<u>Title:</u>

Distribution and predicted climatic refugia for a reef-building cold-water coral on the southeast US margin

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Climate change is reorganizing the planet's biodiversity, necessitating proactive management of species and habitats based on spatiotemporal predictions of distributions across climate scenarios. In marine settings, climatic changes will predominantly manifest via warming, ocean acidification, deoxygenation, and changes in hydrodynamics. Lophelia pertusa, the main reef-forming coral present throughout the deep Atlantic Ocean (> 200m), is particularly sensitive to such stressors with stark reductions in suitable habitat predicted to accrue by 2100 in a business-as-usual scenario. However, with new occurrence data for this species along with higherresolution bathymetry and climate data, it may be possible to locate further climatic refugia. Here, we synthesize new and published biogeographic, geomorphological, and climatic data to build ensemble, multi-scale habitat suitability models for L. pertusa on the continental margin of the southeast United States (SEUS). We then project these models in two timepoints (2050, 2100) and four climate change scenarios to characterize the occurrence probability of this critical cold-water coral (CWC) habitat now and in the future. Our models reveal the extent of reef habitat in the SEUS and corroborate it as the largest currently known essentially continuous CWC reef province on earth, and also predict abundance of *L. pertusa* to identify key areas, including those outside areas currently protected from bottom-contact fishing. Drastic reductions in L. *pertusa* climatic suitability index emerged primarily after 2050 and were concentrated at the shallower end (< ~550 m) of the regional distribution under the Gulf Stream main axis. Our results thus suggest a depth-driven climate refuge effect where deeper, cooler reef sites experience lesser declines. The strength of this effect increases with climate

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scenario severity. Taken together, our study has implications for the regional and global management of this species, portending changes in the biodiversity reliant on CWC habitats and the critical ecosystem services they provide.

Introduction

Climate change is affecting global biodiversity by reorganizing species distributions. Spatial refugia (*sensu* Keppel & Wardell-Johnson 2012) from climate change can dictate complex distributional legacies, which leave behind evidence of faunal origins and climate-induced fluctuations by selecting for resilience or by creating faunal patchworks through extirpation (Horne 1999; McClain & Hardy 2010; Carpenter et al., 2001; Fernandez et al., 2021). Therefore, locating and protecting refugia from additional human pressures is imperative to climate-smart management that is proactive against future changes in biodiversity (Arafeh-Dalmau et al., 2020; Queirós et al., 2021).

In the marine realm, climate change manifests primarily through warming, acidification, deoxygenation, and alterations in hydrodynamics affecting food supply (Doney et al., 2012; Levin & Bris 2015). In surface waters, models predict large expanses with either novel or disappearing climates by 2100 (Lotterhos et al., 2021; Dixon et al., 2022), yet little is known about no-analogue climates in the deep ocean (> 200 m depth) where most of the planet's habitable area exists. However, *in situ* trends (Johnson et al., 2014; Desbruyères et al., 2016; Meinen et al., 2020), and climate models (Sweetman et al., 2017; Morato et al., 2020) suggest rapid ocean change will accumulate at the seafloor this century. Many deep-sea ecosystems may soon experience oceanographic conditions with only distant spatial or temporal analogues. This can cause mobile species' distributions to move deeper towards cooler waters (Pinsky et al., 2013; Chaikin et al., 2021) and reorganization and compression of ecological communities (Mora et al. 2013; Chu & Tunnicliffe 2015; Sato et al., 2017;

Gasbarro et al. 2019). The temporal scales of ocean change make it less likely that slow-growing sessile species will be able to track their climatic niche via larval dispersal.

Amongst the most at-risk taxa are cold-water corals (CWCs), which form the basis for Vulnerable Marine Ecosystems including coral reefs and gardens (FAO 2009). *Lophelia pertusa* (proposed revision to *Desmophyllum pertusum*; Addamo et al., 2016) is the most well-known reef-building coral in the deep ocean. It is found in every major ocean basin and as an autogenic habitat engineer creates coral reefs and carbonate mounds of varying sizes that are formed over thousands of years by generations of coral growth/death leading to mound build-up by coral skeleton and baffled sediment (Raddatz et al., 2014). This species has a long pelagic larval duration, and these planktotrophic larvae readily cross density gradients and may even spend time in the photic zone, suggesting an ability to disperse over great distances (Larsson et al., 2014; Strömberg & Larsson 2017).

Notable *L. pertusa* reefs have been found in the South Atlantic on seamounts (Bridges et al., 2021), on the Brazilian (Cavalcanti et al. 2017), and west African margin (Hanz et al., 2019; Hebbeln et al., 2020); in the North Atlantic off of Florida (Reed et al., 2006) and Greenland (Kenchington et al., 2017), Norway (Fossa et al. 2002), Canada (Buhl-Mortensen et al., 2017), the UK (Howell et al. 2011), and the Mediterranean (Freiwald et al., 2009; Carlier et al., 2009; Mastrototaro et al., 2010). These mounds can cluster into large mound provinces covering up to thousands of square kilometers in areas where oceanographic conditions allow, including in the Gulf of Mexico (Hebbeln et al., 2014; Roberts & Kohl 2018), Northeast Atlantic (Kenyon et al., 2003; Masson et al., 2003; Dullo et al., 2008; Howell et al., 2011; Flögel et al., 2014) and Northwest

Atlantic (Reed et al., 2006). These CWC mounds and the living reefs that often top them are remarkable both in their spatial extent and their functioning as important hotspots of biodiversity (Roberts et al., 2006; Henry & Roberts 2017) and nutrient recycling (van Oevelen et al., 2009; Cathalot et al., 2015). Thus, accurate predictions of their expanse are needed to improve our understanding of the function of ecosystem services of the global ocean.

