



# Arctic Sensitivity? Suitable Habitat for Benthic Taxa Is Surprisingly Robust to Climate Change

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Arctic marine ecosystems are often assumed to be highly vulnerable to ongoing climate change, and are expected to undergo significant shifts in structure and function. Community shifts in benthic fauna are likely to result from changes in key physico-chemical drivers, such as ocean warming, but there is little ecological data on most Arctic species to support any specific predictions as to how vulnerable they are, or how future communities may be structured. We used a species distribution modeling approach (MaxEnt) to project changes over the 21st century in suitable habitat area for different species of benthic fauna by combining presence observations from the OBIS database with environmental data from a coupled climate-ocean model (SINMOD). Projected mean % habitat losses over taxonomic groups were small (0–11%), and no significant differences were found between Arctic, boreal, or Arcto-boreal groups, or between calcifying and non-calcifying groups. However, suitable habitat areas for 14 of 78 taxa were projected a change by over 20%, and several of these taxa are characteristic and/or habitat-forming fauna on some Arctic shelves, suggesting a potential for significant ecosystem impacts. These results highlight the weakness of general statements regarding vulnerability of taxa on biogeographic or presumed physiological grounds, and suggest that more basic biological data on Arctic taxa are needed for improved projections of ecosystem responses to climate change.

**Keywords:** benthic invertebrates, climate warming, multiple stressors, ocean acidification, species-distribution modeling

## INTRODUCTION

Ongoing climate change affects multiple environmental drivers, with direct and indirect implications for marine ecosystems. For example, reductions in sea-ice volume and extent, increasing air and sea-surface temperatures, and freshwater and biogeochemical discharge in the Arctic have already been noted to be occurring at rates higher than the global average (IPCC, 2014; AMAP, 2017) with consequences for regional ocean acidification rates (Bates et al., 2012; Bellerby, 2017). Anthropogenic impacts from fossil fuel CO<sub>2</sub> emissions are linked with changes in these drivers and have led to changes in ocean chemistry. These impacts include reductions in both

ocean pH and carbonate mineral saturation states, which have been implicated in decreased performance of many marine taxa (Pörtner, 2008; Kroeker et al., 2011). Models project that ongoing ocean acidification (OA) will continue over the coming decades, and the degree of OA will be dependent mainly on social and policy decisions (e.g., Bellerby et al., 2014, 2018). The amplification of both warming and acidification observed in the Arctic, in combination with other expected physical and biological impacts, has led to the common viewpoint that Arctic ecosystems are particularly vulnerable to on-going climate changes (Smetacek and Nicol, 2005; Renaud et al., 2007; Brierley and Kingsford, 2009; Doney et al., 2012; Hoppe et al., 2018). Changes in community composition that may result are not merely of academic interest: they may reflect underlying functional responses with broad-reaching implications for ecosystem processes and services.

Laboratory experiments and field sampling have generated numerous but often contradictory predictions of how changes in climate-change-related drivers will affect marine life, from individuals to ecosystem processes. Physiology, phenology and life-history, community structure, and system productivity (Pörtner et al., 2005; Ardyna et al., 2014; Renaud et al., 2015, 2018) are just several examples of impacts that may be felt: from the very basic processes of life to the provisioning of ecosystem services. Until now, however, most studies have focused on effects of single factors, and usually in laboratory or isolated field settings (Pedersen et al., 2016; AMAP, 2018). Recent syntheses have called for investigations of broader thematic and spatial scope, including studying multiple factors in combination for arriving at more realistic predictions (e.g., AMAP, 2018), and adopting a pan-Arctic perspective when identifying trends in ecosystem response (Wassmann, 2015). These perspectives are needed for developing mitigation, adaptation, and management strategies for marine ecosystems across the Arctic (Halpern et al., 2012). Whereas some Arctic taxa are clearly threatened by changing climatic conditions (Kovacs et al., 2011; Doney et al., 2012; Grebmeier, 2012; Beaugrand and Kirby, 2018), a multi-factor investigation of vulnerability of Arctic taxa in general has not been performed.

One of the most intuitive changes that could be expected due to climate warming is the poleward expansion of boreal organisms. This has already been observed for both benthic and pelagic organisms, and is suggested to be largely related to thermal tolerance (e.g., Wetthey and Woodin, 2008; Sunday et al., 2012; Neukermans et al., 2018). Indeed, dominance of boreal and Arctic taxa in seafloor (benthic) communities of the Barents Sea have been shown to fluctuate with temperature in the region over the past 100+ years (Blacker, 1965; Matishov et al., 2012). Niche theory is useful for making predictions as to historical or future distributions in the context of climate change. Single-variable models (thermal-envelope models) have been used to predict, for example, changes in fish distribution with climatic warming (Cheung et al., 2009). In Arctic systems, this technique is problematic due to high uncertainty in either fundamental or realized niches of nearly all taxa in the ecosystem. On-line databases can provide one way around this problem by identifying the characteristics of an organism's

realized niche. This technique has been used to identify benthic taxa that may expand or contract under warming scenarios (Renaud et al., 2015). Although biased by the geographical range of data included in databases, and non-random sampling in general, the method can also provide information for multi-dimensional realized niches. Once current distributions are mapped, a suite of species distribution models may be applied to project future distributions in response to multiple environmental drivers. Such modeling activities have been identified as critical next steps in understanding climate change effects on benthic community structure in Arctic environments (Renaud et al., 2015).

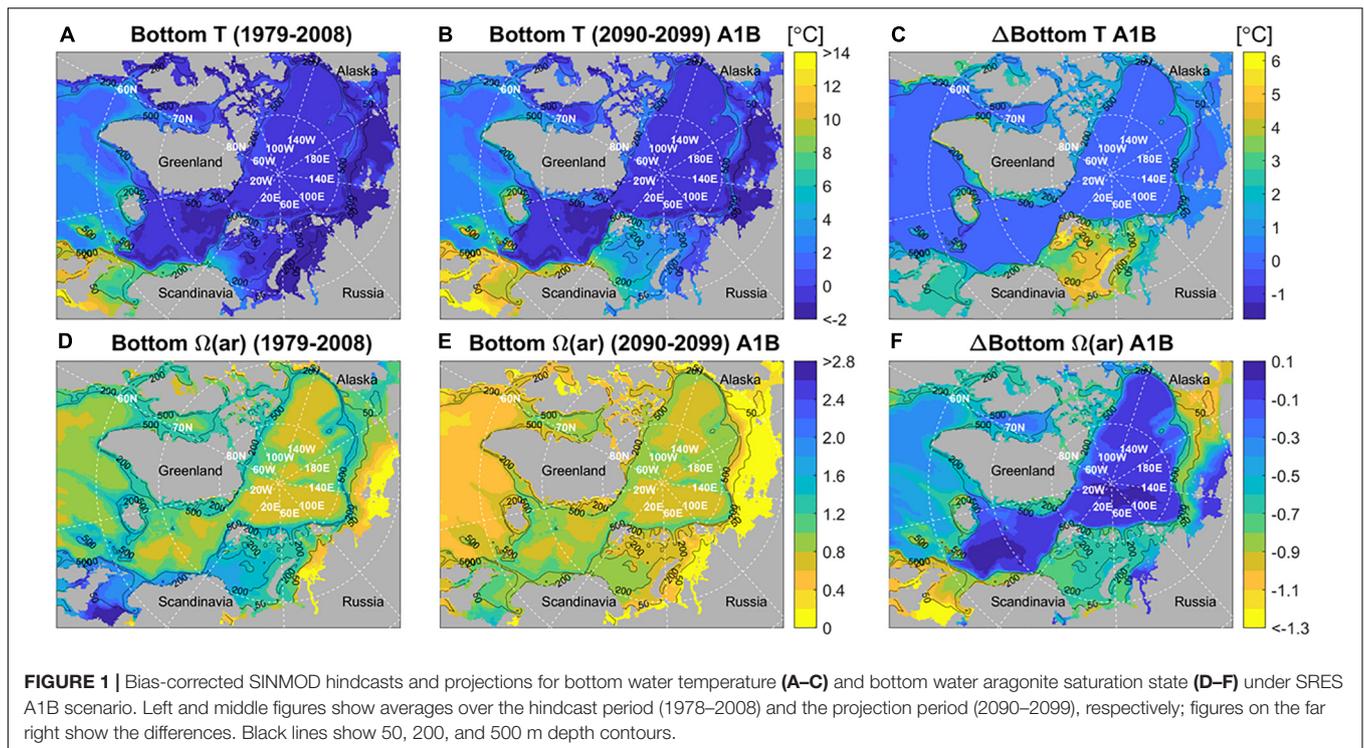
A key requirement for implementing these models is a well resolved and validated hydrographic/ocean chemistry model with realistic climate forcings. This allows for accurate physical characteristics to be assigned to each sampling location in both hindcasting and forecasting modes. One such model has recently been expanded to include the needed spatial and temporal perspectives for the carbonate system, and has been used to upscale results of a mesocosm experiment to other Arctic areas (Bellerby et al., 2012).

