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Past and future decline of tropical pelagic biodiversity

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Abstract: A major research question concerning global pelagic biodiversity remains unanswered: when did the apparent tropical biodiversity depression [i.e., bimodality of latitudinal diversity gradient (LDG)] begin? The bimodal LDG may be a consequence of recent ocean warming, or of deep-time evolutionary speciation and extinction processes. Using rich time-slice datasets of planktonic foraminifers, we show here that a unimodal (or only weakly-bimodal) diversity gradient, with a plateau in the tropics, occurred during the last ice age and has since then developed into a bimodal gradient through species distribution shifts driven by postglacial ocean warming. The bimodal LDG likely emerged before the Anthropocene (here defined as ~1950) and perhaps ~15,000 years ago, indicating a strong environmental control of tropical diversity even before the start of anthropogenic warming. However, our model projections suggest future anthropogenic warming further diminishes tropical pelagic diversity to a level not seen in millions of years.

Significance Statement: We discovered that the tropical oceanic diversity depression is not a recent phenomenon nor very deep-time in origin, by using a comprehensive global dataset of the calcified shells of planktonic foraminifers, abundant unicellular organisms in the world's oceans which are exceptionally well-preserved in marine sediments as fossils. The diversity decline in the lowest latitudes may have started due to rapid post-ice-age warming around 15,000 years ago. Warming may by the end of this century diminish tropical oceanic diversity to an unprecedented level in human history.

One Sentence Summary: The bimodal pelagic latitudinal diversity gradient, at least in planktonic foraminifers, appears to have emerged from the glacial unimodal gradient through species distribution shifts probably driven by postglacial ocean warming.

Introduction

Latitudinal diversity gradients (LDGs), the equatorially-centred parabolic diversity patterns, have been described for over 200 years in terrestrial systems (1-4) and are also well-established in marine environments (5-7). However, there is an increasing recognition that marine LDGs, particularly those in open-ocean systems, tend to have an tropical diversity depression and thus to be bimodal (8-14).

This current tropical depression is consistent with present-day temperatures being beyond the upper physiological thermal tolerances of some species. An inability of species to tolerate high temperatures or sustained physiological stresses may cause shifts of their latitudinal ranges further poleward as the climate warms. Indeed, a near-future tropical biodiversity decline has been predicted with ongoing human-induced climate warming (15-19), and ecosystem-scale impacts of ocean warming are already evident (20-24).

Alternatively, or additionally, the current tropical dip in diversity could be explained through an evolutionary mechanism of higher speciation rates and/or lower extinction rates at the edges of the tropics (8, 13). Distinguishing the ecological and evolutionary time-scale processes

responsible for observed variations in the shape of marine LDGs is critical for assessing the outcome of biotic responses to rapid anthropogenic warming over the coming century (12). However, the lack of a standardized paleoecological baseline for the pelagic LDG has compromised separating whether the observed bimodality is caused by a rapid ecological response to ocean warming, by a longer-term and slower evolutionary process, or both (e.g., 14). While several paleontological studies have shown deep-time bimodal LDGs (25), they are not directly comparable to the present-day pelagic bimodality, or do not answer this question directly for various reasons, including that they tend to be from other (e.g., terrestrial) systems or too deep time to evaluate the hypothesis of rapid ecological response, and/or affected by limited data coverage.

The calcified shells of planktonic foraminifers, abundant and widespread protists in the world's oceans, are well-preserved in marine sediments and can thus provide a baseline for tracking trends in the LDG over the geologic past (26, 27). In addition, the relationship to temperature of planktonic foraminiferal diversity is consistent with that of many other open ocean organisms (5, 11, 28). Here we use global datasets of pre-industrial (broadly representing a late Holocene situation; see Materials and Methods) and last-ice-age planktonic foraminifers as well as a future diversity projection to provide empirical evidence that the tropical diversity depression is neither a recent anthropogenic phenomenon nor of deep-time origin. Rather, it was likely caused by a post-ice-age warming, suggesting a major role for distributional shifts driven by climate.

Results and Discussion

Diversity patterns with latitude and temperature

Our global analysis of planktonic foraminiferal diversity [calculated as species richness (Hill number, q = 0) and effective number of common species (Hill number, q = 1), see Materials and Methods] demonstrates that during the Last Glacial Maximum (LGM, ca. 21 kyr ago), the LDG was unimodal (or only weakly bimodal), whereas the pre-industrial LDG was bimodal with a distinct tropical diversity depression (Fig. 1, SI Appendix, Fig. S1, Tables S1, S2). This indicates that the strength of the bimodal LDG for planktonic foraminifers cannot be entirely due to long-term evolutionary processes because it was minimal during the LGM (Fig. 1, SI Appendix, Fig. S1, Tables S1, S2), and there have been no known global extinctions or speciations of any planktonic foraminiferal species since the LGM (29).

We propose that the cause of the bimodality may then be environmentally-driven extirpation and/or immigration. During warming, any diversity losses at higher latitudes (due to range shifts of species to even higher latitudes) are compensated for by the poleward movements of species from lower latitudes. However, in the tropics, such compensation due to species range shifts is not possible, resulting in an tropical diversity decline (15, 17, 30, 31).

It is unlikely that the tropical diversity depression is a very recent phenomenon originating in the Anthropocene, because we found that the pre-industrial LDG was already bimodal. Thus, the bimodal LDG most likely developed during the post-LGM warming, with a 5.2 % loss in the

mean projected species richness since the LGM at the equator (calculated based on the mean predictions within ±1 degree latitude; Fig. 1).

The LDG exhibited a tropical plateau (or weak bimodality) during the LGM (Fig. 1, SI Appendix, Fig. S1) indicating an approach towards diversity saturation (at or beyond the optimum in the unimodal temperature-diversity relationship; see the next paragraph) with relatively low maximum global sea temperature. The distinct tropical diversity decline may have begun ~15,000 years ago, given that a rapid postglacial warming started at that time (32). The duration of glacial periods has been much longer than that of interglacial periods during the late Quaternary. Therefore, the tropical thermal niches of marine organisms may be optimized to the maximum temperatures of glacial periods, leading to tropical diversity depressions during warm periods, given marine niche conservatism is known to have existed during late Quaternary climate changes (33). As a bimodal LDG is known to be present during the last interglacial (in corals; 34), it is likely that the bimodal LDG has appeared repeatedly during warm interglacial periods during the late Quaternary, and weakened during glacial periods. Species adapted to very warm temperatures existed during the Pliocene, the major previous warmer-than-present period, but significant extinctions of these species are known during the Plio-Pleistocene cooling (27). Note that pre-Plio-Pleistocene Phanerozoic LDG are also known to be dynamic (14, 35-37) though the underlying mechanism may be different.

