

Global distribution and environmental correlates of marine bioturbation

Shuang Zhang^{1,2*}, Martin Solan³, Lidya Tarhan^{2**}

¹Department of Oceanography, Texas A&M University, 3146 TAMU, College Station, TX 77843, USA.

²Department of Earth and Planetary Sciences, Yale University, P.O. Box 208109, New Haven, CT 06520, USA.

³Ocean and Earth Science, National Oceanography Centre Southampton, University of Southampton, Waterfront Campus, European Way, Southampton, SO14 3ZH, UK.

Correspondence: *shuang-zhang@tamu.edu; **lidya.tarhan@yale.edu

Paper's lead contact: shuang-zhang@tamu.edu

X (formerly Twitter): [@caceelab](https://twitter.com/caceelab) (<https://twitter.com/caceelab>)

Summary

The activities of marine sediment-dwelling invertebrates play a fundamental role in mediating major biogeochemical cycles and have profoundly shaped the evolution of marine systems. Yet there remains a paucity of global marine data describing bioturbation intensities and mixed layer depths, and interrogating how these vary with multiple environmental and ecological factors at a system scale. We applied an ensemble of tree-based machine-learning techniques to resolve a global map and determine the environmental and ecological correlates most closely associated with bioturbation. We find that bioturbation intensity and the depth of the sediment mixed layer each reflect different associations with a consortium of environmental and ecological parameters, and that bioturbation intensities are much more readily predicted than sediment mixed layer depths from these correlates. Further, we find that the bioturbation intensity, the depth of the sediment mixed layer, and their environmental and ecological correlates differ between shallow marine and open-ocean settings. Our findings provide new insights into the importance of potential drivers of ancient sediment mixing recorded by geologic archives. These results also highlight that climate change may, in the near future, drive shifts in bioturbation and reciprocal fundamental changes in benthic functioning.

Introduction

Marine sediments form one of the largest and most diverse habitats on Earth and have been critical in regulating global carbon and nutrient cycling both in the modern oceans^{1,2} and across geological timescales^{3,4}. The presence and ecological complexity of communities of burrowing invertebrate fauna are closely associated, in seafloor settings today, with deep and intensive sediment particle mixing which, in turn, directly influences biogeochemical cycling (including of carbon and major macronutrients) as well as both seafloor and water-column ecology^{5–8}. Similarly, the emergence and diversification of ancient bioturbators—burrowing and sediment-mixing animals—have been inferred to have shaped the evolution of biogeochemical cycling in Earth’s past oceans^{9–13}. Yet, despite mounting concern over the extent of biodiversity loss throughout the global ocean^{14,15}, there is currently limited understanding of how bioturbating animals—and the ecosystem and biogeochemical services they provide—will fare under future climate change in human-modified environments^{16–18}. Understanding of how biodiversity and environmental crises will either impact or be amplified by future changes in bioturbation in the modern ocean has been hampered, foremost, by the absence of a robust quantitative framework to reconstruct the global scope of faunal bioturbation activities. Resolving this gap, however, is necessary to determine system responses to planetary change and, at local to regional scales, is critical to setting and implementing management and conservation benchmarks related to biodiversity, climate regulation, food security and other policy-relevant arenas¹⁹.

Investigators have previously sought to identify the environmental and ecological parameters that exert control over faunal bioturbation intensity and sediment mixed layer depth by relating collated measurements of these processes to specific environmental parameters, such as water depth, freshwater discharge, dissolved oxygen levels, sedimentation rates, regional productivity

and availability of sedimentary organic matter^{20–28}. However, these analyses have tended to employ relatively small subsets of available data^{29,30}, consider a limited number of environmental parameters^{21,23}, and rely on extrapolated correlations at coarser spatial scales that exclude regional variations in physical, chemical and ecological conditions^{31,32}. Yet an extensive body of work has established that the contributions sediment-dwelling species make to ecosystem functioning are directly moderated by both biotic and abiotic local factors^{33–36}, including changes in anthropogenic behavior³⁷, which can fundamentally change the character of seafloor habitats and require the implementation of local rather than generic conservation and management strategies³⁸.

Here, we used machine learning to determine the primary environmental and ecological parameters that most strongly predict, and potentially control, variations in bioturbation processes across the modern seafloor. Additionally, we used these data to generate global maps of bioturbation intensity and sediment mixed layer depth at 5-minute spatial resolution. The bioturbation variables we consider include the biodiffusion intensity coefficient (Db , $\text{cm}^2 \text{yr}^{-1}$)—an indicator of biological mixing intensity—and the depth of the sediment mixed layer (L , cm), collected across a spectrum of contemporary marine environments and representing the most extensive bioturbation dataset currently available³⁰ (see “Methods”). We combined these data with a similarly georeferenced environmental and ecological dataset, encompassing a wide variety of parameters from surface waters, bottom waters, and the seafloor, collated from multiple sources (e.g., ref. ^{39–45}) (Table S1). Specifically, the surface-water data include temperature, primary productivity, current velocity, and ocean ecoregion division (cf. ref. ⁴⁰). The bottom-water data consist of temperature, primary productivity, current velocity, dissolved oxygen concentration, and salinity. The seafloor data include minimum distance to shore and

underwater spreading ridge systems, sedimentation rate, water depth, seafloor slope, benthic biomass, sediment total organic carbon (TOC), sediment porosity, and sediment type. These data principally represent environmental conditions but also include ecological (e.g., biomass, ecoregion division) and jointly environmental-ecological observations (e.g., productivity, TOC).