This study investigates the vulnerability of benthic fauna found in the Arctic and its marginal seas to end-of-century climate change. We use outputs from a state-of-the-art ecosystem model to drive species-distribution modeling for a broad range of characteristic macro- and mega- benthic taxa characteristic of the Arctic's shelf seas. We analyse results to determine if Arctic taxa are more susceptible to changes in a combination of environmental drivers than boreal or more widespread taxa. We also investigate whether calcification status or higher taxonomic classification (phylum, class) could indicate vulnerability. Implications of the presence or absence of such patterns are discussed.

## MATERIALS AND METHODS

### Environmental Projections

We ran the ocean biogeochemical model SINMOD (Slagstad et al., 2011, 2015), which includes a CO<sub>2</sub> system module (Bellerby et al., 2012), to estimate bottom water temperature and aragonite saturation state ( $\Omega(\text{ar})$ ). The SINMOD domain is pan-Arctic (see **Figure 1**) with 20 km horizontal grid resolution and 25 fixed depth levels, and calculations were based on biweekly saved output. Saturation state describes the seawater concentrations of carbonate and calcium ions relative to the equilibrium concentrations for that mineral; conditions near or below saturation ( $\Omega = 1$ ) are generally considered unfavorable for production or maintenance of normal calcification. SINMOD was first run to produce a hindcast simulation (1979–2008), and then to produce a projection (2001–2099) under the SRES (Special Report on Emissions Scenarios) scenario A1B (see Slagstad et al., 2015; Wallhead et al., 2017, for more details). SRES A1B is a mid-range business-as-usual scenario assuming a “balanced” use of fossil vs. non-fossil energy sources (Nakicenovic et al., 2000) that results in end-of-century global responses that are comparable to the more recent RCP6.0 scenario and



substantially more conservative than the SRES A2 or RCP8.5 scenarios (Collins et al., 2013). SINMOD bottom-water output was corrected using bias estimates, calculated as a function of spatial position (temperature) or of model bathymetry ( $\Omega(\text{ar})$ ) using a compilation of *in situ* observations and matched model output (see Wallhead et al., 2017). Water depth over the SINMOD grid was estimated by interpolating high-resolution bathymetry products (the 500 m resolution IBCAOv3, Jakobsson et al., 2012, where available, otherwise the 2 min ETOPOv2, National Geophysical Data Center, 2006).

## Niche Description and Climatic Change

Occurrence records for 95 benthic taxa characteristic of Arctic shelf infaunal and epifaunal communities were extracted from the Ocean Biogeographic Information System database (OBIS<sup>1</sup>; extracted February 14, 2017; **Supplementary Table S1**). We chose taxa that were common, ecologically relevant (e.g., ecosystem engineers) and/or characteristic taxa of Arctic shelves, and where biogeographic affinities are well-known (see below). In addition, taxa were chosen to cover the range of factors studied here (biogeographic affiliation, calcification status, and higher taxonomic level). Further, we required that each taxon have at least 30 records within the geographic domain of SINMOD in order to maintain MaxEnt model performance and confidence level. This cutoff appeared to be sufficient since the model performed quite well for taxa with close to 30 observations (e.g., *Parastichopus tremulus* and *Serripes groenlandicus*, see **Supplementary Tables S2, S3**). Bottom water temperature and aragonite saturation state from the bias-corrected SINMOD

hindcast output were matched to the georeferenced benthic fauna distributional records by interpolating from the model grid to the faunal data positions and averaging over the 5 years preceding each faunal observation date. Water depth at the sampling points was mostly (61%) from measurements when plausible values were provided; the remaining 39% were estimated using the high-resolution bathymetry products. Each species was assigned status as calcifying or not, biogeographic affinity (Matishov et al., 2012; World Register of Marine Species)<sup>2</sup>, and higher taxonomic grouping (mollusk, echinoderm, polychaete, crustacean, bryozoan).

In locations where species data have been collected systematically, for example through biological monitoring, both presence and absence of species at each site have been recorded. However, most observations of the OBIS database have been collected non-systematically and are available as presence-only records, and the different gear types deployed and study foci make it difficult to infer absence data from these records. We therefore employed a species distribution modeling approach that requires only presence data, in order to maximize the utility of the database.

From the water depth, temperature, and aragonite saturation state associated with each species record, we estimated three-dimensional realized niches and related them to the projected changes in these parameters through the end of the century. The contribution of each environmental variable to species occurrence probabilities in the Arctic was calculated using the MaxEnt method. MaxEnt is a machine-learning algorithm for modeling species distributions from presence-only

<sup>1</sup>www.iobis.com

<sup>2</sup>www.marinespecies.org

species records. In brief, MaxEnt identifies the environmental characteristics of locations with species occurrences that differ from the environmental setting across the whole geographical region of interest. Based on the observed mismatch, a species' suitable habitat is defined. More specifically, the MaxEnt model minimizes the relative entropy between two probability densities (one estimated from the presence data, and one from the landscape) defined in covariate space compared to a uniform distribution null model (see Merow et al., 2013 for details). MaxEnt's predictive performance is consistently competitive with the highest performing methods. Since becoming available in 2004, it has been utilized extensively for finding correlates of species occurrences, mapping current distributions, and predicting to new times and places across many ecological, evolutionary, conservation and biosecurity applications (Elith et al., 2006).

## Sampling Biases

By default, MaxEnt models assume that: (1) all locations on the landscape (or "background locations") were equally likely to be sampled, and (2) any focal species would have been recorded at the sampled landscape points if it had actually occurred there. Assumption (1) is likely violated in our occurrence datasets, because of logistical and practical constraints on sampling in the Arctic (e.g., due to ice cover). Given such spatiotemporal biases, one cannot differentiate whether species are observed in a particular environmental niche because those locations are preferable or because they receive the largest search effort. Therefore, in order to relax assumption (1), landscape points were only drawn from the "target group sampling" (TGS) locations, which we defined as the total set of sampled locations for all species. However, this does not relax assumption (2), which may be violated where sampling gear and study focus have excluded certain species (Phillips et al., 2009). In theory the gear/study focus biases could be accounted for by partitioning the TGS, but this was not considered feasible for our particular dataset due to insufficient samples.

## Model Fitting and Validation

Multicollinearity can be an issue with MaxEnt when answering if and when environmental variables are of ecological interest. Thus, prior to modeling, a correlation analysis was conducted for environmental variables and the final MaxEnt models included variables that were not significantly correlated with each other at  $p < 0.05$  (aragonite saturation state, temperature, water depth). Calcite saturation state was excluded as it correlates highly with aragonite saturation state.

In this study MaxEnt models were fitted as combinations of basic functions and features. MaxEnt had six feature classes: linear, product, quadratic, hinge, threshold, and categorical. Products were all possible pairwise combinations of covariates, allowing simple interactions to be fitted. Threshold features allowed a "step" in the fitted function, hinge features were similar except they allowed a change in gradient of the response. Many threshold or hinge features were fitted for one covariate, giving a potentially complex function. Nevertheless, the MaxEnt program was allowed to simplify the associations between species and the

environment and in many models only one or a few features were used, e.g., hinge, linear, and quadratic.