Sea surface temperature has been and is unimodal with latitude (Fig. 2d, but see the next paragraph for the equatorial upwelling zone). It is also predicted to remain unimodal under the RCP 8.5 'business-as-usual' climate warming scenario in 2091-2100 (‘2090s’ hereafter), with ~0–4 degrees warming relative to the pre-industrial control (PIC) (Fig. 2). The magnitude of the predicted warming from the PIC to the RCP 8.5 2090s will be larger (and much more rapid) than that from the LGM to PIC (Fig. 2), particularly in the tropics. The unimodal (or only weakly bimodal) LDG during the LGM and the bimodal LDG during the pre-industrial time period reflect a positive temperature-diversity relationship from -2°C to 20°C and a negative relationship beyond that, especially beyond 25°C and for species richness (SI Appendix, Fig. S2). Thus, the present reduction of species diversity in the tropics is likely due to high sea temperatures (SI Appendix, Fig. S2), a thermal response also identified in other pelagic groups (38). Such very high temperatures (those exceeding 25°C) did not exist in any latitudinal band during the LGM (Fig. 2). Supporting our interpretation is the observation that planktonic foraminifer species tend to have optimum temperature ranges at ~20–30°C, with a sharp drop in their growth rates above these temperatures, showing a high-end temperature threshold of thermal performance curves (19, 39, 40). Using the relationship between sea surface temperature and diversity for both time periods (LGM and PIC), we predict a more than 15 % diversity loss at the equator (calculated based on the mean predictions within ± 1 degree latitude) within this century under the ‘business-as-usual’ climate warming scenario (Fig. 1, SI Appendix, Fig. S1). In comparison, only ~5 % diversity loss at the equator has been observed between the LGM and PIC (Fig. 1, SI Appendix, Fig. S1), indicating the potential for a three times greater reduction over the coming century. It is also noteworthy that corals had a bimodal LDG in the last interglacial, a warmer-than-present time period (34). Thus, we may see tropical diversity decline not only in planktonic foraminifers but also in other taxonomic and functional groups with further future warming.

In the equatorial upwelling zone of the eastern Pacific Ocean (especially at ~100–120 ºW; Fig. 2b), sea surface temperature is lower than that in adjacent higher-latitude (e.g., 5–10 ºN and ºS) tropical waters, which may affect species diversity. Indeed, the equatorial diversity is higher than that at 5–10 ºN and ºS in the eastern Pacific at ~100–120 ºW (Fig. 1b). Thus, in the present-day ocean, the equatorial upwelling zone with lower temperature than adjacent higher-latitude tropical waters may be within or close to the optimum temperature range of many species and act as a refugium. In the future warmer ocean, however, temperature will be beyond the optimum temperature range even in the equatorial upwelling zone (Fig. 2c) and the refugium will disappear (Fig. 1c). Nonetheless, the equatorial upwelling zone does not affect our major results, because the low temperature zone related to the equatorial upwelling is limited to a very narrow equatorial band of the eastern Pacific Ocean. The analyses of just the Atlantic Ocean, which lacks a distinct equatorial temperature decline, show the same basic results (see Materials and Methods).

Higher latitude, especially temperate, diversity increases from the LGM to PIC and from the PIC to RCP 8.5 2090s and offsets the tropical diversity decline (Fig. 3). The temperate peaks of diversity shift poleward in the comparison between the PIC and RCP 8.5 2090s (Fig. 3b) relative to that between the LGM and PIC (Fig. 3a), indicating that future warming will further enhance poleward species range shifts. Both the tropical diversity decline and temperate diversity increase from the PIC to 2090s would be reduced with the low-emission scenario RCP 2.6 relative to the business-as-usual scenario RCP 8.5 (Fig. 3c). The subpolar diversity decline from the PIC to 2090s (negative Δ diversity peak at ~50–60 ºN: Fig. 3b, c) is probably due to projected subpolar North Atlantic cooling related to a collapse of the local deep-ocean convection (41-43).

Beta-diversity and the process of diversity change

Beta-diversity quantifies how species composition changes in space and time; for example in response to temperature gradients and ocean warming. We divided beta diversity into turnover and nestedness components (Fig. 4, see Materials and Methods). Turnover occurs with species replacement along an environmental gradient, and nestedness indicates species loss without replacement; i.e., when an assemblage is a subset of a more species rich neighbouring biota. The relative contributions of turnover and nestedness components had positive and negative peaks respectively in the tropics during the LGM, showing unimodal and inverse unimodal LDGs (Fig. 4). Since then the peaks have moved poleward towards the edges of the tropics, showing bimodal and inverse bimodal LDGs during the pre-industrial time (Fig. 4). The tropical peak of the pre-industrial inverse bimodal nestedness LDG is due to a reduction of species, presumably those most sensitive to the warming. In other words, the pre-industrial tropical assemblage has lost species and has become more of a subset of the adjacent higher-latitude tropical assemblages. The peaks in relative contribution of turnover (positive) and nestedness (negative) to beta diversity at the edges of the pre-industrial tropics (Fig. 4) indicate distributional shifts of some of tropical species, which had an equatorial distribution during the LGM, towards higher latitudes (SI Appendix, Fig. S3). Overall, 23 of 27 species extended their interquartile range (75-25 percentile) and shifted southern and northern edges of distributions (97.5-2.5 percentiles)

poleward since the LGM, and 6 of 27 species show much stronger bimodal latitudinal distributions of their occurrence density in the PIC than in the LGM, which is probably responsible for the observed bimodal PIC LDG (SI Appendix, Fig. S3).

Future scenario

A future tropical diversity depression has not only been predicted for planktonic foraminifers but also for other taxonomic and functional groups (Fig. 1; 8, 9, 11, 18, 19). Planktonic foraminifer diversity is known to track marine and especially pelagic diversity (5, 26). Given the exceptional fossil record of planktonic foraminifers used here as an ideal model system and the fact that most marine organisms have poor fossil records, our findings may further apply to other taxonomic groups. For example, Kaschner et al. (44) suggested a reduction of tropical and an increase in temperate diversity in marine mammals under a warming scenario.

In a warmer pelagic world, temperate regions will hold more tropical species, and polar regions more temperate species, as they change their distributions to live within their optimum temperature niches (16). However, tropical regions will have no source for such immigrants (16-18). Our study shows that this tropical dead-end causes a local diversity reduction of planktonic foraminifers between 20 °S and °N. The situation will worsen with continued global warming in the coming decades, particularly without appropriate mitigation of greenhouse gas emissions (Fig. 3c). This tropical pelagic diversity decline likely emerged before industrialization and the Anthropocene and perhaps during the onset of the postglacial warming ~15,000 years ago. Future anthropogenic warming may diminish tropical diversity to a level not seen in millions of years.

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Competing interests: Authors declare no competing interests.