After spatially binning all target variable observations (Db and L) to 5-minute resolution (609 data points for Db and 505 for L) and correlating these target variables to georeferenced predictor variables (see “Methods”), we used a combination of ‘unsupervised’ and ‘supervised’ machine-learning techniques to explore their relationships and to make more highly resolved predictions of the distribution of Db and L values over the global ocean. For unsupervised machine learning, we used principal component analysis (PCA) to explore the relationship between each target variable (Db or L) and numerical predictor variables (environmental and ecological parameters), as well as the relationships between the numerical predictor variables. To quantitatively constrain the relationship between Db or L and these predictor variables, and to predict values of Db and L over the global ocean, we employed the Random Forest (RF) algorithm, a tree-based ensemble supervised machine-learning technique that offers low bias, moderate variance, and performs well in capturing non-linear relationships⁴⁶ (see “Methods”). We explored the relative contributions of each predictor variable to the predicted values of Db or L in a consistent and locally accurate manner by calculating SHAP (SHapley Additive exPlanations) values⁴⁷ for each parameter based on the trained RF model. We further employed our trained RF model to generate maps of the spatial variation in Db and L (as well as their standard errors) for the global ocean at a 5-minute spatial resolution, and estimated the extent to which marine protected areas (MPAs) foster the healthy operation of benthic ecological processes, including types of bioturbation that promote sediment mixing.

Results

Our comparative analyses indicate that, in contrast to common perception^{21,23}, no single dominant environmental or ecological parameter uniquely corresponds to either Db or L. In contrast, a range of parameters emerge that, either negatively or positively, correlate with these metrics of sediment mixing. Further, we find that the parameters that most strongly correlate with Db are, in many cases, not the same as those that most strongly correlate with L. This is important because previous studies have suggested that higher intensities of sediment mixing by benthic organisms promote greater mixed layer depths (e.g., ref. ^{20,22,48}), a framework that assumes that controls on Db and L are equivalent, although this is not always supported by field observations^{20–22,49}. Analysis of the Db and L observations within our database indicate that, globally, Db ranges from 0.17 (25th percentile) to 8.94 (75th percentile) cm² yr⁻¹ and L ranges from 3.8 (25th percentile) to 9.0 (75th percentile) cm (Figure 1; hereafter, all ranges represent 25th–75th percentiles, unless otherwise stated), giving an estimated global volume of actively bioturbated sediment of ~13,700–32,500 km³ for a global seafloor area of ~361 million km². However, these observations also indicate contrasting levels of variability between the shelf (here, all marine settings landward of the shelf break⁴⁰) and the open ocean: Db values are higher in shelf regions⁴⁰ (0.61–18.8 cm² yr⁻¹) relative to the open ocean (0.05–1.0 cm² yr⁻¹), whereas L values are similar across shelf (3.6–11.0 cm) and open-ocean (3.8–8.0 cm) regions. Assessment of these data using a Mann-Whitney U test further indicates that these differences in Db values between shelf and open-ocean settings are statistically significant (p-value < 2.2e-16) whereas differences in L values are not (p-value = 0.054). This distinction in spatial clustering between Db and L is also supported by Moran's I statistics (0.55 for Db and 0.38 for L), suggesting a more pronounced spatial clustering for Db compared to L.

Both Db and L display highly non-linear relationships with numerical environmental and ecological predictor variables (Figure S1), but a stronger correlation is observed between these variables and Db than between these variables and L (Figure S1). We observe that various predictor variables, including seafloor (benthic) biomass (across different trophic and size classes), primary productivity of bottom and surface waters, current velocity of bottom and surface waters, bottom-water temperature, sedimentation rate, seafloor sediment TOC, and distance to spreading ridge, show a significant positive correlation with Db (Figure S1). In contrast, water depth, bottom-water salinity, seafloor slope, seafloor sediment porosity, and distance to shore demonstrate a significant negative correlation with Db. The range of predictor variables that significantly correlate with L, however, is notably narrower (Figure S1). Significant positive correlations with L are limited to seafloor sediment TOC, seafloor macrofauna biomass, and bottom-water primary productivity; no significant negative relationships are evident.