Segment-based (non-gridded) data were modeled using SWD (samples-with-data) format in MaxEnt for both presence and background sites. A separate model was run for each species. A 10-fold cross-validation was used to obtain out-of-sample estimates of predictive performance and estimates of uncertainty around fitted functions. In order to reduce model overfitting, a balance between accurate prediction (model fit) and generality (model complexity) was sought by maximizing the penalized maximum likelihood function, i.e., the gain function. When doing so, regularization or the LASSO penalty was applied by exploring a range of regularization parameter values and choosing a value that maximizes measures of fit on a cross-validation data set. The LASSO penalty is based on the rationale that features with larger variance should incur a larger penalty and, thus be less likely to be included in the model (Hastie et al., 2009). For model validation a random selection of 25% of the overall localities of species occurrences were used. The percent contributions of individual variables to the final model were identified with jackknife tests. The jackknife test evaluates how each variable contributes to the "gain" of the MaxEnt's model (i.e., improvement in penalized average log likelihood compared to null model) (Elith et al., 2011). We used the raw output (default) as this output is the closest estimate of the probability that the species is present (Elith et al., 2011).

## Model Performance and Predictions

We used the area under the receiver-operating-characteristic curve (AUC) as a single measure of overall model accuracy that is not dependent upon a particular threshold. The value of the AUC is between 0.5 and 1.0 with AUC = 1.0 indicating that the model has a perfect match and AUC = 0.5 indicating that model is no better than random (Fielding and Bell, 1997).

The MaxEnt model was used to predict suitable habitat for each species under current and future environmental conditions, and differences between these predictions were used to assess the response to climate change of potential habitat area for each species. We are aware that projecting future species distributions usually involves extrapolating models to novel combinations of environmental variables, and such projections should be treated with extreme caution. However, a comparison of current and future environmental niche space indicated that only 20% of the whole study area would enter a completely novel environment, as defined by moving beyond the 3D convex hull of the present-day niche space distribution. We therefore consider any artifacts of extrapolation will have only a minor effect on our results.

## Statistical Analyses of Model Results

Change in suitable habitat area from the present (1978–2008) to the future (2090–2099) was calculated from the MaxEnt model for each taxon. One-way analyses of variance (ANOVA) were performed for each of the three factors [calcification (as a *t*-test), biogeography, higher taxonomic level] on the change in habitat area (both percentage and absolute) after testing for homogeneity of variances and normality among factor levels (in each case these results were non-significant). Since single-factor analyses

may obscure interaction effects or even effects of one factor when effects of a second factor dominate the variability, we also ran a 3-factor ANOVA on the data. Since we lacked degrees of freedom to run a fully factorial test, we used a main-effects model. Analyses were performed in Statistica v. 13.

## RESULTS

### Environmental Settings Projections

Projections for bottom water temperature (T) and aragonite saturation state ( $\Omega(\text{ar})$ ) indicate substantial changes over the 21st century (Figure 1). The warmer Atlantic Water-influenced zone expands northward and eastward from its current core in the southern and southwestern Barents Sea (Figures 1A,B), and benthic habitat in the Barents Sea warms by up to 6°C between 1979–2008 and 2090–2099 (Figure 1C). Over the same period, bottom  $\Omega(\text{ar})$  on the shelves (<500 m depth) decreases, mostly by 0.6–1.1 units from around 1–2 (weakly saturated) to 0–1 (undersaturated) (Figures 1D–F). Exceptions include the shallow Russian shelf regions (<50 m depth), which are already strongly undersaturated ( $\Omega(\text{ar}) < 0.4$ ; Figure 1D) and cannot become much more undersaturated in the future (Figures 1E,F). The largest decreases in  $\Omega(\text{ar})$  (~1 unit) are in the North, Bering, and Chukchi Seas, and on the East Siberian Shelf at ~50–500 m (Figure 1F). Arctic water-dominated areas of the northern and eastern Barents/Kara Seas remain relatively stable in temperature (Figure 1C), but experience  $\Omega(\text{ar})$  reductions as large as in the central Barents Sea (Figure 1F). Deeper areas (>500 m) in the Arctic and North Atlantic basins, and in Baffin Bay, see relatively small changes in benthic environment (Figures 1C,F).

### Depicting Realized and Predicted Niches of Benthic Taxa

Of the 78 taxa with sufficient (>30) records for evaluation, the mean change in suitable habitat was small: approximately a 5% loss in suitable habitat based on MaxEnt. The range, however, was high, from a 42% increase in habitat (the amphipod *Byblis gaimardii*) to a 53% decrease (the sea cucumber *P. tremulus*) according to the 3-variable model (Table 1, Supplementary Table S2, and Figures 2, 3). MaxEnt performed very well for most (~70%) taxa, with AUC > 0.9 (Supplementary Table S3). Temperature contributed most to defining the niche space for 72% of the taxa, while depth and aragonite saturation state were the largest contributors for 19 and 8% of the taxa, respectively (Supplementary Table S3).

Surprisingly, both calcifiers (in general and by phylum) and Arctic (compared to Arcto-boreal or Boreal) taxa were found to be resilient to the projected environmental changes in terms of group mean response. Calcifiers were slightly more vulnerable on average than non-calcifiers (6.1 vs. 2.8% mean habitat loss) but the difference was not significant due to high intra-group variability (*t*-test,  $P = 0.18$ , Figure 4). Neither biogeographic affinity nor higher taxonomic level had a significant influence on the group-mean change in habitat (one-way ANOVA tests,  $P = 0.39$  and  $0.80$ , Figure 4). When Arcto-Boreal and Arctic taxa were combined, their mean habitat

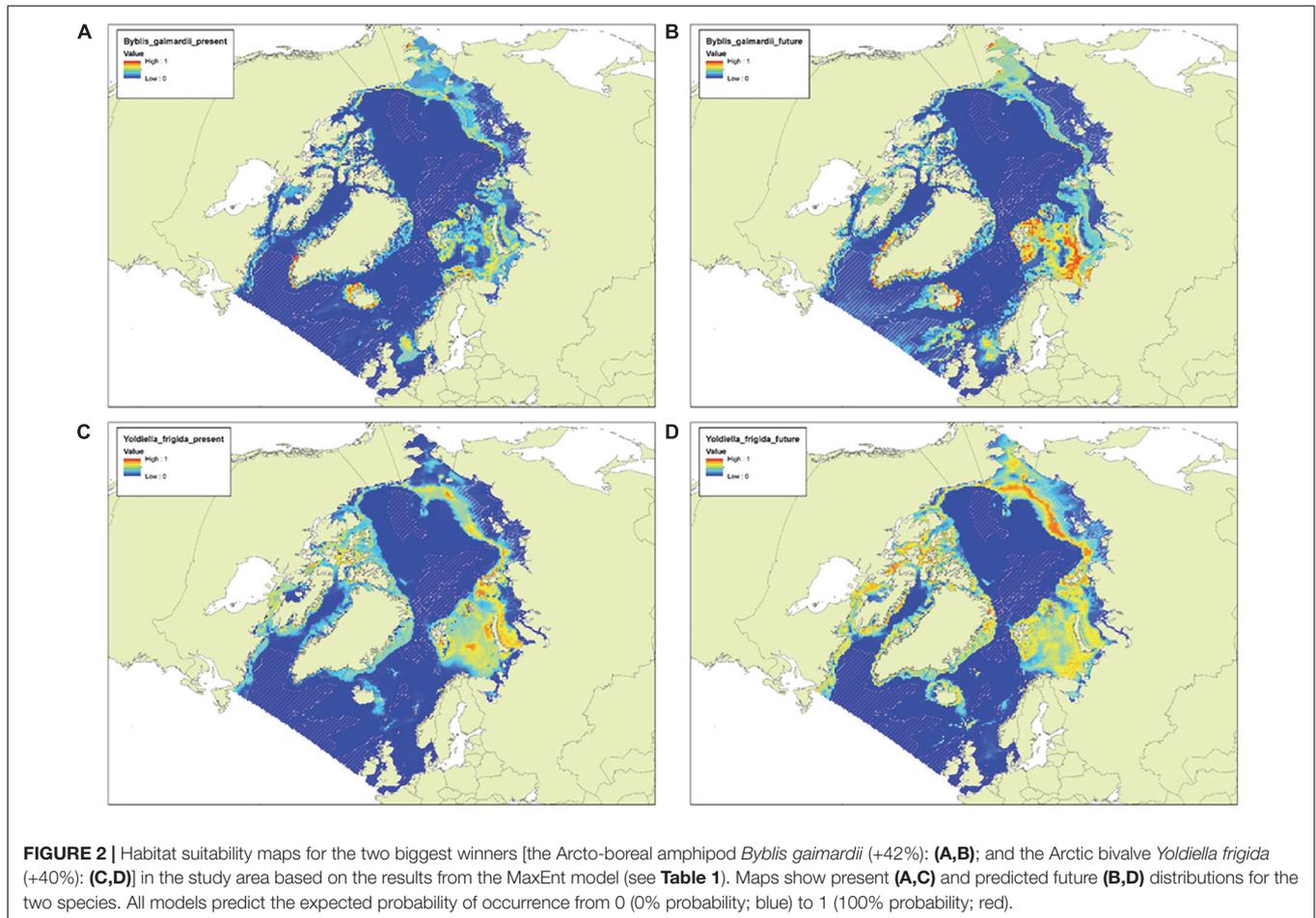
**TABLE 1** | Winners (>20% increase in suitable habitat, upper panel) and losers (>20% decrease in suitable habitat, lower panel) as predicted by the MaxEnt model.