Data and materials availability: All data is available through the main text, the supplementary materials, and references therein.

Materials and Methods

Foraminifera

We used exceptionally comprehensive global census datasets of planktonic foraminifera, the ForCenS (45) and the MARGO (46, 47) compilations, for "present-day" pre-industrial (see below) and LGM LDG reconstructions, respectively. The databases comprise specimens collected using a constant 150 µm sieve size (see Yasuhara et al., 2012 for discussion on the sieve size). We consider *Globigerinoides ruber* pink and white as separate species. We merged *Globorotalia menardii* and *Globorotalia tumida*. P/D integrade is merged with *Neogloboquadrina incompta*. Otherwise, we used species only and did not use subspecies or categories including multiple species. *Globorotalia crassula* was removed from the datasets, because it already became extinct ~0.9 million years ago (28, 29). We also removed small, rare, and/or taxonomically obscure species (*Tenuitella iota*, *Berggrenia pumilio*, *Dentigloborotalia anfracta*, *Globorotalia cavernula*, *Globigerinita minuta*, and *Globorotalia ungulata*) following Siccha and Kucera (45). Eventually, we used these 34 species: *Beella digitata*, *Candeina nitida*, *Globigerina bulloides*, *Globigerina falconensis*, *Globigerinella adamsi*, *Globigerinella calida*, *Globigerinella siphonifera*, *Globigerinita glutinata*, *Globigerinita uvula*, *Globigerinoides conglobatus*, *Globigerinoides ruber* pink, *Globigerinoides ruber* white, *Globigerinoides tenellus*, *Globoconella inflata*, *Globoquadrina conglomerata*, *Globorotalia crassaformis*, *Globorotalia hirsuta*, *Globorotalia menardii+tumida*, *Globorotalia scitula*, *Globorotalia theyeri*, *Globorotalia truncatulinoides*, *Globorotaloides hexagonus*, *Globoturborotalita rubescens*, *Hastigerina pelagica*, *Hastigerinella digitata*, *Neogloboquadrina dutertrei*, *Neogloboquadrina incompta*,

Neogloboquadrina pachyderma, *Orbulina universa*, *Pulleniatina obliquiloculata*, *Sphaeroidinella dehiscens*, *Trilobatus sacculifer*, *Turborotalita humilis*, *Turborotalita quinqueloba*. After removing duplicated samples, the pre-industrial ForCenS and LGM MARGO datasets include 4138 and 1442 samples, respectively, with >~300 specimens per sample for most samples (45-47). Given generally slow sedimentation rate in the deep sea, the ForCenS coretop "present day" dataset probably represents mostly the late Holocene (= the last few thousand years) but pre-industrial and pre-Anthropocene (48). Although small proportion of specimens would be from the Anthropocene, they should be negligible, given time averaging of a few thousand years. It is unlikely that the bimodal LDGs are artifacts of sampling biases (49), because tropical regions are well sampled in our datasets (Fig. 1, SI Appendix, Fig. S1), and the less sampled LGM dataset does not show a remarkably bimodal LDG.

Temperature

We used the 3-D, fully coupled Earth system models GISS-E2-R from the Goddard Institute for Space Studies, IPSL-CM5A-LR from the Institut Pierre Simon Laplace, and MPI-ESM-P from the Max Planck Institute to calculate the ensemble average of annual mean sea surface temperature (SST) for the last 100 years during the LGM (50-52). For the last 100 years of the pre-industrial control (PIC) scenario and the years 2091-2100 (2090s) projections (RCP 8.5 a.k.a. business-as-usual scenario and RCP 2.6 with appropriate mitigations of carbon dioxide emission), we use earth system model simulations from the Geophysical Fluid Dynamics Laboratory's GFDL-ESM-2G, the Institut Pierre Simon Laplace's IPSL-CM5A-MR, and the Max Planck Institute's MPI-ESM-MR to calculate the ensemble average of the annual mean SST (51, 53, 54). We re-projected the SST layer of each Earth system model to 0.5 by 0.5-degree grids based on bilinear interpolation and then calculated the multi-model average of each interpolated grid. All Earth system models are part of the Coupled Model Intercomparison Project Phase 5 (CMIP5) and were downloaded from the Earth System Grid Federation (ESGF) Peer-to-Peer (P2P) enterprise system (https://esgf.llnl.gov/).

Statistical modeling

For diversity measures, we used Hill numbers (55), ${}^{q}D = (\sum_{i=1}^{S} p_i^q)^{1/(1-q)}$, where S is the number of species in a site and p_i is the relative abundance of the ith species. The (larger) value of order q discounts the rare species and thus emphasizes the abundant species. When q = 0, all species have equal weight and ${}^{q}D$ is equivalent to species richness. Where q approaches 1, the derived mathematical expression of Hill numbers (${}^{1}D$) is given as ${}^{1}D = exp\left(-\sum_{i=1}^{S} p_i^q log(p_i)\right)$. Because the equation gives more weight to common species (with higher relative abundance), it can be interpreted as the effective number of equally-abundant and common (typical) species in a community (56). Both measures were very similar in our results, so we present the Hill number of order q = 0 in the main text because species richness is the most intuitive and commonly used measure of diversity, and q = 1 (the exponential form of the Shannon index) in the SI Appendix.

We decomposed beta-diversity (multiple-site Sorensen dissimilarity) which is influenced by turnover and species richness, into spatial turnover (also called Simpson's dissimilarity index) and nestedness components (57, 58). The beta-diversity measures and partitions were conducted over a one-degree-latitude moving window. Within each moving window, five sites were randomly resampled (with replacement) for 1000 times to estimate the mean and standard deviation. Windows with less than five sites were omitted from the calculations. The same

analyses were tested across one- to five-degree-latitude moving window and show consistent latitudinal patterns in beta diversity (SI Appendix, Fig. S4).

The latitudinal gradients of diversity were fitted by a generalized additive model (GAM) with a quasipoisson error distribution, and a thin plate regression spline for the LGM and PIC datasets. We also used a GAM to fit the LGM or PIC SST to their observed Hill numbers (e.g., species richness or effective number of common species) to visualize the thermal gradient of diversity. Finally, we constructed a third type of GAM using SST, longitude, latitude (and their interaction), the ocean basins (i.e., Atlantic, Pacific, Indian, Arctic, and Southern Oceans) where the samples were collected, and time (LGM and PIC) as predictor variables to account for spatial and temporal diversity variations and to project the future distribution of species richness based on the ensemble average of projected SST under RCP 8.5 and RCP 2.6 in the 2090s. The basis dimensions in the GAMs was chosen ($k = 5$ or 6) to generate smooth curve fit for ease of interpretation; nevertheless, the fitted lines are in general agree to the GAM with automatic selection of k.