To further explore correlations among numerical predictor variables and to facilitate the identification of a suitable parameter set for supervised machine learning, we computed the Variance Inflation Factor (VIF) for each predictor variable in the Db and L dataset (Figure S2A, B). The majority of variables exhibit relatively low VIF values (below 10), suggesting relatively low multicollinearity⁵⁰. However, both seafloor macrofauna biomass and megafauna biomass demonstrate relatively high VIF values (>10), highlighting their substantial correlation with other parameters. To mitigate potential multicollinearity, we removed the biomass parameters before running the Random Forest machine-learning model (see below). The VIF values for all remaining parameters are below 10 (Figure S2C, D), indicating relatively low multicollinearity

and enhancing the reliability of the rankings of feature importance produced by machine learning.

Spearman's correlation analyses (Figure 2) assessing the relationship between the numerical predictor variables and each target variable reveal a more pronounced correlation between Db and the environmental and ecological parameters we consider (the ten largest absolute correlation coefficients are greater than 0.3) than emerges for L (all absolute correlation coefficients are less than 0.2). In agreement with some previous findings²⁰, the environmental and ecological parameters that have the highest degree of correlation with Db are distinct from those that have the highest degree of correlation with L (Figure 2). Of the ten most strongly correlated environmental and ecological predictor variables, seafloor (benthic) biomass (across various trophic and size classes), bottom- and surface-water primary productivity, bottom-water current velocity, and seafloor sediment TOC exhibit a strong positive correlation with Db, whereas seafloor macrofauna biomass, bottom-water dissolved oxygen, bottom-water primary productivity, bottom-water current velocity, seafloor megafauna biomass, and sedimentation rate are positively (but weakly) correlated with L (Figure 2). Similarly, water depth and distance to shore negatively correlate with Db, whereas seafloor sediment TOC, seafloor slope, and surface-water current velocity, albeit weakly, negatively correlate with L (Figure 2). A negative correlation between seafloor sediment TOC and L has been previously reported²⁶ but, notably, we find that the correlation coefficients for L are considerably weaker than those for Db. We also find variable degrees of interdependency between a range of environmental and ecological parameters (Figure S3).

Our PCA analysis (Figure 3) of numerical predictor variables reveals that seafloor biomass, bottom- and surface-water primary productivity, sedimentation rate, sediment TOC, and bottom-water temperature, each characterized by relatively higher PC1 values, can be grouped together and are associated with higher Db values (Figure 3A). Conversely, water depth, sediment porosity, distance to shore, and surface-water temperature, characterized by lower PC1 values, can be grouped together and are associated with lower Db values (Figure 3A). A similar grouping of predictor variables is observed for the dataset linked to L. However, similar L values occur in both the negative and positive PC1 fields of our predictor variables (Figure 3B), reflecting weak relationships between L and these environmental and ecological parameters. This disparity underscores the potentially strong variability in the extent to which environmental and ecological parameters shape Db and L. Beyond the scope of PCA, we also observe that Db is characterized by greater disparity, relative to L, between distinct sediment types, with higher Db values observed in areas of the seafloor dominated by siliciclastic sands, silts and clays and lower Db values observed in sediments dominated by biogenic calcareous oozes (Figure S4).

To predict the finer-scale distribution of variability in Db and L (beyond the resolution afforded by previously available empirical datasets), we built a machine-learning model by selecting an initial set of fifteen mainly environmental parameters as predictor variables (Table S1) (based on knowledge accrued during our correlation and unsupervised machine-learning analyses) and using Db and L as target variables; we consider this our baseline scenario. We employed a 10-fold cross-validation method⁵¹ to evaluate the predictive capability of our machine-learning model on unseen data. We selected the Random Forest (RF) model for its robust handling of high-dimensional data with complex non-linear relationships and its better predictive accuracy for our dataset, as evidenced by its higher R^2 values compared to other machine-learning

algorithms (Table S2). Our RF model represents a significant advance in global-scale predictive capability for Db (the 10-fold cross-validation R^2 value is 0.54; Figure 4A) relative to previous prediction efforts (e.g., R^2 values of 0.22 in ref. ²¹ and 0.43 in ref. ²³). For L, we obtained a 10-fold cross-validation R^2 value of 0.29 (Figure 4B). Previous efforts to predict L vary widely, with R^2 values ranging from 0.001 in ref. ²¹ to 0.47 in ref. ²⁶ or 0.88 in ref. ²⁵, but these analyses did not have the same observation density, spatial coverage or spatial resolution as our dataset. Larger R^2 values for the prediction of L emergent from previous studies^{25,26} suggest that smaller datasets do not adequately capture the global-scale heterogeneity of values and environmental correlates that emerges from our larger and more robustly validated (e.g., using 10-fold cross-validation) dataset. The relative importance of each variable, as determined by SHAP value (Figure 4C, D), indicates that the top five parameters that most strongly correlate with Db are water depth (negative correlation; Figure S5A–D), surface-water primary productivity (positive correlation), bottom-water primary productivity (positive correlation), sediment type (categorical variable, so no numerical correlation), and distance to shore (negative correlation). The top five parameters that most strongly correlate with L are sediment TOC (negative correlation; Figure S5E–I), surface-water primary productivity (positive correlation), water depth (negative correlation in shallower water and positive correlation in deeper water), bottom-water salinity (negative correlation), and distance to shore (negative correlation). It is notable that the mean |SHAP| values for all parameters, when predicting L, are substantially lower compared to those associated with prediction of Db, and thus lower confidence should be attributed to predictions of L from correlative environmental and ecological parameters (Figure 4C, D).