Taxon	Change in habitat (%)	N	Higher taxon	Calcify?	Biogeography
<i>Byblis gaimardii</i>	+42	356	Crustacean	No	ArctoBoreal
<i>Yoldiella frigida</i>	+40	41	Bivalve	Yes	Arctic
<i>Astarte crenata</i>	+22	115	Bivalve	Yes	ArctoBoreal
<i>Parastichopus tremulus</i>	-53	30	Echinoderm	Yes	Boreal
<i>Notoproctus</i> sp.	-36	210	Polychaeta	No	ArctoBoreal
<i>Haploops tubicola</i>	-28	294	Crustacean	No	ArctoBoreal
<i>Mya truncata</i>	-27	655	Bivalve	Yes	ArctoBoreal
<i>Balanus balanus</i>	-26	764	Crustacean	Yes	Boreal
<i>Pagurus pubescens</i>	-25	222	Crustacean	Yes	ArctoBoreal
<i>Yoldiella solidula</i>	-25	64	Bivalve	Yes	Arctic
<i>Scoletoma impatiens</i>	-24	292	Polychaeta	No	Boreal
<i>Astarte montagui</i>	-23	409	Bivalve	Yes	ArctoBoreal
<i>Haploops</i> sp.	-22	579	Crustacean	No	ArctoBoreal
<i>Yoldiella nana</i>	-21	342	Bivalve	Yes	ArctoBoreal

The number of records extracted from OBIS for each taxon (N), higher taxon, calcification status, and biogeographical zones are indicated for each taxon. See Figures 2, 3 for maps of present (1978–2008) and future (2090–2099) distributions for the two biggest winners and losers.

loss was less than for Boreal taxa (2.9 vs. 10.9% habitat loss,  $P = 0.043$ ). Typically, heavily calcified taxa from the Mollusca and Echinodermata phyla have been suggested to suffer greatest from acidification (Hale et al., 2011). Our analysis suggested that only 24 and 25% of the taxa in these phyla, respectively, would experience range reductions, compared to 45–100% for other phyla (Supplementary Table S2). Similarly, only 12% of Arctic taxa lost habitat, compared with nearly 50% of Arcto-boreal and Boreal taxa (Supplementary Table S2). Results were the same for all analyses whether the absolute or percent change in suitable habitat was analyzed. The 3-factor main effects ANOVA also indicated no significant factor effects (biogeography:  $F = 2.1$ ,  $df = 2$ ,  $P = 0.14$ ; calcification:  $F = 1.9$ ,  $df = 1$ ,  $P = 0.17$ ; higher taxonomic level:  $F = 0.70$ ,  $df = 4$ ,  $P = 0.59$ ).

The data we extracted from the OBIS database were at both the species and genus levels. In most cases, genus-level taxa were assigned to broader biogeographic zones (e.g., ArctoBoreal) because they include several species. We tested for this potential bias by analyzing only the data for taxa identified to the species-level or where it was reasonably certain that the taxa found could be ascribed to either Boreal or Arctic zones. Results of these analyses with 23 fewer taxa were nearly identical to those for the entire data set (3-factor main-effects ANOVA: biogeography:  $F = 2.4$ ,  $df = 2$ ,  $P = 0.10$ ; calcification:  $F = 0.89$ ,  $df = 1$ ,  $P = 0.35$ ; higher taxonomic level:  $F = 0.23$ ,  $df = 4$ ,  $P = 0.92$ ).



## DISCUSSION

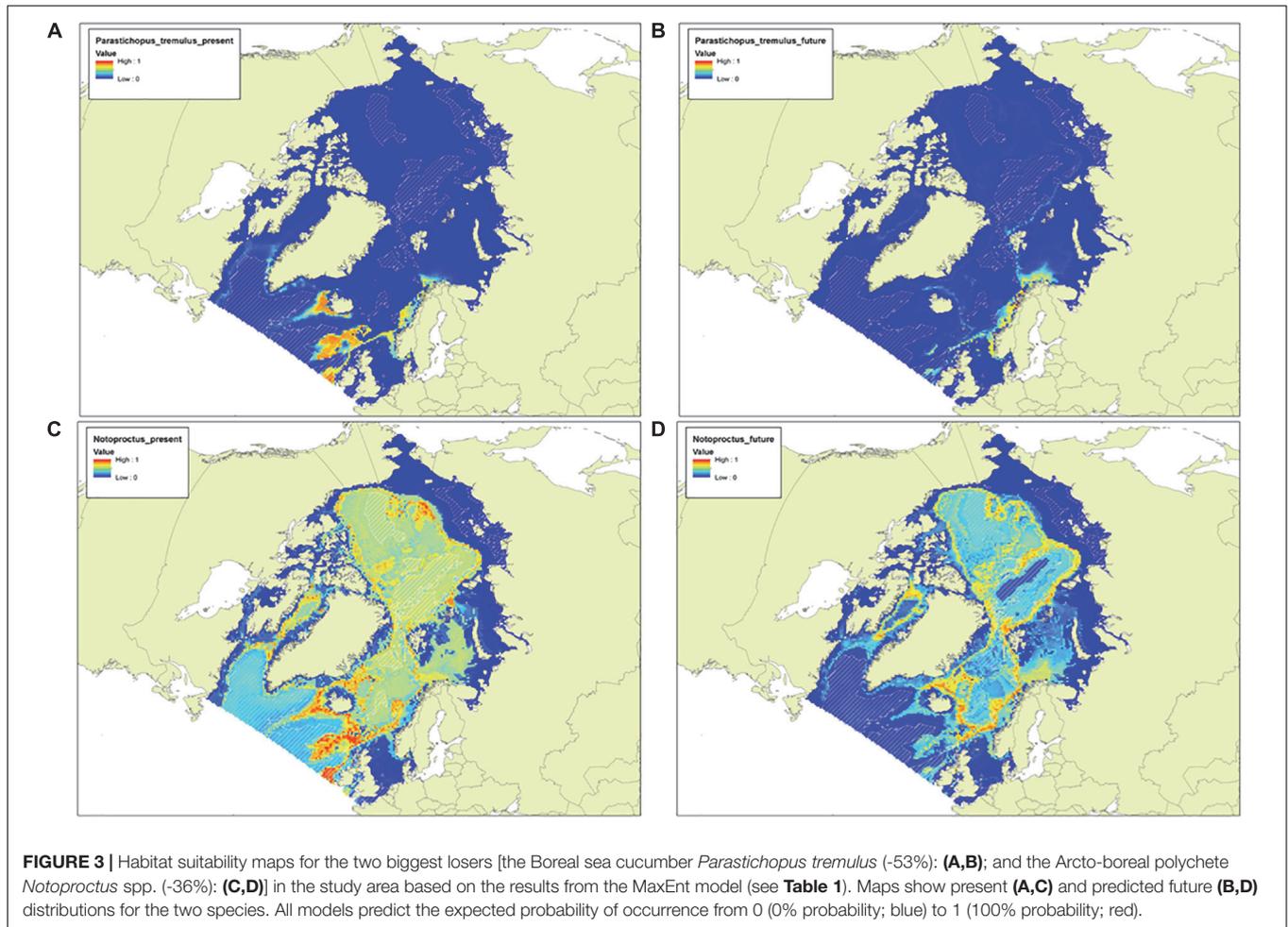
We found little evidence to support the belief that Arctic taxa are more sensitive than taxa with more southerly or cosmopolitan distributions, nor did we find that calcifiers as a group are more sensitive to projected climate-related changes than non-calcifiers. Models did indicate, however, that individual taxa can experience considerable changes in suitable habitat under future scenarios, with potentially significant ecological consequences.

