ribbons.)

A close up of a map

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*Figure 4. Abundance-at-size (abundance spectrum) of sprat, herring and cod in 2050 relative to projection with constant temperature after 2007 (see Figure S10) assuming different fishing mortalities, here implemented as scalars to average FMSY from stock assessments and size-spectrum models. The top row corresponds to no temperature increase, hence the pink line which shows the average FMSY scenario is at y=1. The middle and bottom rows show scenarios without and with temperature depended resource dynamics, respectively.*

* The rate of fishing changes the abundance spectrum quite drastically - higher fishing means removal of large fish and more biomass of medium sized fish, where’s low fishing rates increase the abundance of large fish.
* The qualitative and quantitative effect of warming on the size spectra depends on how the resources are temperature-dependent (more so than the growth rates do).
  + If growth rates do not increase with temperature and the carrying capacity is unchanged, warming leads to a lower abundance-at-size for all species given the same fishing effort – especially for the large fish (the pink lines increase initially but fall down after a certain size along the size-axis).
  + If resource growth rates increase with temperature abundance increases for medium sized fish but decreases for large and small fish.

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*Figure 5. Spawning stock biomass (SSB) and fishery yield (assuming knife edge selectivity at maturation size) under constant temperatures at steady state. Cold = , i.e. no temperature effects, and warm is +2.*

* The rate of fishing changes the abundance spectrum quite drastically - higher fishing means removal of large fish and more biomass of medium sized fish, where’s low fishing rates increase the abundance of large fish.

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*Figure 6. Fisheries yield relative to a scenario with constant temperature () and average FMSY. The x-axis shows the scaling factor to and the y-axis is a scalar to the average FMSY from model and stock assessment.*

* I need to look into this more…
  + Remember how I used the overall average FMSY in the projections? (See comment on Figure 1). That is not very good I think because cod SSM-FMSY is below the assessment-FMSY, but the other two are above. So the difference between species here is likely due to the scalar putting the species away from their SSM-FMSY in slightly different directions.
  + For this figure, I will simply just use the model FMSY to make the cross-species comparison relevant. I will also do it with 2\*3 plot (probably), corresponding to models with and without temperature dependence on resource dynamics…
* BUT! What seems to come out is that warming reduces yield. This is despite it enhancing growth rates (Figure 3), and is likely connected to the reductions in SSB and abundance-at-most-sizes (especially larger ones or above maturation size which is the selection cut off.)
* Note also that there might be some variation between the time-varying temperature and effort scenarios and the constant ones. I’m not entirely sure yet but will look into it more. Guessing it has something to do with starting values. Anyway, I don’t think I’m comfortable comparing this too much with Figures 3-4 (which are from time-varying projections), because of this. So I will maybe redo this figure based on time-varying projections.

*Extra analysis*

* Mean body size of populations (variations on Fig. 3)
* Look at mortality similar to how we look at growth (we have growth and size spectra already but need to show mortality as well).
* Plot not only relative size-spectra but the actual size-spectra. Can also dig out equivalent plots from data as a means of validating…
* What I initially wanted to look into also:
  + See if the effect of fishing on e.g. abundance spectrum, depends on temperature (have not gotten this far)
  + What is the effect of selectivity here? It has been suggested that selectivity has increased density dependence by truncating the size-distribution… Have not thought too much about this however.

*Preliminary conclusions*

* Top down and bottom up effects of warming both shape food webs.
* An increase in temperature with 1.5C positively affects size-at-age for all species, but the positive effect diminishes with age and often becomes negative for cod aged 5+. This result is not very sensitive to temperature-driven changes in background resources.
* The effect of temperature on resources (bottom up) partly determines whether or not abundance-at-size increases or decreases with warming

**Discussion**

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**Author contributions**

The code was first developed from mizer (Scott *et al.* 2019) code by AA, all authors contributed to tailoring it for the purpose of this study. ML conceived the idea. All authors contributed to study design. ML performed analysis and wrote the first draft. All authors contributed to writing the paper and to revisions.

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