To examine the potential for enhancing model accuracy with more predictor variables, we incorporated additional ecological parameters into the RF model, including benthic biomass and

biogeographical ecoregion division. As RF predictions of Db and L using this more comprehensive set of global environmental and ecological parameters did not increase model fit (R^2 values: Db, 0.51; L, 0.26) relative to RF results using our smaller subset of variables, and given the risk of introducing more multicollinearity into the parameter space, we omitted these additional ecological parameters from subsequent machine-learning analyses. To examine the influence of sampling bias—stemming from the uneven distribution of Db and L observations across the seafloor (Figure 1; ref. ³⁰)—on model accuracy and interpretation, we employed a randomized sub-sampling technique. Specifically, we randomly sampled 50% of the entire Db or L dataset, with the selection probability scaled by the inverse density of samples within each $10\times 10^\circ$ grid, across the entire global seafloor (Figure S6A, C). This method markedly diminished sample clustering in particular regions, serving as a robust sensitivity analysis for evaluating model performance. Applying the RF model to these Db and L subsets and the corresponding subset of environmental and ecological parameters did not substantively change model outcomes (10-fold cross-validation R^2 : Db, 0.55; L, 0.22) or the rank importance of predictive parameters (Db subset: water depth, followed by surface-water primary productivity, bottom-water primary productivity, and sediment type; L subset: sediment TOC followed by surface-water primary productivity) (Figure S6B, D). To investigate the potential for methodological differences in collection of bioturbation data to shape model outcomes—particularly given the disparate time scales of integration represented by different bioturbation proxies (e.g., common use of different radioisotope systems as well as particulate tracers and textural proxies, cf. ref. ³⁰), we subsampled all Db and L data originally derived from ^{210}Pb observations, which represent a substantial share of the whole dataset (62% for Db and 62% for L). We observe that even when constrained to a single (and shared) methodology, the disparity between Db and L persists.

Specifically, when using only the ^{210}Pb data, machine learning still performs better for Db ($R^2 = 0.57$) than for L ($R^2 = 0.3$), with R^2 values similar to those of our baseline scenario ($R^2 = 0.54$ for Db and 0.29 for L). Additionally, we find congruence in four of the top five most important features for Db, and four of the top five most important features for L, between the ^{210}Pb -based analyses and our baseline scenario (Figure S7; Figure 4C, D).

Next, to explore whether the environmental and ecological parameters vary in their relative importance in predicting variations in Db and L between different environmental settings, we independently trained RF models to predict Db and L in shelf and open-ocean regions. We find that the predictability either diminished or remained comparable relative to model performance using the full global ocean dataset (R^2 : shelf Db = 0.40, L = 0.28; open-ocean Db = 0.44, L = 0.33). The ranking of environmental and ecological parameter importance for shelf Db aligns with findings emergent from the full dataset, with water depth, sediment type, bottom-water primary productivity, and surface-water primary productivity retaining their status as the parameters with the greatest predictive power (Figure 5A). Conversely, for open-ocean Db, surface-water primary productivity, distance to spreading ridge, surface-water current velocity, and sediment TOC emerge as the strongest environmental and ecological predictors, whereas water depth, distance to shore, and bottom-water productivity emerge as relatively unimportant (Figure 5C). For L, the environmental and ecological parameters with the highest predictive power in shelf and open-ocean settings are similar to those identified for the full dataset, with surface-water primary productivity, distance to shore, sediment TOC, and water depth emerging as the strongest environmental and ecological correlates in shelf settings (Figure 5B), and surface-water primary productivity, sediment TOC, bottom-water dissolved oxygen, and water depth holding the greatest predictive power in open-ocean settings (Figure 5D).

After applying the RF model (trained using the full dataset) to our selected suite of 15 environmental and ecological parameters across the global ocean, we generated high-resolution maps of predicted Db and L (and associated standard error) across the global seafloor (Figure 6). Because Random Forest learns and makes predictions based on the patterns and relationships found within the sample dataset, its performance can be limited when predicting conditions that significantly deviate from those encountered during training. The value range of the training predictor variables encompasses approximately 97–99% of the values of predictor variables across the global ocean, indicating that the trained RF model can derive robust global patterns with minimal extrapolation. These maps show that predicted Db values range from 0.12 to 0.50 $\text{cm}^2 \text{yr}^{-1}$ and that L ranges from 5.6 to 8.7 cm (25th and 75th percentiles, respectively). Mirroring the broad-scale patterns that emerged from the empirical dataset, predicted Db values are higher in shelf settings (0.69–3.78 $\text{cm}^2 \text{yr}^{-1}$) relative to open-ocean regions (0.11–0.30 $\text{cm}^2 \text{yr}^{-1}$), whereas L values (shelf, 5.0–7.1 cm; open-ocean, 5.8–9.0 cm) do not show this dichotomy.