### Roles of Niche-Defining Parameters

When fitting MaxEnt models, temperature was the most important factor determining current suitable habitat for more than two-thirds of the taxa studied. This should come as little surprise as temperature is one of the main factors used to delineate biogeographical provinces (Longhurst, 1998), so studies over a large spatial scale with strong temperature gradients such as this are likely to show an important temperature effect. Temperature was also identified (by MaxEnt modeling) as the most important factor for mollusk and crustacean species distributions at local scales in an Arctic fjord (Drewnik et al., 2017). Temperature can act in many ways, and may be viewed as a “master parameter” that works directly through thermal tolerance for survival and/or reproduction (e.g., Wetthey and

Woodin, 2008), or indirectly through its impacts on, e.g., primary production, ice cover, and zooplankton and microbial grazing pressure, all of which could affect benthic food supply (Maar and Hansen, 2011). Shifts in community structure of Barents Sea benthos and fish have been noted, with the southern limit of Arctic communities moving northward during warming and retreating again southward during cooling periods (Blacker, 1965; Matishov et al., 2012; Fossheim et al., 2015). These changes were attributed to temperature, but it is unclear whether they were direct tolerance effects or something more indirect. It is important to note, however, that some high-latitude taxa have very narrow temperature ranges over which they occur, making them particularly vulnerable to climate-driven changes (Pörtner et al., 2014; Morley et al., 2019).

We used aragonite saturation state to represent the degree of potential species response to ocean acidification, but we acknowledge that this may not be the only relevant carbonate system driver. Whereas most of the calcifying taxa we studied produce aragonitic skeletons (Mollusca: Kukliński, unpublished data), there is also a large number of organisms included in our study which use calcite with varying degrees of Mg incorporation (Echinodermata, Bryozoa, Cirripedia: Kukliński and Taylor, 2009; Iglukowska et al., 2017, 2018). In addition, some species may be more sensitive to pH or pCO<sub>2</sub> or combinations of OA stress.



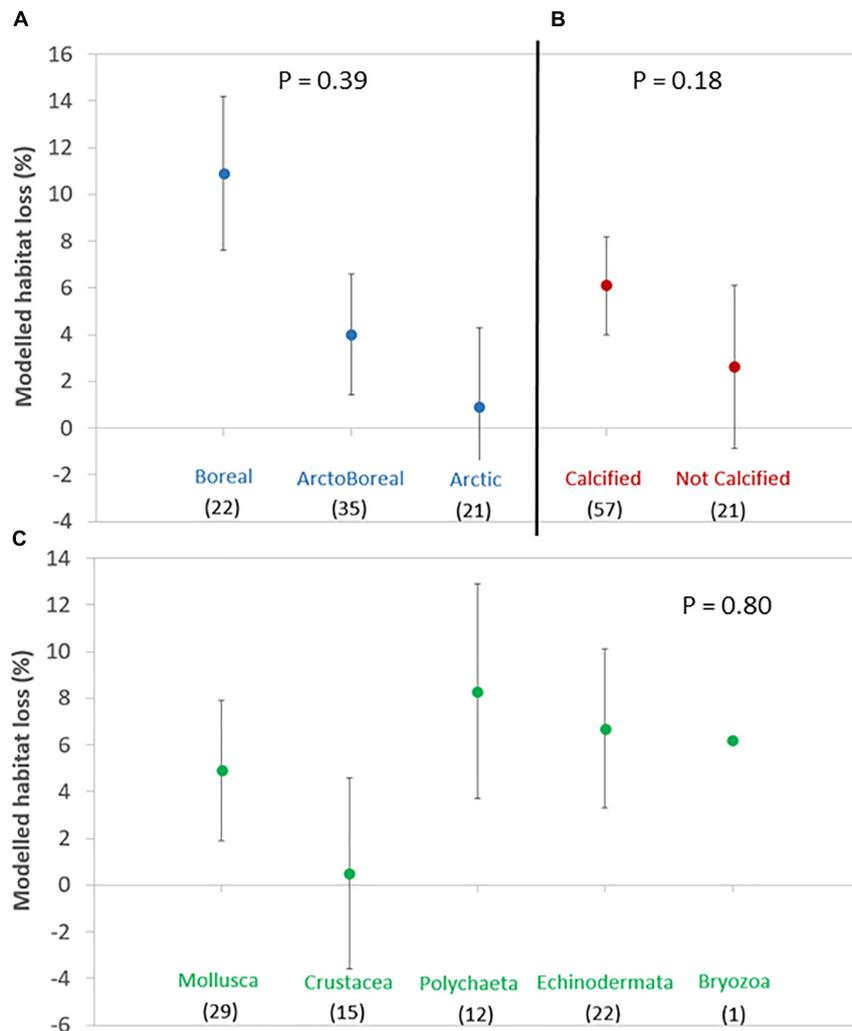
However, we suspect that using only one parameter (aragonite saturation state) provides a reasonable first approximation, since most changes are expected to be quite strongly correlated between different carbonate system parameters (e.g., Figure 9 in Wallhead et al., 2017).

We included depth in our distribution models, even though it will not change significantly in the near future. Depth can often reflect a number of linked parameters important for an organism's habitat state, such as sediment grain size, bottom currents, and vertical flux of particulate organic carbon (i.e., a proxy for food for many benthic organisms). Maintaining homeostasis under conditions of ocean acidification, for example, is thought to require more energy, and, as such, may only be possible when sufficient resources are available for affected organisms (Pansch et al., 2014). Increased metabolic rates due to higher temperatures may enhance this challenge (Pörtner, 2008). Primary production in the Barents Sea is predicted to change, both quantitatively and spatially, due to climate change (Slagstad et al., 2015). Clearly, inclusion of depth in the model will not account for all of this, but given the strong relationship between depth and vertical flux, we expect use of depth as one of the niche parameters to some extent accounted for spatial differences

in food supply, in addition to partially controlling for other habitat parameters.

### Accuracy of Environmental Data

The pan-Arctic SINMOD model, used here to provide bottom water temperature and aragonite saturation state, has been extensively tested and bias-corrected using field measurements (Slagstad et al., 2015; Wallhead et al., 2017). We therefore consider that the hindcast data used to define faunal niches are unlikely to have been a major source of bias. Projected changes for the 2090s are, however, subject to numerous model uncertainties and an ensemble approach would likely be required to quantify these. Nevertheless, we know that SINMOD gives bottom water projections that are comparable with global and other downscaling models, and it is sufficient for this study that the model gives plausible projections for the chosen climate change scenario (SRES A1B). For end-of-century simulations, the dominant source of uncertainty is likely to be the climate change scenario itself (Hawkins and Sutton, 2009). Water depth estimates for the pan-Arctic region are also not without uncertainty (e.g., the absolute differences between plausible measured values and the interpolated bathymetry products had median = 6 m, 95th percentile = 103 m) but it seems