All statistical analyses were performed with R version 3.5.1 (59). Hill numbers and multivariate analysis used the *vegan* package (60), beta-diversity the *betapart* packages (61), GAMs the *mgcv* package (62), and GIS mapping and data visualization used the *raster*, *sp* and *ggplot2* packages (63-65). A significance level of $\alpha = 0.05$ was applied to all statistical tests. All model residuals were checked by standard diagnostic plots (i.e., residual vs. fitted values, qq plot) for assumptions of homogeneity, independence and normal distribution, and by Moran's I test, Moran's I spatial correlogram and variogram for spatial autocorrelation (66). The assumptions of homogeneity, independence and normal distribution were reasonably met. Spatial autocorrelations in the model residuals were detected at distances up to 2791 km for the LGM and 1696 km for PIC species richness (Hill numbers of order $q = 0$) and up to 785 km for the LGM and 1229 km for PIC effective number of common species (Hill numbers of order $q = 1$).

Dissolution of planktonic foraminiferal shells and upwelling may affect diversity. To demonstrate that the diversity patterns were not affected by dissolution or upwelling, we ran the same analysis for three subsets; namely samples with water depth less than 3000 m, those from Atlantic Ocean only, and those excluding all coastal ecoregions, and thus coastal upwelling areas (67). The shallow-depths and Atlantic subsets have higher calcium carbonate saturation state, and thus better foraminiferal preservation (than the whole dataset including deeper depths and other oceans than the Atlantic). The Atlantic Ocean does not have distinct low temperature zone related to the equatorial upwelling, compared to the Pacific Ocean (Fig. 2). The results of these subsets remain qualitatively the same (SI Appendix, Figs S5, S6, S7), showing that our results are not artefacts of preservation or affected by upwelling.

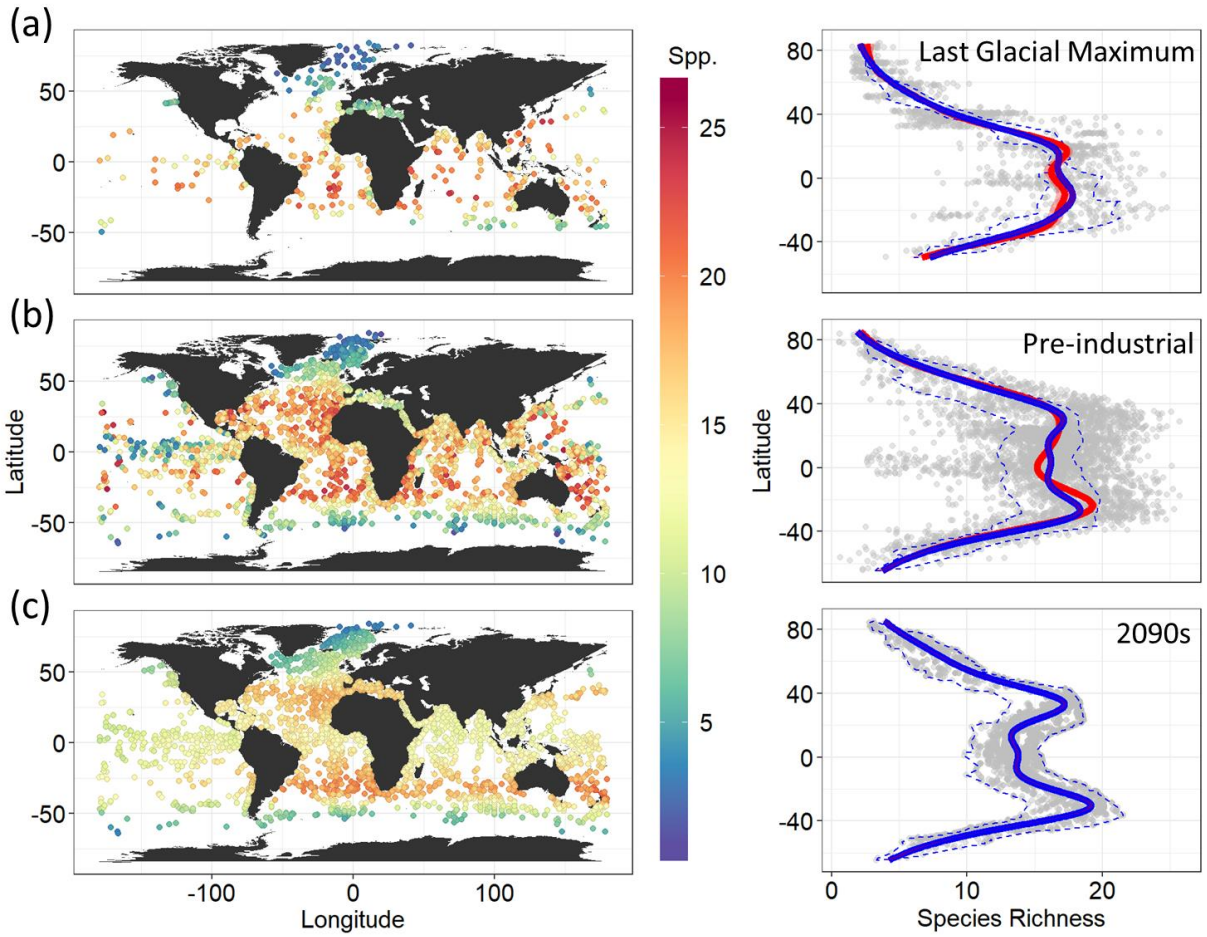


Fig. 1. Species richness of planktonic foraminifers during the (a) Last Glacial Maximum (LGM), (b) pre-industrial control (PIC), and for (c) 2091-2100 (2090s) as maps and latitudinal gradients. The observed diversities (LGM & PIC, gray symbols) were modeled by sea surface temperature, coordinates, and ocean basin using a Generalized Additive Model (GAM). The diversities in 2090s (gray symbols) were predicted from the same set of variables with future sea surface temperature (based on RCP 8.5). The predicted latitudinal diversities for the three time periods (enclosed by blue dashed lines) were smoothed by a GAM to show latitudinal diversity gradients (blue lines). The latitudinal gradient of observed diversities during the LGM and PIC were also fitted by a GAM and shown as the red lines with the shaded areas indicating the 95% confidence intervals (small and not visible in the PIC panel). LGM and PIC observations points have been had a small amount of jitter added on the x-axis to make them visible when overlapping. See SI Appendix Fig. S1 for empirical and projected diversities using a Hill number of order q = 1.