Utilizing our predicted global maps of Db and L, we assessed whether the implementation of marine protected areas (MPAs)⁵², in general, promote higher levels of bioturbation and thus the healthy functioning of benthic ecological processes. By comparing values of Db and L within designated MPAs (“within MPA”, Figure 7) with those determined for an equivalently sized and immediately adjacent non-protected area (“surrounding MPA”, Figure 7), we find that values of Db and L within (Db: 3.1–9.7 $\text{cm}^2 \text{yr}^{-1}$; L: 5.5–7.6 cm) and surrounding MPAs (Db: 2.5–9.4 $\text{cm}^2 \text{yr}^{-1}$; L: 5.5–7.5 cm) are broadly comparable (Figure 7B, C). The relative percentage change for either Db or L between MPAs and surrounding areas is limited (Db, median change is 0.54% and -2.9%–6.3% for 25th to 75th percentile; L, median change is 0.11% and -1.2%–2.2% for 25th to 75th percentile; Figure 7B, C). Wilcoxon Signed-Rank analyses reject the null hypothesis of no

relative change, for both Db and L, between MPAs and surrounding areas ($p = 1.5e-12$ for Db and $8e-9$ for L), indicating a detectable albeit limited disparity between MPAs and surrounding unprotected areas; median percent change values are nonetheless close to 0%. We also find that the relative change (either positive or negative) between MPA and non-MPA pairs does not appear to vary systematically with geographic location. To quantify the impact of machine-learning uncertainty on these estimates of percentage change, we employed Monte Carlo resampling (1000 iterations) to randomly sample both Db and L values for each pixel, considering the standard error, and calculate the relative percentage change between MPAs and unprotected surrounding areas. These analyses yield percentage changes ranging from -5.1% to 9.1% for Db (25th and 75th percentile, with a median of 1.1%) and from -2.8% to 4.1% for L (with a median of 0.49%), further underscoring that most protection measures do not appear to be sufficiently targeted to substantially impact bioturbation, as predicted by its environmental and ecological correlates.

Discussion

Our unsupervised and supervised machine-learning models applied to the full global inventory of Db and L highlight the variability inherent in each of these bioturbation metrics (and the processes they represent). Therefore, when inferring underlying patterns in distribution—whether local, regional or global—it is critical to consider the broadest possible representation of bioturbation. Our analyses also strongly suggest that a wide range of environmental and ecological factors⁵³—and not individual factors such as sediment accumulation rate^{21,54,55} or water depth²³ alone—are strongly correlated with and may play a significant role in shaping bioturbation intensity and sediment mixed layer depth on a global scale. The positive multivariate relationships we have identified may indicate that bioturbation intensities and depths

are linked strongly to parameters relating directly to food supply and availability, productivity and benthic biodiversity, whereas the negative multivariate relationships may potentially reflect either changes in body size and other species traits with water depth^{44,56} or reductions in surface-ocean or benthic productivity or export productivity associated with changes in nutrient or light availability^{30,57} that coincide with increasing water depth or distance from land. However, the interdependencies between these variables (Figure S3) may hamper precise mechanistic evaluation, and it is important to emphasize that, within our identified subsets of correlates, the relative importance of specific abiotic and biotic variables are likely to vary with location, timing, environmental setting and ecology³⁵.

Given that several of the environmental and ecological parameters that emerge from our analyses as particularly strong correlates (and potential drivers) of bioturbation intensity have been proposed to have varied considerably across the Phanerozoic—most notably productivity and benthic biomass^{58–60}—this suggests that Phanerozoic increases in productivity and benthic metabolic expenditure may have contributed to observed secular increases in bioturbation intensity, as previous studies have suggested (e.g., ref. ^{58,61–64}). Importantly, however, evolutionary changes in bioturbation intensity may not have translated into directional changes in sediment mixed layer depth. Although not environmentally invariant, as has previously been suggested²⁰, from our analyses L values do not appear to group strongly on the basis of either water depth or sedimentation rate, nor to be as readily predicted as Db by any correlative environmental and ecological parameters, whether alone or in combination (Figure 2B, 3B, 4) and regardless of proxy system employed (Figure S7). Although both Db and L are linked to the bioturbation activities of burrowing infauna^{20,22,48}, this disparity strongly suggests that environmental and ecological processes shaping intensities and depths of faunal-mediated