The Blake Plateau is a large depositional feature on the Southeast US (SEUS) margin, shaped by the influence of the Gulf Stream and associated eddies. Vibrant CWC ecosystems flourish underneath the Gulf Stream due to elevated currents and pulses of fresh phytodetritus to the seafloor (Mienis et al., 2014). Since 2018, numerous research and exploratory expeditions to CWC habitats in this region have built upon historical baseline surveys (Stetson et al., 1962; Reed et al., 2006) with mapping efforts revealing the largest nearly continuous mound province on Earth with tens of thousands of apparent CWC mounds extending over approximately 700 km from offshore Florida to Virginia (Sowers 2020), including a remarkable coral reef complex over 200 km in total length off of Charleston, South Carolina herein referred to as the "Richardson Reef Complex" (Figure 1). Coral rubble appears to be the dominant substrate on even the smallest mounds in the region, indicating CWC habitability at some point in the Holocene, if not currently. It is unknown how many of the mounds are topped with living coral and, if so, how much.

These sites face imminent climate threats, as climate change causes the Gulf Stream to deliver increasingly warm and acidic subtropical waters to their depth range (~400-900 m) (Saba et al., 2016) but potentially at lower velocities (Boers et al., 2021;

Caesar et al., 2021), simultaneously lowering the rate of food delivery to less optimal levels. However, all CWC ecosystems of this region may not be affected equally by climatic changes, as many shallower sites near the shelf break sit directly underneath the Gulf Stream's velocity core while deeper sites on the eastern Blake Plateau are affected more intermittently by Gulf Stream meanders. Paleoecological reconstructions of past climates from a Cape Lookout coral mound indicate that CWCs in the SEUS proliferated most recently following hydrographic changes in the Gulf Stream system ~7 kya that changed the oceanographic conditions experienced by CWCs (Matos et al., 2015), suggesting that these ecosystems are sensitive to changes in climate.

Here, we present a novel biogeographic, geomorphological, and climatic data synthesis to build ensemble, multi-scale habitat suitability models for *Lophelia pertusa*. We project these models onto the present-day SEUS margin and two future timepoints (2050, 2100) under four climate change scenarios in order to characterize the extent of critical CWC ecosystems in the region and test several hypotheses regarding their future. First, our models test whether the smallest of the coral mounds mapped in the region are likely to contain *L. pertusa*, and whether terrain with high habitat suitability index (HSI) scores will be concentrated on one or more landforms associated with these mounds (i.e. peaks, ridges, and slopes). We also test whether greater declines in climatic suitability index (CSI) scores occur in more severe emissions scenarios, and whether the time of emergence for notable declines in CSI will be before 2050. Finally, we test for regional disparities in climate change outcomes by depth, predicting that shallower populations will face greater CSI and abundance declines due to their proximity to the Gulf Stream, with deeper sites serving as spatial climate refugia with

more modest changes. We also calculate climate change velocities to test this depthdisparity hypothesis, predicting higher velocities at shallower seafloor depths and isotherm trajectories will show a general trend towards deeper locations. Taken together, our results provide timely information required for both mitigation and adaptive spatial management of these ecosystems and the critical services they provide.

Materials & Methods

Occurrence Data

L. pertusa occurrence points were generated from submersible dive imagery from 11 dives completed in 2018-19 (Table S1.1). Dive video was annotated for *Lophelia* presence and georeferenced by timestamp. Additional presence points were downloaded from the NOAA Deep-sea Coral Research & Technology Program database (v102020; <u>https://deepseacoraldata.noaa.gov/</u>). Points with location error > 1000 m were removed, which generally excludes older, less reliable records (e.g. trawl records) that likely have lower fidelity to gridded environmental data. All points falling within a grid cell were merged to curtail pseudoreplications.

Abundance (= % cover) data were generated from one-minute video segments during the above dives. Video segments were chosen where the submersible was transecting the seafloor and image quality was high (i.e. not obscured by the submersible or water column turbidity). Five random screenshots were taken for each segment by selecting random numbers between 1 and 60 (1 representing the first second of the 1-minute segment), and then 50 random points overlain on each screenshot in ImageJ software using a custom-written macro. Points falling on live *Lophelia* were counted, giving an estimate of percent cover for each screenshot and

mean/standard deviations for each segment. In total, 516 segments were used to train abundance models (Table S1.1).

Terrain & Climate Data

We used the bathymetric dataset covering much of the Blake Plateau generated by Sowers (2020) to create bathymetric terrain variables at the highest available resolution for this area (3.5×10^{-40} or ~35 m). Slope, aspect [both cosine (N-S) and sine (E-W)], three types of curvature, fine (100 m) and broad-scale (1000 m) bathymetric position indices (BPI) comprised the initial set of terrain variables (Table 1). BPI was calculated with an inner radius of one cell. All terrain variables were generated from the bathymetric data with Benthic Terrain Modeler v3.0 (Walbridge et al., 2018) in ESRI Arcmap version 10.8. In addition, we used a 'bathymorphon' (underwater landform) classification scheme (Sowers 2020) of valley, flat, slope, ridge, and peak bathymorphons *post-hoc* to test whether *L. pertusa* occurrence probability was higher on particular bathymorphons. This classification was achieved using the Bathymetryand Reflectivity-based Estimator for Seafloor Segmentation (Masetti et al., 2018) using an inner and outer radius of one and six cells, respectively, and a flatness parameter of 1.5°.

A number of variables were used to model climatic suitability for *L. pertusa* (Table S1.2). Global Atmosphere/Ocean General Circulation Model data were downloaded in four-dimensional (X, Y, depth, time) netCDF format from the Earth System Grid Federation as either monthly or annual means. All data come from Climate Model Intercomparison Project 6 (CMIP6; Eyring et al., 2016), allowing for higher-resolution models than basin-scale models for *L. pertusa* based on CMIP5-era data

(e.g. Morato et al., 2020). CMIP6 data included the variables pH, mean dissolved oxygen, and export carbon from the high-resolution implementation of the Max Planck Institute's MPI-ESM1-2 model (Müller et al., 2018). Bottom temperature data were available from several modeling groups and thus a multi-model ensemble was used (see Table S2 for climate metadata).