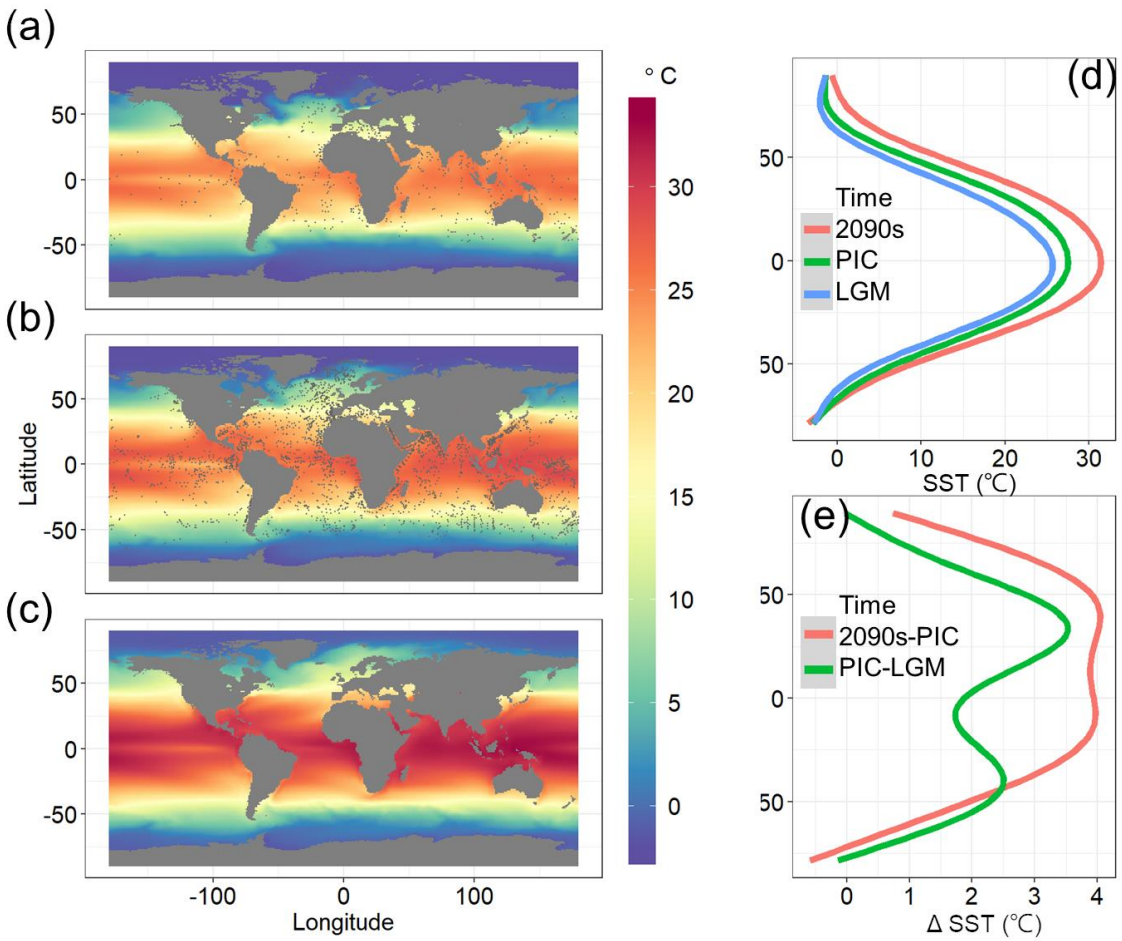


Figure 2. Maps and latitudinal gradients of the projected ocean sea surface temperature (SST) during the (a) LGM, (b) pre-industrial control (PIC), and (c) 2091-2100 (2090s) based on RCP 8.5. The latitudinal SST (LGM: blue; PIC: green; 2090s: red) and ΔSST (warming from the LGM to PIC as green and from the PIC to 2090s as red) are smoothed by a GAM and shown in panels (d) and (e), respectively. Grey dots in panels (a) and (b) indicate sample locations.

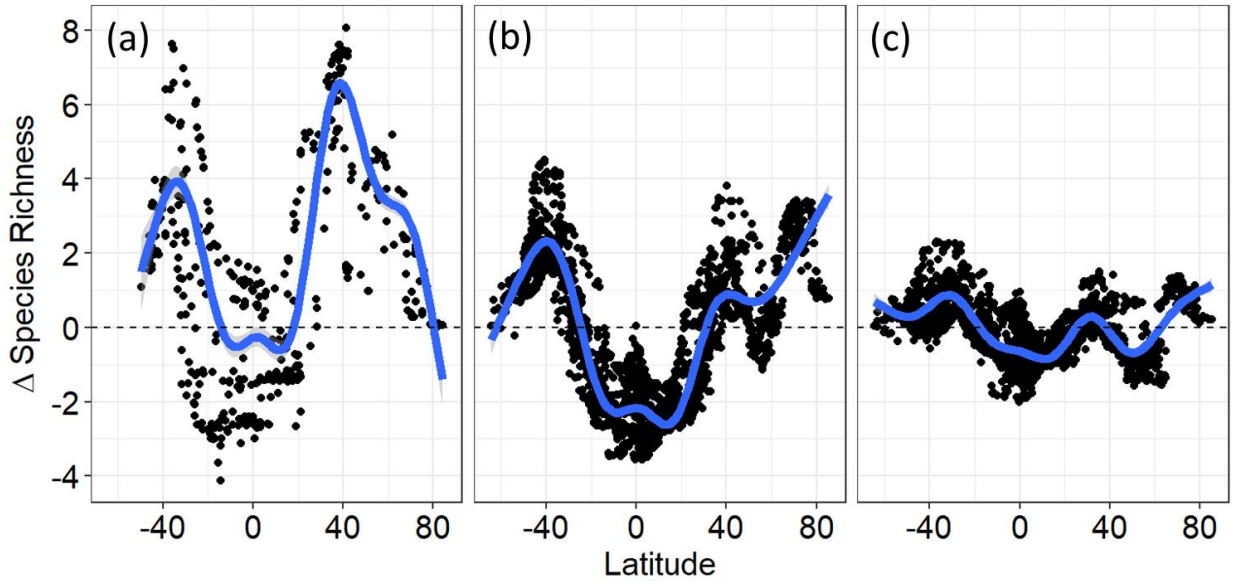


Figure 3. Changes in predicted species richness (Δ species richness) from the LGM to PIC (a), from the PIC to RCP 8.5 2090s (b), and from the PIC to RCP 2.6 2090s (c). Species richness was predicted using sea surface temperature for LGM, PIC, RCP 8.5 2090s, and RCP 2.6 2090s. The Δ species richness were calculated for the LGM samples in panel (a) and PIC samples in panel (b) and (c), and smoothed by a GAM [blue lines with the gray shaded areas indicating the 95% confidence intervals that are small and not visible in the panels (b) and (c)].