sediment mixing are likely decoupled. These findings emphasize the need for greater mechanistic understanding of the processes that influence the development and depth of the sediment mixed layer^{25,53} and how these differ from processes that shape biodiffusion intensity. Previous studies have also noted disparities between environmental patterns of Db and L and have suggested that L may be limited by the high metabolic costs of burrowing deeper as sediments become increasingly compacted, and by the declining availability of labile organic matter^{20,21}. However, we do not observe a strong correlation between L and either sediment type, porosity, or sedimentation rate (Figure 3, S3, S4), and recent work has highlighted that burrowing invertebrates adjust the mechanical efficiency of their burrowing behavior, allowing burrow networks and vertical mixing to extend to depths beyond expectations based on the physical properties of the sediment profile⁶⁵. Similarly, bottom-water primary productivity and sediment TOC are strongly correlated with Db, but are notably weaker correlates of L. We conclude, therefore, that although the rheology and organic content of seafloor sediments may indeed influence styles and intensities of burrowing (and may have additionally shaped the Phanerozoic evolution of bioturbation depths and intensities, as previous studies have suggested; e.g., ref. ^{61–64,66}), they do not appear to sufficiently account for the variable distribution of mixed layer depths in the modern oceans.

The dichotomy we observe between patterns of correlation for Db and L may potentially reflect the inability of our compiled global environmental and ecological dataset to capture transient or rapid community responses to environmental variation. Both bioturbation and many of the environmental and ecological parameters represented in our dataset vary dramatically on seasonal or even shorter time scales^{48,67–70}, as well as along environmental gradients⁷¹. The inability of our model to more robustly predict L suggests that mixed layer depth may be

comparatively more varied, on relatively fine spatial or temporal scales, than mixing intensity. If these observations hold true, this implies that Db may more closely track the behavioral and functional diversity of the community (which, in turn, may be regulated by longer-term environmental conditions), whereas L may be more sensitive to the impact of seasonally variable local environmental conditions on the abundance and biomass of bioturbating fauna. We suggest that coupled collection of bioturbation and environmental data at a higher temporal resolution should be a priority for future studies. We cannot rule out, however, that variability in Db and/or L may reflect other factors not yet determined at a global scale¹⁹, in particular ecological responses to change (e.g., intraspecific variability, dominance, composition, and compensation)^{36,72–74}. This underscores the potential vulnerability of seafloor level-bottom communities to short-term climate and environmental perturbations; it is anticipated that accelerating rates of not only coastal but also open-ocean and deep-water warming⁷⁵ will affect species' behavior^{35,76}, and reductions in seafloor biomass associated with climate change^{16,17} will minimize benthic community contributions. Further, the non-linear relationships we observe between our environmental and ecological predictor variables and Db and L (Figure S1) suggest a heightened risk that abrupt fundamental shifts in processes critical to benthic functioning, such as bioturbation, will occur as environmental thresholds are crossed under climate forcing^{77,78}, as has already begun in regions of climate amplification⁷¹.

The diminished significance of water depth and bottom-water primary productivity in predicting open-ocean Db, as opposed to shelf Db (Figure 5), may reflect the inherent heterogeneity (and more productive nature) of shallow-water seafloor systems. In shelf settings, water depth (and associated hydrodynamic energy) varies considerably relative to seafloor area and appears to be associated with substantial variability in Db. In contrast, in open-ocean settings, the association

between water depth and Db is comparatively weak. Bioturbation intensities in open-ocean settings may, therefore, be comparatively less responsive, respond on longer timescales, or be ‘buffered’ against the effects of environmental variability.

Despite our model's global scale, its high resolution (5-minute pixels) enables the identification of smaller-scale patterns. Our findings on the relationship between MPAs and bioturbation suggest that management of benthic resources to promote seafloor ecosystem health may be insufficiently targeted across a variety of geographic and environmental settings, presumably because protection measures are commonly not tailored to seafloor biodiversity and functioning and do not comprehensively protect the seafloor from all salient critical pressures^{79–82}. If we are to mitigate the effects of anthropogenic activity and climate change on seafloor functioning, better understanding will likely be essential to more robustly evaluate when and how the contributions of bioturbating species are compromised⁸³, how altered biodiversity⁸ and post-perturbation compensatory mechanisms alter biodiversity-ecosystem function relations^{17,73}, and how the ecology of benthic environments can deliver transformative change to enhance and deliver innovative adaptation plans^{84,85}.

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Author contributions

S.Z., L.T., and M.S. designed research; S.Z., M.S., and L.T. provided methodology; S.Z. performed research, analyzed data, and contributed new modeling tools; S.Z., L.T., and M.S. wrote the paper.

Declaration of interests

The authors declare no competing interests.

Figures Legends

Figure 1. Spatial distribution of bioturbation intensity (Db) and sediment mixed layer depth (L) values over the global ocean. (A) The spatial distribution of $\log_{10}(\text{Db})$ ($n = 609$). (B) The spatial distribution of $\log_{10}(\text{L})$ ($n = 505$). In each display panel, points represent spatially binned information at a spatial resolution of 5×5 minutes. The insets display histogram distributions of Db and L, with data for each bioturbation metric split into 30 equal bins.