CMIP6 models used in this study were forced under Shared Socioeconomic Pathways (SSPs). These SSPs range from the SSP1-2.6 "Sustainable Future" to the SSP5-8.5 "Business As Usual" scenarios with two intermediate pathways (SSP2-4.5, SSP3-7.0). The SSPs incorporate revised emission and land use pathways than those in the Representative Concentration Pathways of CMIP5 (see O'Neill et al., 2016; Riahi et al., 2017). Data for each SSP were averaged into current (1995-2014), mid-century (2031-2050), and end-of-century (2081-2100) timepoints, giving a rough estimate of the time of emergence for suitability changes. Climate Data Operators (Schulzweida 2019) was used to convert data from each model to annual means where necessary, re-grid from curvilinear to lat/long grids, and to extract bottom-most (= benthic) data at their native resolution (Table S1.2). These data were cropped and re-gridded at 0.007° (~700m) for the SEUS using bilinear interpolation.

CMIP6 bottom temperature data for the SEUS were compared to data from Alexander et al., (2020)'s dynamically downscaled model data that covered the study area with comparable data for the historical and 2100 business-as-usual scenarios at a higher native resolution (see Supplementary Materials 2). We chose to not include the Alexander et al., (2020) temperature data in the models due to the differences in methodology and native resolution between these and CMIP6 data, and thus used the

Alexander et al., (2020) data to explore possible biases in the CMIP6 data for the region. We did, however, use these data to calculate distance-based climate change velocity, i.e. the distance to the geographically closest thermal analogue in the SEUS in the year 2100 using the R package 'VoCC' (Molinos et al., 2019) due to its higher native resolution. A climate tolerance of 0.25°C and margin-wide search radius was used to find analogous cells. Thermal trajectories were overlain on the velocity map in order to visualize isotherm movements.

Habitat Suitability and Abundance Modeling

Variables with Variance Inflation Factors > 5 and/or highly correlated (Pearson's R > 0.9) variables were sequentially removed before the modeling process in order to minimize multicollinearity (Figure S3.1). The final variable set (Table 1) still included a few correlated variables (temperature, pH, O₂) that were included in the model for their ecological relevance to *L. pertusa* (Brooke et al., 2013; Hennige et al., 2014; Georgian et al., 2016a).

A multi-scale framework was used to model the distribution of *L pertusa* to incorporate the variation in the species' ecological response to both terrain and climate at relevant scales. That is, we captured the fine-scale (~35 m) landscape morphology that is fundamental to predicting CWC distributions by proxying hydrodynamics (Rengstorf et al., 2013), while separately modeling climatic suitability at a downscaled resolution (~700m). Suitability of both terrain and climate is a prerequisite for *L. pertusa* presence, and so decreasing the terrain resolution to match the climatic data would smooth over important information and decrease model precision (Miyamoto et al., 2017). The climatic niche of *L. pertusa* was first constructed for the whole of the North

Atlantic on the native resolution grid, then projected onto each scenario climate grid for the SEUS (n = 9), as the entirety of their niche was not represented in the SEUS. For example, pH is relatively high in the region. Thus, SEUS data would not capture a lower limit for *L. pertusa* suitability and a model trained solely on current SEUS climatology would not predict an influence of pH declines in future scenarios, despite the known relevance of pH in this range to the species' distribution and survival (Davies et al., 2008; Georgian et al., 2016a; Lunden et al., 2014; Hennige et al., 2020; Morato et al., 2020). Thus, there may be regional variability in *L. pertusa* climate-stressor tolerance in the SEUS that was not captured in our models (e.g. Georgian et al., 2016b).

Suitability scores were computed with one regression [Generalized Linear Models (GLM)] and two machine-learning algorithms [Random Forest (RF), Gradient Boosting Machines (GBM)]. For both the terrain and climate models, three presence/background replicates were run with 10,000 pseudoabsences each selected from <1000 km from presence points. This distance was chosen to reflect this species' capability of long-distance dispersal (Strömberg & Larsson 2017). Models were evaluated with repeat split-sample cross validation (70-30 testing/training splits). Five evaluation runs were computed for each pseudoabsence set and algorithm combination for a total of 45 models for each of terrain and climate. Response curves were generated according to Elith et al., (2005) and inspected for biologically plausible responses for each algorithm. Then, we calculated ensemble means and variances from model runs across algorithms to minimize individual model biases (Buisson et al., 2010). Individual and ensemble model discrimination accuracy was assessed with True Skill Statistic (TSS; Lawson et al., 2014) and area under the Receiver Operating

Characteristic (ROC) metrics. Relative variable importance in the models was estimated by the variable randomization approach recommended by Guisan et al., (2017). In addition, the degree to which the models make predictions outside of the environmental envelope of present-day *L. pertusa* was assessed using the Extrapolation Detection Tool of Bouchet et al., (2020).

Projected scenario climatic suitability index scores were compared to scores from the present-day model and suitability scores at known *L. pertusa* sites were regressed against depth to test the hypothesis that the deepest, eastward sites containing *L. pertusa* may act as climate refugia. The combined terrain-climatic suitability scores were then used to model the areal percent cover for each scenario using the Random Forest regression as in Hill et al., (2017). All computations were performed in R version 4.0.5 (R Core Team, 2021) on Temple University's High-Performance Computing server cluster. The R packages 'biomod2' (v. 3.5.1; Thuiller et al., 2021) and 'randomForest' (Liaw & Wiener, 2002) were used for terrain/climatic suitability and abundance modeling, respectively.

Results

Terrain Suitability

GBM, GLM, and RF runs were incorporated into the ensemble terrain model, with TSS and ROC scores showing excellent discrimination capacity (both > 0.9 on average) in recreating the *L. pertusa* distribution (Figure 2). Broad (1 km) BPI, slope, and depth were respectively the most important variables in the models, with high BPIs and slopes favored along with a depth range of ~200-900 m (Figure 3).

Broad expanses of habitat with high habitat suitability index (HSI) were predicted by the model (Figure 4a), including the ~200 km length of the Richardson Reef Complex (Figure 4b), notable mound provinces with many mounds topped by live *L. pertusa* (Figure 4c-d), and ridge and terrace features along the southwestern portion of the mapped area (Figure 4a). A portion of this habitat fell outside of the South Atlantic Fisheries Management Council's 'Stetson Banks-Miami Terrace Habitat Area of Particular Concern' (HAPC) (Figure 4a). Flats and valleys typically had lower HSI scores, while peaks, ridges, and slopes each contained a sizeable portion of the areas with HSI > 0.75 (Figure 5). Peaks, which included the tops of mounds of all size ranges shown in Figure 1, generally had the highest HSI.