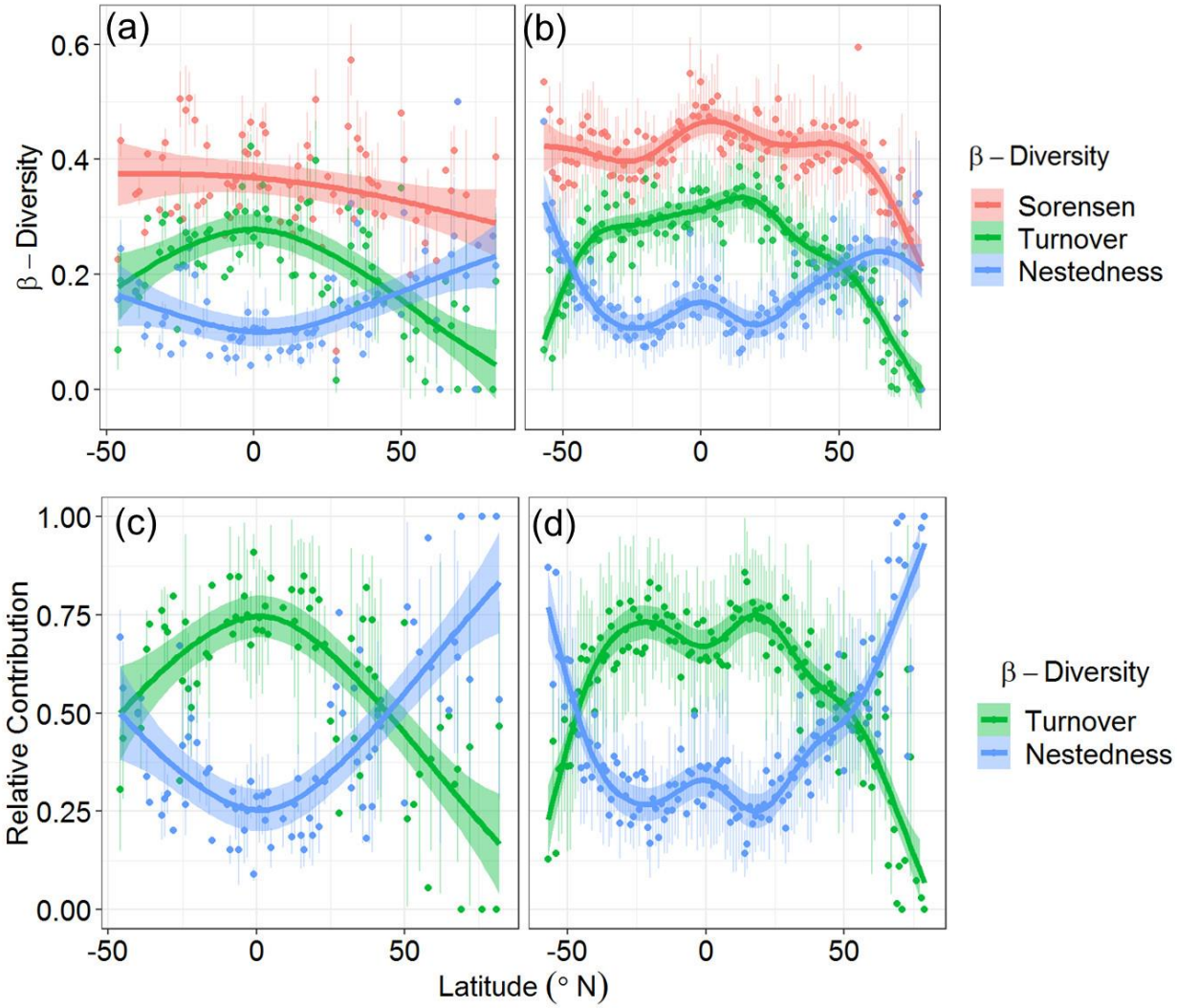
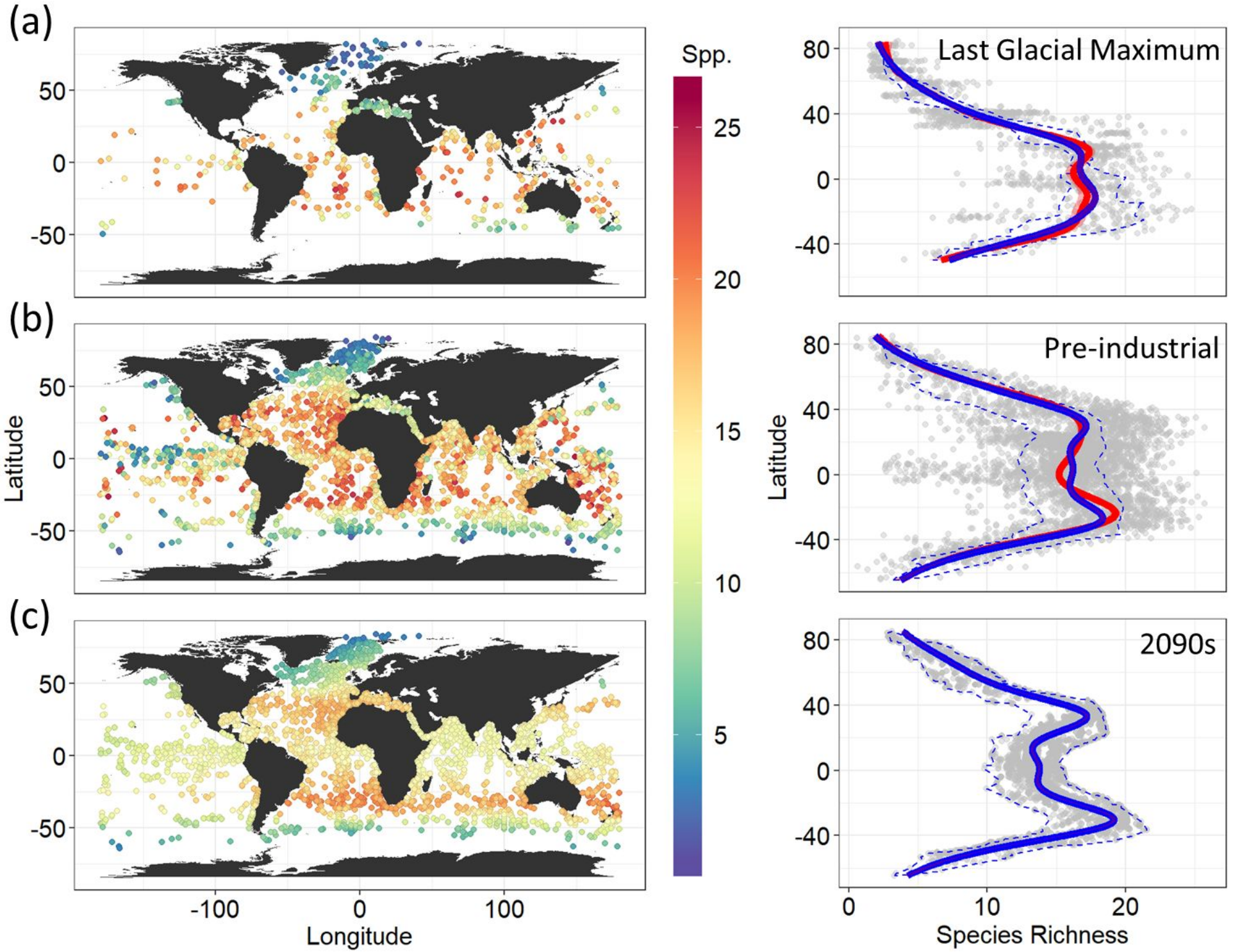
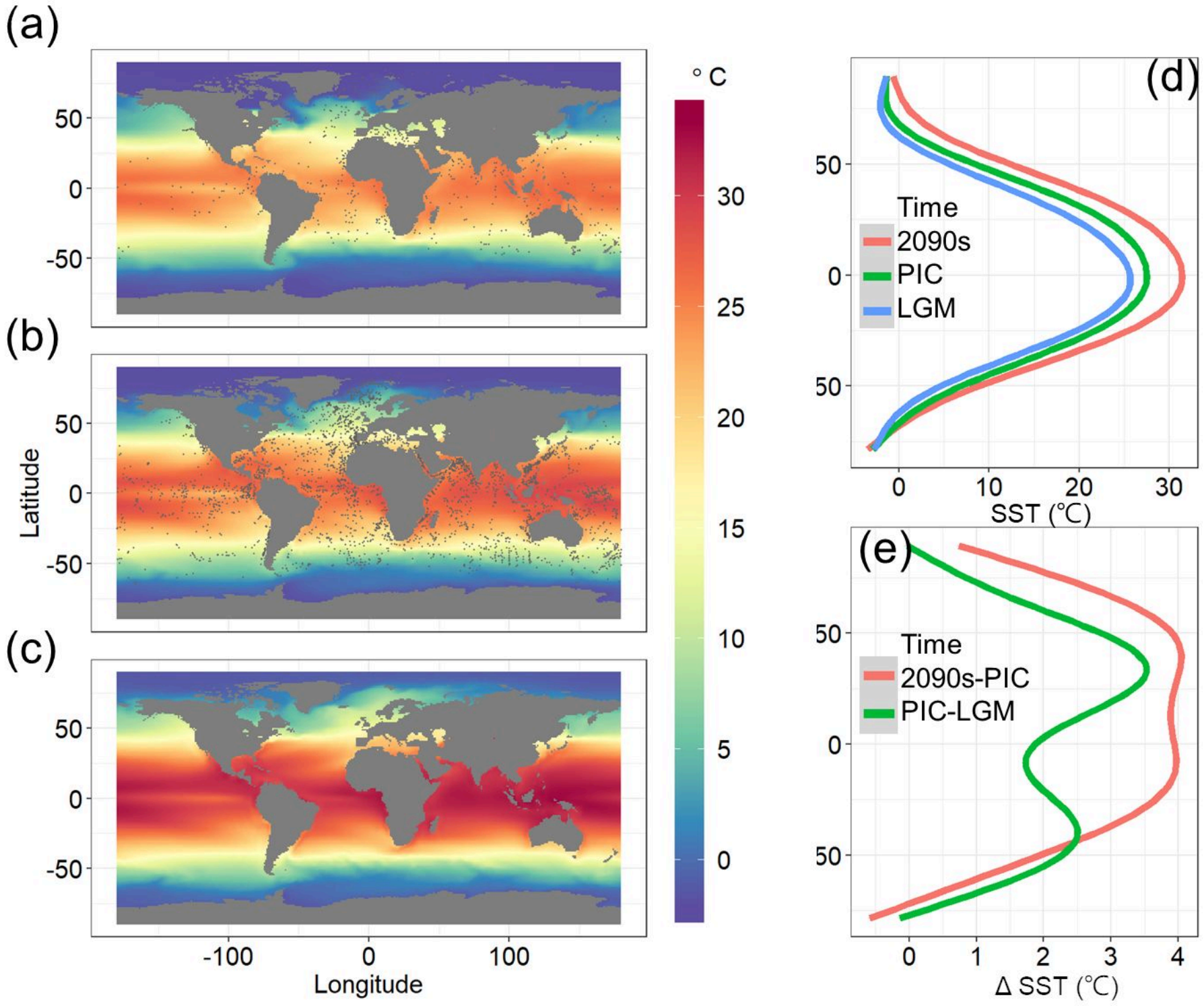