Figure 2. Spearman's rank correlation coefficient between bioturbation and the environmental and ecological parameters. (A) Bioturbation intensity (Db). (B) Sediment mixed layer depth (L). Red bars indicate a positive correlation, whereas blue bars signify a negative correlation. The absolute value of the correlation coefficient decreases from top to bottom of each diagram. See also Figure S1, S2 and S3.

Figure 3. Projection of the environmental and ecological parameters on the first and second principal components. Observations are shown as points in the plane formed by two principal

components (PC1 and PC2) and are colored according to values of (A) bioturbation intensity (Db) or (B) sediment mixed layer depth (L). The arrows represent the PCA loadings of the environmental and ecological parameters. See also Figure S4.

Figure 4. Machine-learning (Random Forest model) outcomes for bioturbation intensity (Db) and sediment mixed layer depth (L). The correspondence between predicted and

observed values of (A) Db and (B) L values are shown for the 10-fold hold-out test dataset (data unseen during the training stage). The red line represents a hypothetical 1:1 correspondence. (C, D) The mean absolute SHAP values (displayed in decreasing order, from top to bottom) associated with calculation of target variables (Db and L) from predictor variables (i.e., environmental and ecological parameters). See also Figure S5, S6, and S7.

Figure 5. Machine-learning outcomes for Db and L for samples located in the shelf region and in the open-ocean region. (A, B) The mean absolute SHAP associated with calculation of target variables Db and L from predictor variables in the shelf region. (C, D) The mean absolute SHAP associated with calculation of target variables Db and L from predictor variables in the open-ocean region.

Figure 6. Random Forest-generated predictions of the global distribution of modern marine bioturbation intensity (Db) and sediment mixed layer depth (L). (A, C) Predicted value at 5-minute resolution by the Random Forest model from geospatially correlative environmental and ecological variables. (B, D) Standard error of the predicted value at 5-minute resolution. Marine white regions represent areas lacking predictor variable data.

Figure 7. Comparison of predicted Db and L values between marine protected areas (MPAs) and equivalent adjacent non-protected areas. (A) Locations of each of the 5056 marine protected areas (red polygons) alongside the area-equivalent adjacent non-protected areas (blue polygons). (B, C) distribution of the predicted mean bioturbation intensity (Db) and predicted mean sediment mixed layer depth (L) for each marine protected area and the equivalent adjacent non-protected area, for the subset of MPAs exceeding 2 pixels in size. Insets indicate the distribution of percentage change of predicted mean bioturbation intensity (Db) or predicted mean sediment mixed layer depth (L) values between each marine protected area and the paired equivalent and adjacent non-protected area. Note that extreme percentage change values (the uppermost 0.5% and lowermost 0.5%) are omitted from the insets of panels B and C for clarity.

STAR Methods

RESOURCE AVAILABILITY

Lead contact

Further information and requests for resources and reagents should be directed to and will be fulfilled by the lead contact, Shuang Zhang (shuang-zhang@tamu.edu).

Materials availability

No unique reagents were generated in this study.

Data and code availability

All raw and spatially binned Db and L data needed to evaluate the conclusions in the paper are present in the Supplemental Information. The raw Db and L data are also available at <https://doi.org/10.7910/DVN/GBELFW>. The code used to analyze the data, run the machine-learning model, and produce the global Db and L maps, as well as additional supporting information, have been posted to the Zenodo platform (<https://zenodo.org/records/11069160>).

METHOD DETAILS

Establishing a dataset of global patterns in bioturbation

We utilized a recently published fully georeferenced compilation of marine bioturbation data which includes reported values of the biodiffusion intensity coefficient (Db, $\text{cm}^2 \text{yr}^{-1}$) and the depth of the sediment mixed layer (L, cm) as published in peer-reviewed literature³⁰. This bioturbation dataset was meticulously compiled to exclude areas disturbed by human activities, particularly those subjected to extensive trawling. As this dataset encompasses a wide range of methods for measuring Db and L that demonstrate methodological dependency^{27,85–89}, we constrained our analysis to data obtained using downcore profiles of radioisotopes, chlorophyll *a*, and particulate tracers, such as luminophores and tek beads (for both Db and L); glass beads (Db only); organic carbon, luminescence and x-ray profiles, as well as values calculated from modeling of geochemical profile data (L only). These data were further processed by averaging intra-locality data (georeferenced to the same latitude and longitude) through time for individual localities. As both the Db and L data are right-skewed, we applied a \log_{10} transformation to mitigate the skewness, stabilize the variance, and reduce any impact of outliers (Figure 1). To reduce the impact of outliers on our analyses we then removed the lowermost and uppermost 1% of observations from each of these two bioturbation datasets. To match point measurements of

Db and L with associated environmental and ecological parameters, we binned our Db and L data to a spatial resolution of 5×5 arc minutes. Specifically, multiple data points within 5 arc minutes were averaged. The final dataset contains 609 values for Db and 505 values for L.