Climatic Suitability

Notable changes in temperature and pH at *L. pertusa* sites were observed in the CMIP6 data by 2100, with more modest changes in dissolved oxygen and export carbon (Figure 6). In each scenario besides SSP 1-2.6, all sites with current *L. pertusa* populations experienced pH levels below which there is an analogue in the present-day SEUS. These conditions are highly unfavorable for the species based on the climatic response curves for the N. Atlantic (Figure 7). Only RF and GBM runs algorithms were used in the ensemble model due to biologically implausible responses in the GLM runs (Fig. S3.5). Despite this, TSS and ROC scores were > 0.9 (Figure 2c-d).

Ensemble projection onto the SEUS showed that almost all of the region from the shelf break to 1000 m currently has high climatic suitability index (CSI) scores (Figure 8). This was not the case in the climate change scenarios, however, which displayed spatiotemporal patterns of CSI declines (Figure 9). In the 2050 timepoint, ~20%

declines in CSI on the Blake Plateau were apparent near the shelf break and appeared regardless of SSP. Lesser CSI declines extended onto the Blake Plateau in a similar way in all scenarios, with the southernmost Blake Plateau a notable exception. At the 2100 timepoint, CSI declines were more severe and widespread. Near the shelf break, 80% declines were projected, and the extent of these declines increased with increasing emissions in the SSPs. In SSP1-2.6, however, the spatial extent and severity of these declines was notably lesser than the other three scenarios (Figure 9). A spatial pattern emerged with CSI declines significantly correlated with depth at both known L. pertusa sites (Linear Regression, p < 0.05; Figure 10a) and peak grid cells (Linear Regression, p <0.05; Figure 10b). The slope and variance explained by this relationship increased in each successively higher emissions scenario. At known Lophelia sites the regression slopes (i.e. the change in Δ CSI m⁻¹ depth) ranged from -2x10⁻⁴ in SSP 2-4.5 to -5.5x10⁻⁴ in SSP 5-8.5 while R² values in those were respectively 0.10 and 0.51; this corresponds to 2 and 5.5% worse respective declines in occurrence probability for every 100m in depth. At all 'peak' bathymorphon grid cells (n = 262,822), the regression slopes ranged from -2.8 x 10⁻⁴ in SSP 1-2.6 to -7.3 x 10⁻⁴ in SSP 5-8.5 while R² values in those respectively were 0.09 and 0.72; this corresponds to 2.8 and 7.3% worse respective declines in occurrence probability per 100m depth. A notable exception to this pattern occurred in SSP1-2.6 at known *Lophelia* sites where the relationship between depth and CSI declines were not significant (p > 0.05).

Abundance of L. pertusa

We found a wide range of coral cover in one-minute video segments from sites on the SEUS margin, ranging from zero to 80.4% cover at Blake Plateau Knolls (see

Figure 1d). Observed abundances were correlated with those predicted by terrainclimate suitability indices (RF Regression; p <<0.01, $R^2 = 0.51$, RMSE = 9.9; Figure 11a). Sites predicted to have > ~20% coral cover cease to exist in SSP5-8.5, but are still present in SSP1-2.6 (Figure 11b). The sites retaining relatively high cover were concentrated on the eastern Blake Plateau, with the Richardson Reef Complex maintaining the largest areal extent of high cover while more isolated individual mounds maintained lesser abundances (Figure S3.8).

Climate Velocity

Climate velocities varied throughout the region, with some of the highest velocities occurring near the shelf break where many of the shallower *L. pertusa* sites occur (Figure 12). Here, isotherms moved at approximately four kilometers per year (km·y⁻¹) from 2015 to 2100, while isotherms at sites on the eastern Blake Plateau (e.g. Richardson Reef Complex, Central Plateau Mounds) moved <1 km·y⁻¹. Velocities were also lower to the north, with sites off Cape Fear and Cape Lookout, Virginia experiencing similarly low velocities. While these sites are predicted to experience severe warming, they are closer to deep areas, lowering their velocities. Isotherm trajectories generally moved towards the eastern Blake Plateau, which is predicted to offer a more similar thermal environment to what currently occurs at the *Lophelia* sites < 500m.

Discussion

Habitat Extent

Our models project a previously underappreciated extent of *L. pertusa* habitat in the SEUS, including outside of the boundaries of protected areas that are closed to bottom-contact fishing due to the presence of CWC habitat (coral HAPC; Figure 1). While there are > 50,000 apparent CWC mounds in the roughly 500 km by 100 km region of highest suitability index scores - with many predicted to be topped with live coral by our models (Figure 4) – the coral mounds are predominantly unvisited. Most of the areas highly likely to contain live *L. pertusa* fell on peak and ridge bathymorphons (Figure 5), highlighting the importance of these features for conservation and exploration. Future expeditions to the region will help ground-truth the models presented in this study. It should be noted that preliminary versions of these models were used to predict the presence of the mounds in Central Plateau Mounds and Blake Plateau Knolls areas following the discovery of the Richardson Reef Complex in 2018, and the predictions were later confirmed on expeditions by the NOAA Ship Okeanos Explorer (Table S1.1). The spatiotemporal predictions arising from our models, in conjunction with the differences observed between bathymorphon types (Figure 5), allow for most complete broad-scale characterization of *Lophelia* habitat in the SEUS to date. Climate-driven distribution changes

Currently, most of the SEUS is has high CSI scores for *L*. pertusa (Figure 8). However, the waters overlying most of the suitable terrain will become oceanographically unfavorable by 2100 (Figure 9) with conditions outside of the current climate domain of the species (Figure S1.2). Shallower sites near the shelf break will be the first to feel climate-induced stress and will experience the most acute long-term reductions in CSI. These shallow sites retain relatively high CSI scores at the 2050

timepoint, however, suggesting that they could still be important transitory habitats as the spatiotemporal connectivity of the deep, eastern sites and the shallower sites diminishes. The relevance of such transitory habitats remains an open question but may prove important with increased temporal resolution in future HSM iterations (Huang et al., 2020).