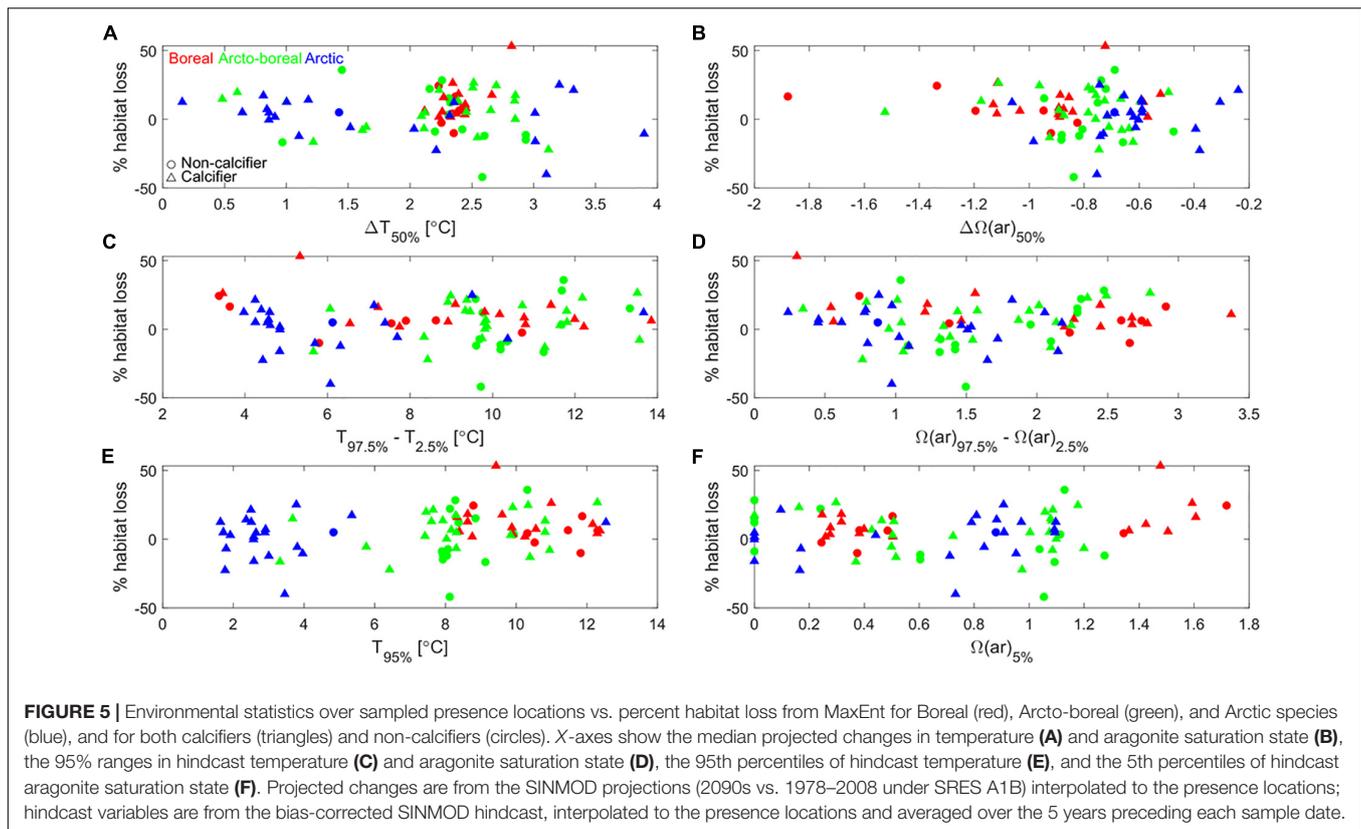
**FIGURE 4 |** Mean ( $\pm 1$  SE) percent habitat loss for benthic taxa between the present (1978–2008) and the end of the 21st century (2090–2099). *P*-values of analyses of variance are included for comparisons of species grouped into: **(A)** biogeographic zone, **(B)** calcification status, and **(C)** higher taxonomic group. Number of taxa in each group is indicated by the number in parentheses under the group name.

unlikely that such errors could significantly bias our estimates of mean habitat loss.

## Surprising Robustness of Habitat Suitability

Given the reported higher vulnerability of polar organisms to ocean warming (IPCC *medium confidence*, Pörtner et al., 2014) and of highly calcified organisms to ocean acidification (IPCC *high confidence*, Ibid), the fact that our projections for benthic taxa at high northern latitudes show no clear increase in habitat loss for Arctic or calcifying species may seem surprising. It leads us to question whether something could have been lost in the complexities of the MaxEnt analysis, and if the results might be understood or corroborated by simpler analyses based on the statistics of the environment where the different species were found.

A first reason to expect higher Arctic vulnerability is that environmental changes for Arctic species may be larger under climate change, due to rapid loss of ice cover/thickness and stronger warming in polar regions (Smetacek and Nicol, 2005; Brierley and Kingsford, 2009; Doney et al., 2012). However, if we consider the median projected bottom water temperature change at the sampled presence points for each species (see **Figure 5A**), the species-mean warming is slightly higher for non-Arctic species (2.3 vs. 1.8°C,  $P = 0.04$ , *t*-test, unequal variances). **Figure 5A** also shows that the warming is much more consistent for Boreal species than for non-Boreal species (standard deviation = 0.15 vs. 0.87°C;  $P < 10^{-11}$ , *F*-test). Regarding the median decreases in aragonite saturation state at sampled points (**Figure 5B**), the mean decreases are again slightly larger for non-Arctic species (0.87 vs. 0.63,  $P < 10^{-3}$ , *t*-test). So at least for bottom water temperature and saturation state, it seems the environmental changes are not larger for Arctic species, as



one might have anticipated from **Figure 1**. Furthermore, there are no significant correlations between the median environmental changes and the % habitat loss (**Figures 5A,B**).

A second basis for high Arctic sensitivity is the idea that polar organisms have narrow environmental tolerance ranges, e.g., a “stenothermal” characteristic due to low energy-demand lifestyles and specific adaptations to cold water (Pörtner et al., 2014; Morley et al., 2019). We can test this by considering the 2.5–97.5% ranges in bottom water temperature and aragonite saturation state at the sampled presence points as tolerance metrics for each species (**Figures 5C,D**). Here we find that Arctic species do have somewhat narrower ranges than non-Arctic species on average (6.0 vs. 9.6°C, 1.2 vs. 1.8;  $P < 10^{-3}$ , *t*-tests). However, variability in tolerance ranges is large within each group (**Figures 5C,D**), and there are no significant correlations between % habitat loss and tolerance ranges for temperature or saturation state (or indeed water depth, not shown). The rather moderate differences in mean tolerance range may in part reflect a higher level of historical variability in Arctic vs. Antarctic systems, at least for temperature (Pörtner et al., 2014). Arctic shelf systems have experienced, even in the past 150 years, several cycles of warming and cooling within the range we are observing today, and Arctic benthic fauna have expanded or retreated on decadal scales accordingly (e.g., Blacker, 1965).

A third basis for high Arctic sensitivity is the idea that polar organisms will have their habitat “squeezed out” under global change, either because they cannot move to higher latitudes in response to warming (Pörtner et al., 2014; Renaud et al., 2015), or

because critical thresholds of saturation state will be approached first in the Arctic under ocean acidification (Feely et al., 2009; Steinacher et al., 2009; Ciais et al., 2013). If we consider the 95th percentile of bottom water temperature at the presence locations as a simple metric of upper thermal limit (**Figure 5E**), we find that Arctic species do indeed have a significantly lower upper thermal limit on average (3.3 vs. 9.2°C,  $P < 10^{-12}$ , two-sample *t*-test). However, there is only a marginally significant correlation between % habitat loss and upper thermal limit (Pearson’s  $r = 0.23$ ,  $P = 0.04$ ). Considering the 5th percentile of aragonite saturation state at presence locations as a lower tolerance limit and potential predictor of vulnerability to acidification, we find no significant difference between Arctic vs. non-Arctic species, and no significant overall correlation with % habitat loss (**Figure 5F**).