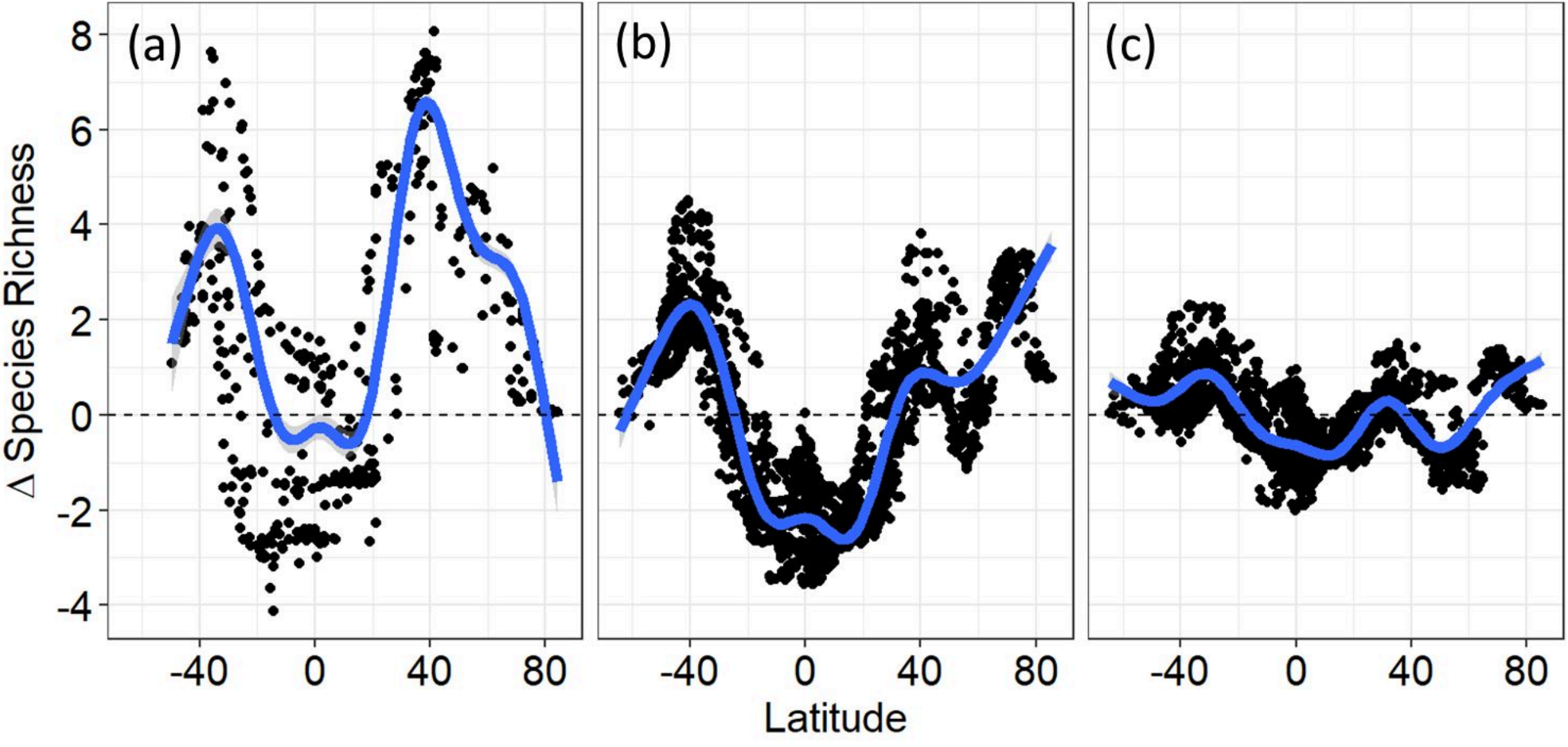
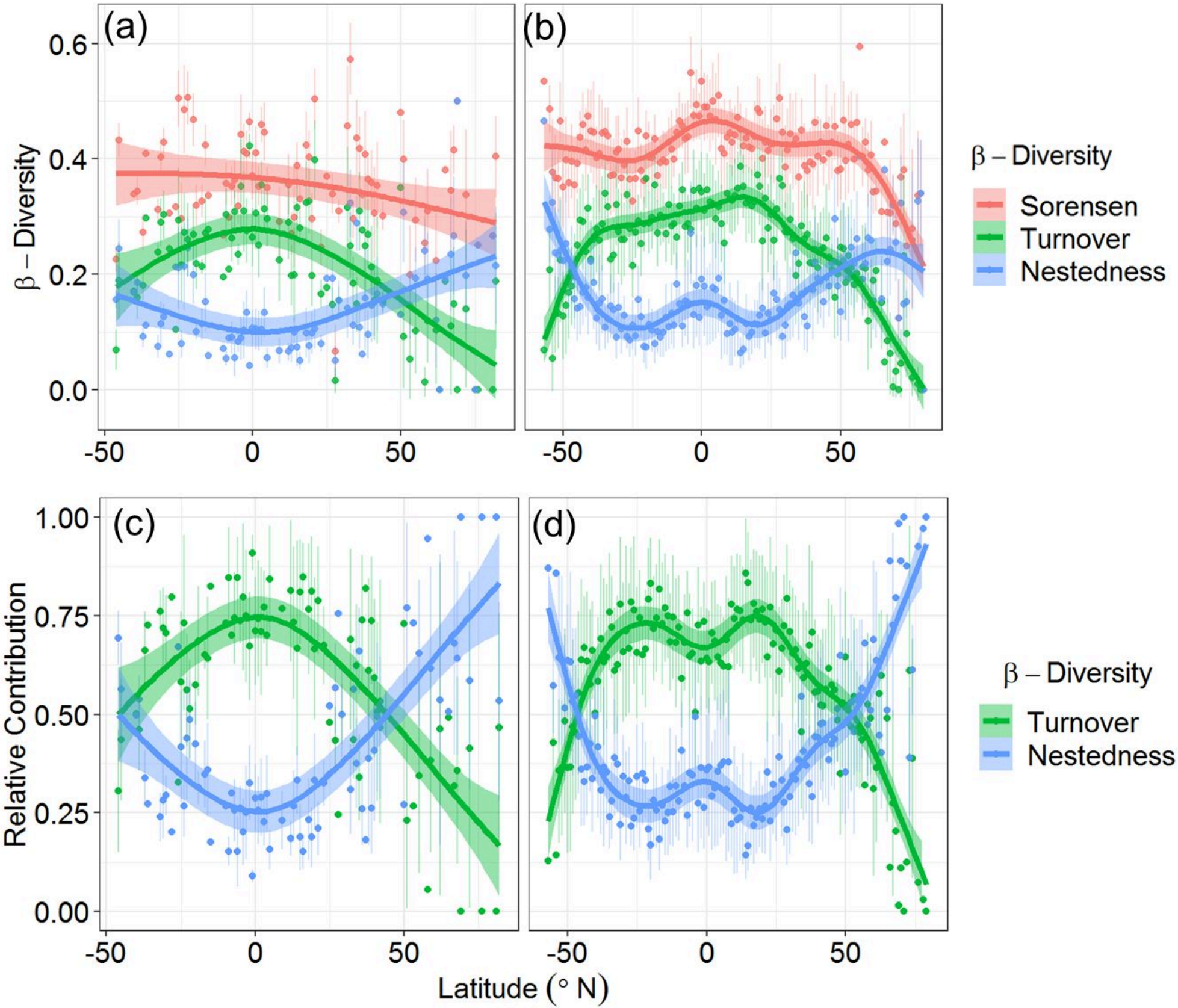