Our results are consistent with the growing consensus on temperature as a major biodiversity driver in the ocean (Hunt et al., 2005; Yasuhara & Danovaro 2016; Doi et al., 2020). The Northwestern Atlantic is a warming hotspot (Saba et al., 2016). Accordingly, our models that a warming-driven deepening of the *L. pertusa* distribution will occur this century, and that the eastern Blake Plateau sites could serve as critical climate refugia. Interestingly, at the 2050 timepoint some CSI declines emerged but notable differences between scenarios did not (Figure 8-9). By 2100, the depth-driven refugia effect increased with increasing emissions. In fact, SSP1-2.6 was the only scenario in which many of the Blake Plateau coral sites retained CSI scores > 0.7 (Figure 8) and cover of ~20% (Figure 11). This suggests flourishing reef habitats typified by high live coral cover may face regional extirpation outside of this best-case scenario.

In other regions, e.g. the Mediterranean or Northeast Atlantic, losses of *L. pertusa* could be somewhat compensated by the proliferation of *Madrepora oculata*, which co-occurs at high densities with and is more thermally resilient than *L. pertusa* (Chapron et al., 2021). However, in the SEUS, *M. oculata* occurrences are sparser and growth of this species is unlikely to keep pace with *L. pertusa* declines. Relaxation of spatial competition from *L. pertusa* may also positively impact hexactinellid sponges

such as *Vazella pourtalesii*, which grows primarily on coral rubble in the SEUS and is predicted to expand its range with climate change (Beazley et al., 2021).

Coral mounds in the northern parts of the region e.g. known reefs off Cape Lookout and Cape Fear, Virginia are predicted to be amongst the sites with the fastest warming. These sites also currently experience frequent short-term (i.e. days-to-weeks) temperature spikes associated with Gulf Stream dynamics and/or downfluxes of surface-waters (Mienis et al., 2014). While similar temperature spikes have been observed at the deeper reefs in the region, their lower mean temperatures give them a buffer against shorter-term spikes. In 7-day experimental exposures to a temperature increase from 8°C to 14°C, corals from the Richardson Reef Complex exhibited increased respiration and excretion rates as well as higher rates of protein catabolism, indicating a distinct stress response, although all of the corals survived (Gómez et al. 2022). The incorporation of short-term variability in environmental data into models may reveal further vulnerable and/or refuge areas for CWCs but is currently beyond the purview of large-scale correlative models such as the ones we present here.

The deep, eastern Blake Plateau sites are especially notable because they may not only be climate refugia, but they currently contain thriving reef habitat with the highest coral cover and megafaunal biodiversity observed in the SEUS (Table S1.2; Gasbarro et al., unpubl.). This biodiversity may be impacted by our predicted declines, as it can correlate with the percent cover of live coral (Rowden et al., 2020; Price et al., 2021), although a mix of live and dead standing coral framework can also foster high biodiversity (Mortensen et al. 1995, Cordes et al. 2008). The losses of living coral we project may also lead to positive feedbacks of declining geomorphological complexity,

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as dead skeleton is more susceptible to dissolution and colonization by bioeroding species (Freiward & Wilson 1998; Hennige et al., 2020), flattening the habitat and making it even less suitable for CWC growth. While there was a significant positive relationship between our abundance predictions and those observed, there is a notable amount of variance, particularly at high cover values. This may be caused by species interactions or finer-scale environmental dynamics not incorporated in the present study. Despite these caveats, our models of *L. pertusa* abundance could have further utility when combined with laboratory and/or *in situ* carbon and nitrogen flux measurements to refine first-order estimates of the contributions of these ecosystems to margin- and even basin-wide functioning (De Clippele et al., 2020, 2021), and to better quantify the ecosystem services lost with projected coral declines (Cordes et al. 2021).

Somewhat surprisingly, oxygen and export carbon flux were not significant environmental drivers in our models. CWCs may be more resilient to deoxygenation if other conditions such as food supply are well met (Hebbeln et al., 2020), or oxygen limitation may be a regional phenomenon where availability is lower than in the North Atlantic, e.g. the North Pacific (Chu et al., 2019, Auscavitch et al., 2020). However, low oxygen concentrations have an effect on *L. pertusa* metabolic function (Dodds et al., 2007, Lunden et al., 2014) and the window for metabolically viable habitat will be narrowed by deoxygenation and warming in concert (Pörtner 2017; Deutsch et al., 2015; Ern 2019). Thus, more research is needed into the additive and/or synergistic effects of deoxygenation and warming. Our data suggests that the waters overlying the SEUS will remain normoxic until at least 2100, however.

Export carbon, on the other hand, will likely decline, particularly under the velocity core of the Gulf Stream (Figure 6; Figure S3.4). Despite energy availability's pervasive influence on seafloor biodiversity (Woolley et al., 2016), the degree to which CWC distributions are influenced by large-scale export carbon patterns is debated (Raddatz et al., 2014; Roberts & Kohl 2018, Portilho-Ramos et al., 2022). In addition, projected declines in open ocean productivity may be somewhat alleviated by increasing terrigenous carbon inputs (Lacroix et al., 2021). pH did not have as strong an effect on CSI as temperature. Its importance may have been masked by the correlation between the two, however (Figure S1.1). In the Mediterranean, where temperatures are also high, pH does appear to be a critical driver *L. pertusa* reef persistence (Matos et al., 2021). Indeed, there is both *in situ* and laboratory evidence of ocean acidification affecting the physiology of *L. pertusa* (Hennige et al., 2014; Kurman et al., 2017; Gómez et al., 2018; Dorey et al., 2020) and eroding the dead skeleton that forms the basis for their mound habitat (Hennige et al., 2015, 2020).

Our models generalize the climate response of the species from the whole of North Atlantic to the SEUS. It is possible that, because *L. pertusa* in this region – especially near the shelf break – are already exposed to some of the warmest temperatures recorded for this species, there may be a degree of acclimatization or adaptation as hypothesized in the Gulf of Mexico for acidification (Georgian et al., 2016b; Kurman et al., 2017) or for hypoxia on the west Angolan margin (Hebbeln et al., 2020). While acclimatization for marine calcifiers does not appear to be a viable strategy in shallow waters (Comeau et al., 2019), it may be achieved deeper where climate change proceeds slower. CWCs at these sites are possibly compensated for increased

metabolic demand in high temperatures by current-driven high food supply. Further slowing (Boers et al., 2021) or eastward movement (Matos et al., 2015) of the Gulf Stream may tip this delicate balance.