With regard to the paradigm of higher sensitivity of calcifying species, we find no significant difference in species-mean 5th percentiles of aragonite saturation state at presence locations for calcifiers vs. non-calcifiers (0.71 vs. 0.67,  $P = 0.76$ , *t*-test), and both categories contain species with 5th percentiles much less than 1, and in several cases  $< 0.2$  (**Figure 5F**). Such tolerance of corrosive water may be partly explained by physiological mechanisms associated with the calcification process, particularly in these often heavily calcified taxa, which are already adapted for internal control of carbonate saturation state at the site of precipitation (Hendriks et al., 2015). Considerable variability in response to ocean acidification has been shown in experimental manipulations (Kroeker et al., 2011; Vihtakari et al., 2016), and

naturally high environmental variability in some habitats (shallow waters, vegetated substrate) may also help to promote tolerance via phenotypic plasticity and local adaptations (Hofmann et al., 2010). Such mechanisms may make the species less susceptible to modest decreases in saturation state, even at under-saturation levels, although food supply may mediate the ability of an organism to regulate conditions for calcification (Ramajo et al., 2016).

## Analysis of Habitat Losses in Niche Space

It is worth noting that neither large environmental changes nor narrow tolerance ranges will necessarily result in high species habitat loss; if climate change produces enough new habitat, the net loss of habitat may be zero (or indeed negative) no matter how large the changes or narrow the tolerances. Rather, large changes and narrow tolerances will likely result in large (negative or positive) changes in suitable habitat area. To better understand the projections, it can be helpful to consider the calculation of % habitat loss in niche space – here a 3D space of (temperature, saturation state, water depth). The niche space can be divided into a grid of subvolumes, where the  $i$ th subvolume “contains” a physical area of habitat  $A_i^{(1)}$  in the present and  $A_i^{(2)}$  in the future. The MaxEnt algorithm essentially determines an occupancy probability  $P_{ji}$  for the  $j$ th species and  $i$ th niche-space subvolume by fitting a complex preference function to the present-day niche occupancy data (Merow et al., 2013). Assuming no change over time in this niche-space preference function (ruling out e.g., genetic adaptation), the net change in habitat area  $H_j$  for species  $j$  is given by:

$$\Delta H_j = \sum_i P_{ji} (A_i^{(2)} - A_i^{(1)})$$

It follows that the projected habitat losses can in large part be explained by the changes in niche subvolume area content  $\Delta A_i = A_i^{(2)} - A_i^{(1)}$ . These latter are entirely independent of species and the MaxEnt procedure, and depend only on the present and future environment (here based on the SINMOD biogeochemical model and the high-resolution bathymetry products).

The habitat area changes over niche space can be visualized on a 2D grid over temperature ( $T$ ) and aragonite saturation state ( $\Omega(\text{ar})$ ) by integrating over the water depth dimension (Figure 6A). The largest losses are in the red cells around ( $-2 \leq T < 1^\circ\text{C}$ ,  $1 \leq \Omega(\text{ar}) < 1.5$ ), while the largest gains are in the blue cells around ( $-2 \leq T < 1^\circ\text{C}$ ,  $0 \leq \Omega(\text{ar}) < 0.2$ ) and ( $0 \leq T < 5^\circ\text{C}$ ,  $0.4 \leq \Omega(\text{ar}) < 1$ ). In order to be a “winner” under the projected environmental change, a species needs to have an occupancy probability or niche preference function  $P_{ji}$  that favors the niche subvolumes that gain habitat area. This requires that its present-day sampling distribution over niche space gives evidence that the species can tolerate the winning blue cells. For example, the biggest winner, the amphipod *B. gaimardii* (+42% habitat area), has a widespread niche space distribution (magenta crosses). Comparing this to the sampling distribution over all species (the TGS, black dots), the MaxEnt algorithm infers a strong tolerance of low  $\Omega(\text{ar})$  and a thermal window that

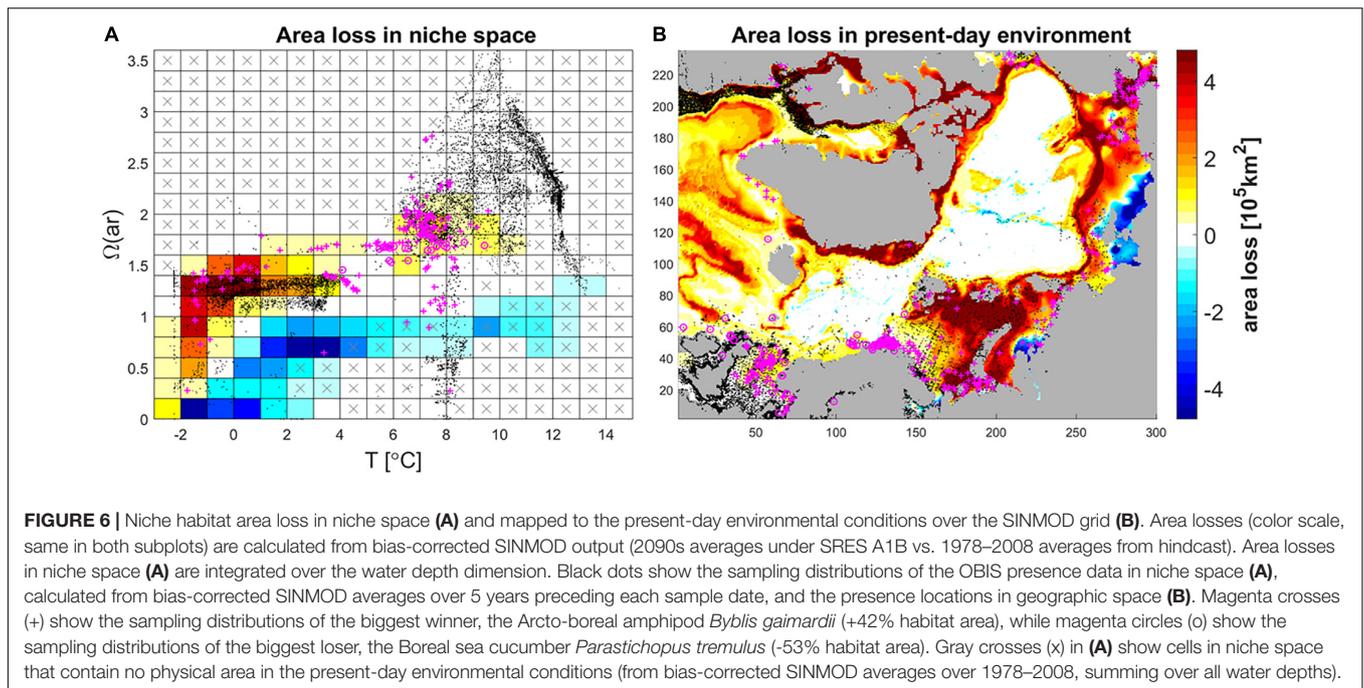
comfortably contains the winning range ( $0 \leq T < 5^\circ\text{C}$ ). This allows *B. gaimardii* to make large area gains in the future Barents Sea, Bering Strait, Chukchi Sea, and on the East Siberian shelf between 50 and 500 m (Figures 1B,E, 2A,B). By contrast, the biggest loser, the sea cucumber *P. tremulus* (-53% habitat area), has a much more confined sampling distribution in niche space (magenta circles), suggesting limited tolerance of undersaturated water ( $\Omega(\text{ar}) < 1$ ) and a thermal window of roughly 4–10°C. Consequently, *P. tremulus* loses habitat in the North Atlantic and becomes increasingly confined to the Norwegian shelf shallower than 500 m (Figures 3A,B).

Figure 6A also sounds a note of caution regarding our projections: the sampling distribution of presences (black dots) is highly non-uniform over niche space, and in particular the winning niche cells (blue colors) are strongly undersampled. For several winning cells, the present-day niche area is zero (indicated by a gray “x”); these are new niches that do not exist yet in the pan-Arctic region. Although the MaxEnt procedure is rarely strictly extrapolating with respect to the 3D convex hull (see section “Materials and Methods”) it is often being used to make bold interpolations based on limited data. This issue, plus other potential sources of bias (e.g., gear biases), reduce our confidence in the projections for individual species habitat loss, although we suspect a lesser impact of sampling bias on the significance of differences between group means. Our first attempt at running the analysis used the default MaxEnt assumption of uniform sampling effort over the pan-Arctic grid (a clearly untenable assumption, see Figure 6B); this led to much larger group-mean habitat losses (59–82%), but mean losses of Arctic species (75%) were still not significantly different to those of boreal (59%) or Arcto-boreal species (82%). The results presented in this paper account for this first assumption.