Figure 4. The latitudinal gradients in beta diversity during (a) the LGM and (b) the pre-industrial periods. The total beta-diversity, i.e. Sorensen dissimilarity (red), was separated into turnover (green) and nestedness (blue) components. Panels (c) and (d) show the relative contribution of the turnover (green) and nestedness (blue) components to total dissimilarity for the LGM and pre-industrial periods, respectively.


(a)
(b)
(c)
Spp.
25
20
15
10
5
Last Glacial Maximum
Pre-industrial
2090s
Latitude
Longitude
Species Richness
80
40
0
-40
50
0
-50
-100
0
100
10
20


(a)
(b)
(c)
(d)
(e)
°C
30
25
20
15
10
5
0
50
0
-50
Latitude
-100
0
100
Longitude
Time
2090s
PIC
LGM
SST (℃)
Time
2090s-PIC
PIC-LGM
Δ SST (℃)


(a)
(b)
(c)
Δ Species Richness
8
6
4
2
0
-2
-4
-40
0
40
80
Latitude


(a)
(b)
(c)
(d)
β – Diversity
Relative Contribution
Latitude (° N)
0.6
0.4
0.2
0.0
1.00
0.75
0.50
0.25
0.00
-50
0
50
β – Diversity
Sorensen
Turnover
Nestedness
β – Diversity
Turnover
Nestedness