It is unknown whether climate corridors represented by the climate velocity trajectories will match dispersal corridors. That is, will larvae be able to track the lines in Figure 12b at sufficient speeds to allow recruitment to more accommodating habitat, or will the western-most populations face extirpation? Particle release models of the hexactinellid sponge Vazella pourtalesii, which commonly co-occurs with L. pertusa, suggest that larval exchange may be partially split among the eastern Blake Plateau and the shallower sites on the upper shelf (Wang et al., 2021). However, genetic evidence indicates that *L. pertusa* populations on the Blake Plateau are mixed (Morrison et al., 2011), suggesting connectivity whereas Northeast Atlantic populations appear to recruit locally with differentiation apparent between certain sub-populations (e.g. fjords versus offshore banks; Le-Goff Vitry et al., 2004). Larval dispersal models incorporating realistic ontogenetic behaviors have shown the connection between Lophelia reefs and oil rigs in the North Sea (Henry et al., 2018), and may be of use in describing connections between CWC habitat in the SEUS. If there is indeed genetic exchange between populations and a spatial gradient in the magnitude of climate change, the eastern Blake Plateau reefs meet both assumptions of the 'deep reef refuge hypothesis' that posits mesophotic coral ecosystems can be larval sources for shallow reefs (Bongaerts et al., 2010), suggesting the potential for an analogous process between CWC habitats separated by depth. Further studies of community similarity and identification of distinct faunal assemblages (e.g. Murillo et al., 2018) between reefs in

the region could also provide further tests of our hypothesized deep reef refuges and the utility of this framework in the deep ocean.

Our notable findings of potential climate refugia on the eastern Blake Plateau warrant additional protections including the extension of the HAPC to unprotected mound features (Figure 1), in order to prevent damaging these valuable ecosystems with bottom-contact fishing gear. Our study also highlights the utility of regional models in refining habitat suitability projections from coarser models (e.g. Davies et al., 2008; Morato et al., 2020), and of using multiple climate scenarios rather than a single business-as-usual scenario that is unlikely to occur (Burgess et al., 2022; Meinshausen et al., 2022). Indeed, visual comparison of our model outputs with those from Morato et al. (2020) suggest that more of the eastern Blake Plateau is within the current climatic niche of L. pertusa (Figure S3.11), which may also be due in part to our inclusion of additional occurrence locations. Our results corroborate their findings that a businessas-usual scenario (e.g. RCP 8.5 or SSP 5-8.5) would lead to precipitous declines in area on the Blake Plateau that L. pertusa is likely to occupy. We hope that this iterative refinement of model predictions continues as the spatiotemporal resolution of both climatic and mapping data increases, potentially revealing more localized refugia and aiding in conservation and exploration of the seafloor. Ultimately, this will lead to better accounting of the ecosystem services and vulnerable marine ecosystems both in US waters and other exclusive economic zones. Despite limited evidence that active restoration of CWC habitats is possible (Montseny et al., 2021), our results suggest that sharply curtailing greenhouse gas emissions is the only reasonable way in which the extent and health of these ecosystems will be maintained into the 22nd century.

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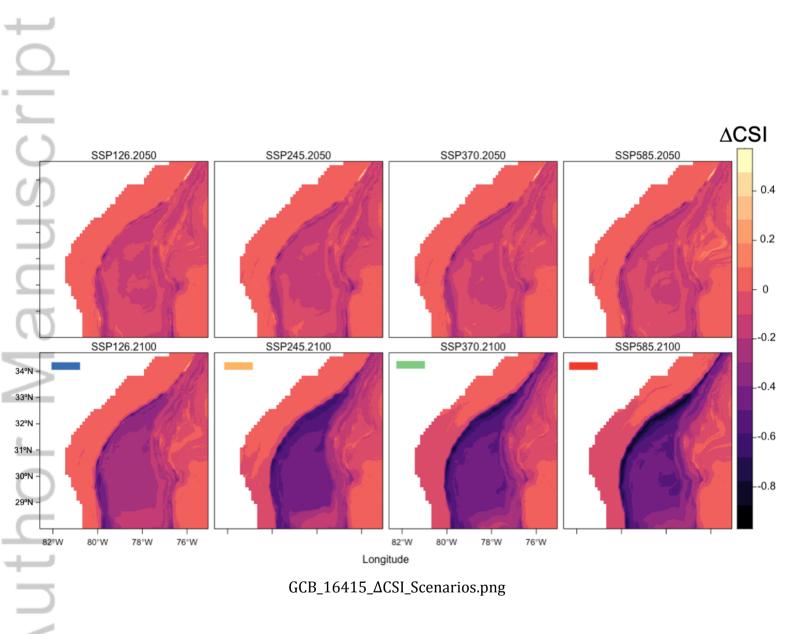
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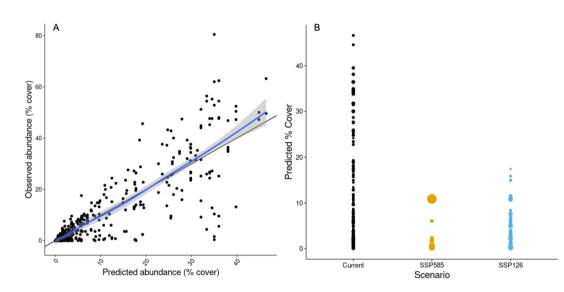
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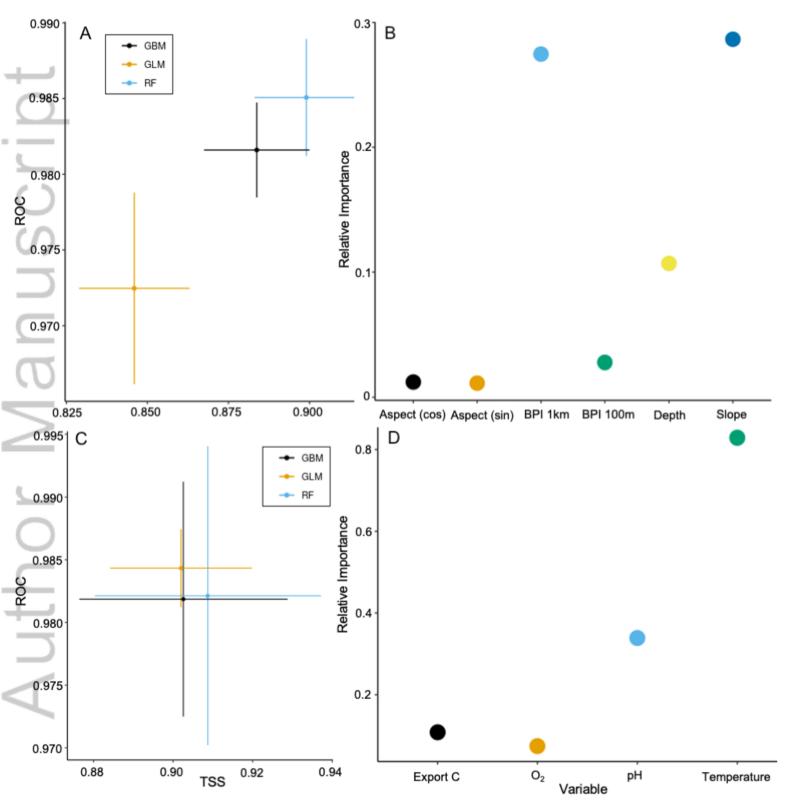
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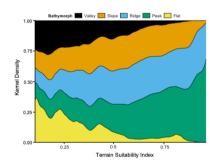