Further insights into vulnerability and sampling needs are provided by interpolating the 3D niche space area changes onto the present-day (1978–2008 average) values over the SINMOD grid (Figure 6B). Here, any species whose present-day geographical distribution is confined to areas of niche area loss (yellow/red colors) is set to be a loser under the projected changes. Hence, we can project, independently of any MaxEnt analysis, that species with environmental tolerances that confine them to e.g., the northern Barents Sea, or the East Greenland Shelf, are likely to lose habitat under climate change. Conversely, any species that tolerates present-day conditions in the Kara Sea near the Gulf of Ob, or the shallow areas (<50 m) of the eastern Laptev Sea and East Siberian Sea, is set to win. It is thus unfortunate that relatively few species presence records from these latter areas have made it into the OBIS database, and future attempts to project future species distributions would likely be well served by more sampling effort in and/or data recovery from these particular regions.

## Ecological Consequences of Distributional Shifts

Approximately half the taxa were projected to experience substantial changes in suitable habitat area (>10% increase or decrease), and some of these species are characteristic



or dominant on Arctic continental shelves. The polychetes *Chaetozone setosa* and *Spiochaetopterus typicus*, the bivalve *Medicula ferruginosa*, the barnacle *Balanus balanus*, and the decapods *Pagurus pubescens*, *Hyas araneus*, and *Pandalus borealis* are common and characteristic taxa in the open Barents Sea and adjacent fjords. *S. typicus* (12% increase in habitat) and *B. balanus* (26% loss) are key ecosystem engineers in that the physical structure they provide is important for ecosystem function. *Pandalus* is an important commercial shrimp in Arctic and boreal waters, and a 17% increase in habitat was predicted. If these taxa are as strongly affected as our analyses suggest, the impact on ecosystem structure and services in the Barents region could be significant over the 21st century.

Surprisingly, one of the biggest “winners” in our analyses, the Arctic bivalve *Yoldiella frigida* (40% habitat increase), was described as one likely to experience substantial range retraction in an analysis based solely on temperature ranges (Renaud et al., 2015). However, the Renaud et al. (2015) projection considered only the Barents Sea. The MaxEnt algorithm predicts a much more modest increase for the Barents Sea (9%), with most of the pan-Arctic habitat increase driven by gains on the Chukchi/Russian shelves and coastal waters off Canada, Greenland, and Iceland (Figure 2D). Also, Renaud et al. (2015) assumed a hard upper temperature limit of 4°C; the MaxEnt algorithm infers some tolerance of warmer temperatures (thus allowing for the slight increase in the future Barents Sea) based on the limited presence records (41) within the hindcast timeframe (1978–2008). In fact the histogram shown in Figure 5 of Renaud et al. (2015) also suggests some tolerance to warmer waters, although this latter is based on August-only SINMOD output for 2000–2009 without bias correction or matching to sample date.

One caveat of our study is the exclusive focus on bottom water environment, while many benthic organisms reproduce via planktonic larvae that live for hours to months in the pelagic zone. A recent study of an Antarctic barnacle indicated the need to consider both stages of multi-phasic species since conditions are likely to change differently in the two ocean realms (Gallego et al., 2017). Registered presence in the OBIS database, as used in our study, integrates environmental influence on multiple life-history stages/processes from fertilization, to development and dispersal, to recruitment and juvenile stages, any of which can be acted upon by ecological (including acidification) interactions in both the water column and sea floor. Another possible caveat is our use of only two physical/chemical parameters. Many other such parameters (e.g., carbon flux, bottom-water oxygen content) may define where an organism can exist, and ecological interactions cannot be expected to be unchanged as species ranges change. Kroeker et al. (2012) found ocean acidification to alter competitive hierarchies of macroalgae, and this could have consequences for associated fauna. In contrast to what has been observed for Barents Sea fish communities (Fossheim et al., 2015), benthic community change is not likely to progress via wholesale replacement of Arctic fauna with boreal fauna. Highly mobile fish are expected to move northward and southward in response to changing extent of Atlantic Waters, whereas benthic taxa have limited mobility and range changes are more likely to vary by taxa as a consequence of life-span, dispersal ability, etc. New species combinations will likely lead to novel and unpredictable species interactions, further affecting species distributions (Pech et al., 2017). Finally, our models are based on fixed niches so that populations follow the environmental change by shifting their geographical distributions. However, populations may adapt to

tolerate novel conditions through selection by evolving traits that provide tolerance in the new environment. Although it is tempting to project future distributions that take into account evolutionary responses, scientists seldom know if, or how quickly, the climate-sensitive traits of populations can evolve (Merilä and Hendry, 2014; Jezkova and Wiens, 2016).

## CONCLUSION

Despite acknowledged issues and uncertainties with using species distribution models, especially in poorly sampled areas like the Arctic, such models are accepted as useful in providing a first approximation of how climate change will affect biodiversity (Pearson and Dawson, 2003; Wiens et al., 2009). This study is not meant to emphasize fine-scale species-specific changes, but rather to delineate expected shifts in broad patterns. The findings reported here are surprising in the context of current paradigms regarding susceptibility to climate change. In particular, we found that for benthic species under end-of-century ocean warming and acidification: (i) mean habitat losses over taxonomic groups were generally small (0–11%), (ii) Arctic benthic species were not significantly more vulnerable than boreal or Arcto-boreal species, (iii) calcifying species were not significantly more vulnerable than non-calcifiers. The lack of sensitivity of our results to such taxonomic groupings was consistent with simpler analyses based on magnitude of environmental change and observed tolerance windows, although such criteria were individually poor predictors of the habitat loss results from MaxEnt. Sampling bias in the presence data was a major issue, and in general we expect projections from benthic species distribution models for high northern latitudes to be strongly sensitive to how this is dealt with. Analysis of niche space area changes, dependent only on the environmental data, suggested that largest future habitat losses will be for taxa now inhabiting cold/oversaturated niches ( $-2 < T < 3^{\circ}\text{C}$ ,  $1 < \Omega(\text{ar}) < 1.5$ , e.g., the present-day northern Barents Sea and East Greenland Shelf) while the largest gains will be for those currently found in cold/undersaturated niches ( $-2 < T < 5^{\circ}\text{C}$ ,  $\Omega(\text{ar}) < 1$ , e.g., the present-day eastern Laptev and East Siberian Seas). The latter “winning” niches were undersampled in our dataset, and we expect that further field sampling and data recovery for these niches will be especially helpful for projecting future species distributions.

There are few similar studies from high latitudes (but see Morley et al., 2019 for Antarctic waters), mainly due to limited information about species ranges and physiological tolerances, and regional models powerful enough to project environmental parameters into the future. As more data become available and models are developed, the tools we used here can be improved, and predictions refined. As in other predictive modeling approaches, MaxEnt is a process that uses data mining and probability to forecast outcomes. When doing so, the established relationships are only based on statistical dependence between environmental and distributional data, and therefore may fail to account for physiological limits and biological interactions. Development of hybrid modeling platforms that integrate species distribution models and experimental results

can facilitate higher resolution distribution models for individual taxa, which can be particularly useful for economically or ecologically influential species (Kotta et al., 2019). This work is important as changes in ranges of some of these species may have substantial impacts on ecosystem services provided by benthic communities (Pecl et al., 2017).

## DATA AVAILABILITY

The datasets generated for this study are available on request to the corresponding author. Model output data are available on Zenodo (<https://zenodo.org/record/3245446>).

## AUTHOR CONTRIBUTIONS

PR, MW-K, and PK developed the study idea. JK and MR performed MaxEnt modeling. RB and DS developed the SINMOD simulations. PW analyzed and bias-corrected the SINMOD output. PK was the project leader. PR led work on the manuscript. All authors contributed to the interpretation of results and editing of the manuscript.

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## SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fmars.2019.00538/full#supplementary-material>

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**Conflict of Interest Statement:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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