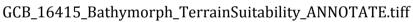


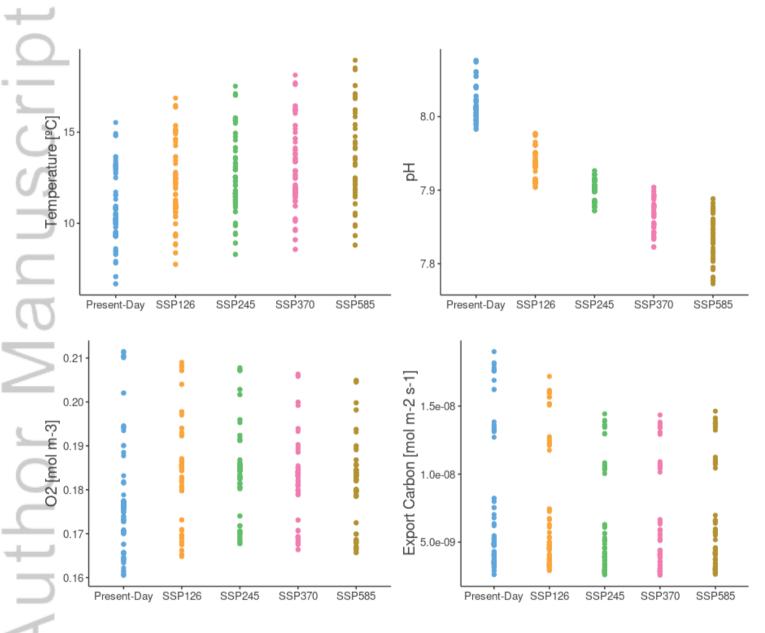
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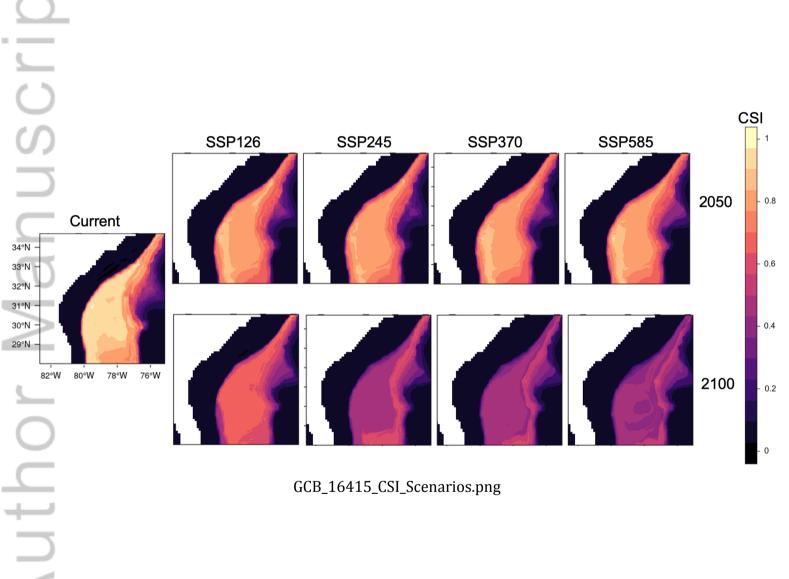
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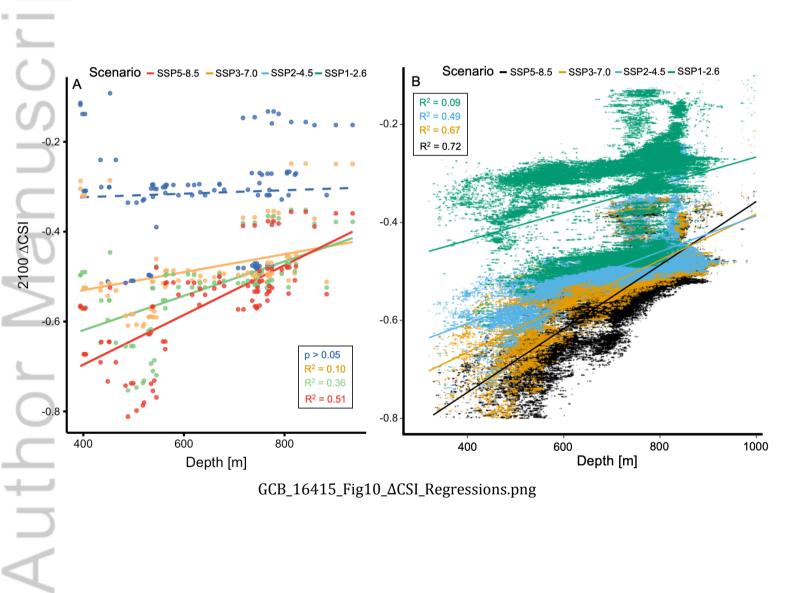


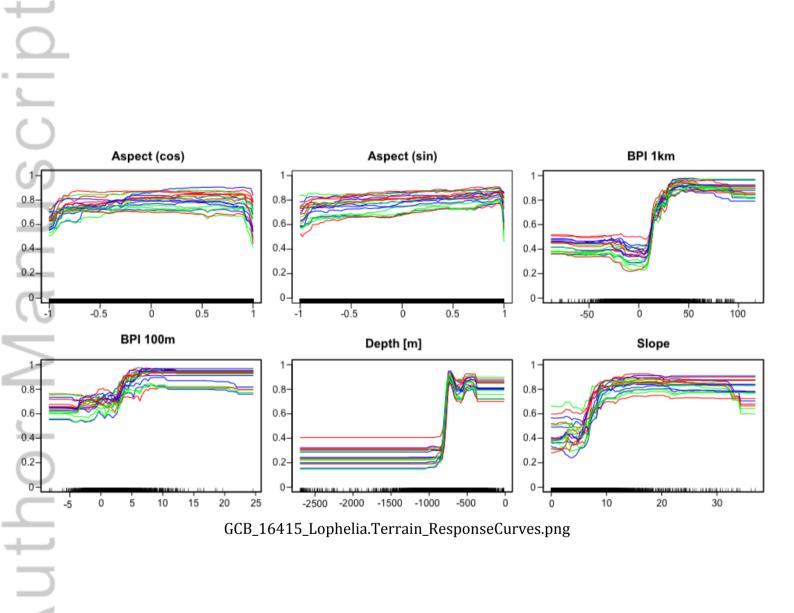


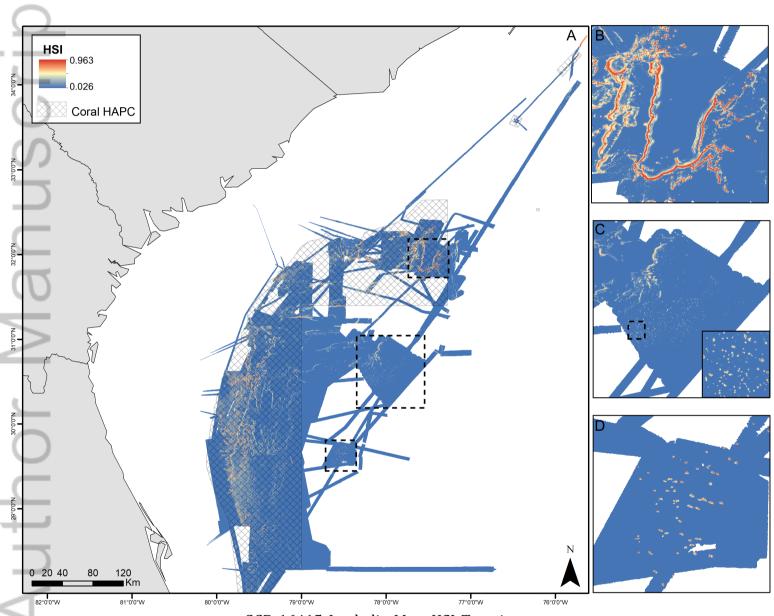
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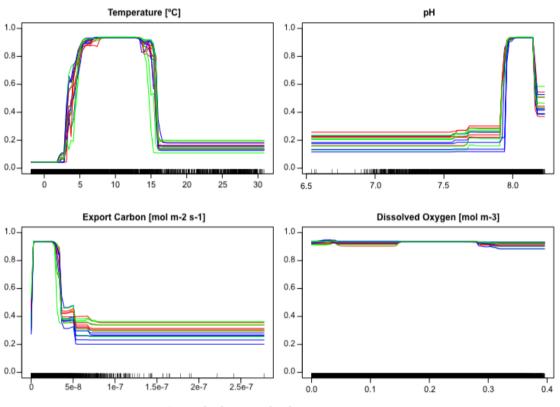


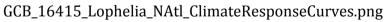


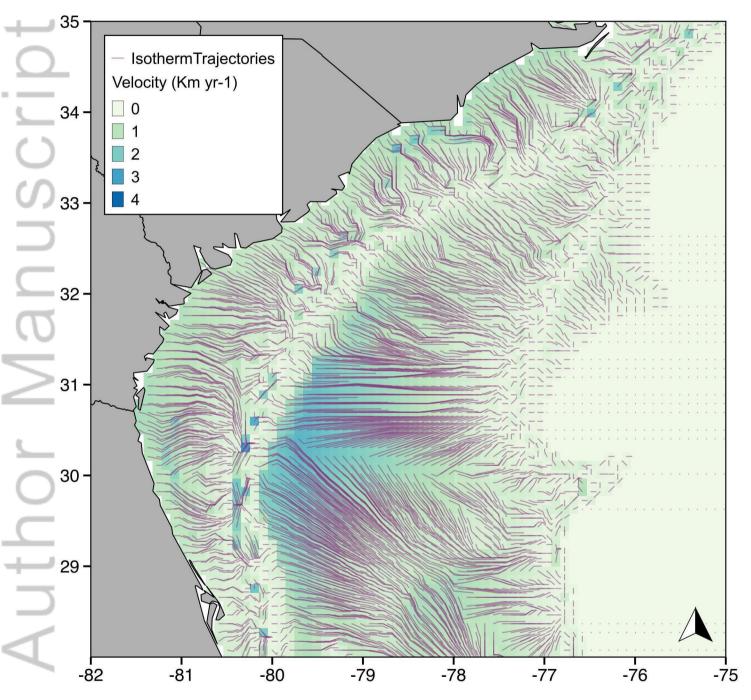


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