Linking Db and L to spatially correlative environmental and ecological data

We assessed the relationship between Db or L values and a range of marine environmental and ecological parameters from the same georeferenced areas. These parameters (for data resolution and sources, see Table S1) include surface ocean features (including ocean ecoregion division, surface-water temperature, surface-water primary productivity, and surface current velocity); bottom-water features (including bottom-water temperature, bottom-water primary productivity, bottom-water current velocity, bottom-water dissolved oxygen concentration, and bottom-water salinity); and seafloor features (including minimum distances to shore and underwater spreading ridge systems, sedimentation rate, water depth, seafloor slope, benthic biomass, sedimentary total organic carbon (TOC), sediment porosity, and sediment type).

To account for differences in environmental conditions, we determined 15 divisions (in a shapefile vector format) from Bailey's ecoregions of the oceans⁴⁰: Shelf, Equatorial Countercurrent Division, Equatorial Trades Division, Trade Winds Division, Poleward Trades Division, High-Salinity Tropical Monsoon Division, Tropical Monsoon Division, Poleward Monsoon Division, Jet Stream Division, High-Salinity Subtropical Division, Subtropical Division, Equatorial Westerlies Division, Poleward Westerlies Division, Outer Polar Division, and Inner Polar Division. For seafloor sediment type, data were converted to a shapefile vector format (from the original raster format) and are comprised of 13 sediment types⁴¹, including

gravel and coarser, sand, silt, clay, ash and volcanic sand/gravel, fine-grained calcareous sediment, siliceous mud, calcareous ooze, radiolarian ooze, diatom ooze, sponge spicules, mixed calcareous-siliceous ooze, and shells and coral fragments. Ecoregion division and sediment type are both categorical variables. To associate Db and L values with categorical values for each ecoregion division and seafloor sediment type, we calculated the distance between each Db or L pixel and different divisions and sediment types and then determined the nearest division and sediment type and assigned those to each Db or L pixel. The geospatial analyses described above were accomplished using GRASS GIS⁹⁰ and R⁹¹.

Principal component analysis (PCA)

We conducted the PCA analysis on the numerical predictor variables in R using the PCA() function from the FactoMineR package⁹². The arrows in Figure 3 represent the PCA loadings of the environmental and ecological parameters. The length of each arrow indicates the strength of correlation between each environmental and ecological parameter and the principal components: a longer arrow signifies a stronger correlation with the principal components, whereas a shorter arrow signifies a weaker correlation. Each arrow's direction reflects the relationship between that parameter and the axes, indicating that the parameter positively or negatively influences the principal components. Accordingly, parameters represented by arrows pointing in similar directions are positively correlated with each other, whereas those pointing in opposite directions are negatively correlated with each other. Parameters represented by arrows that are perpendicular to one another are not correlated.

Supervised machine learning

To quantitatively predict and map Db and L over the global ocean, we employed the Random Forest (RF) algorithm⁴⁶. RF is based on a decision tree, which is a non-parametric supervised learning algorithm used for classification or regression and that captures non-linear relationships in the data and is robust to input outliers⁹³. However, decision trees can also be prone to overfitting (i.e., high variance), be relatively low in predictive accuracy and may not yield optimal solutions. RF can be thought of as an ensemble of many trees (the ‘bagging’ technique), thereby reducing the high variance experienced by a single decision tree. Unlike simple bagging, RF further decorrelates the trees, using a random set of predictor variables to divide the training data during each split. All these modifications contribute to the high predictive accuracy (low bias and moderate variance) of RF. The RF algorithm has been widely used in geoscience research^{94,95} and is particularly suited, as here, to making predictions using high-dimensional data with complex non-linear relationships (Figure S1). The construction of the RF model was conducted in R⁹¹ using the “range” package⁹⁶. Optimal values of hyperparameters (i.e., number of trees and minimum node size) in the RF model were generated by implementing a 10-fold cross-validation coupled with a grid search in the training process. Specifically, we employed a random partitioning strategy to divide the entire dataset into 10 subsets. We set aside one of these subsets for evaluation purposes (i.e., the hold-out dataset), while we used the remaining nine subsets to train the model. We assessed model performance across a range of number of trees (from 500 to 1500) and conducted sensitivity analyses of minimum node size, ranging from 1 to 10. Mean-squared-error (MSE) was used as an indicator of performance. The tuned parameters that yield the smallest average MSE of the hold-out test data (i.e., test set in the 10-fold cross-validation) were selected as the best parameters. We then used the best set of hyperparameters to train our RF model on the entire dataset to generate the final model.

Compared with linear models, the RF model excels at capturing non-linear relationships between variables and identifying complex patterns that linear models may overlook. We performed the same task with a range of different machine-learning algorithms to independently verify the suitability of applying RF to our dataset; the lower R^2 values for Db and L emergent from other algorithms (Table S2) confirms the appropriateness of our selection of the RF algorithm. Moreover, the consistent disparity in predictive power between Db and L across various algorithms supports the robustness of our interpretations regarding the challenges associated with using environmental and ecological parameters to predict L relative to Db (see Table S2 for details about these algorithms). The uncertainty (standard error) of the RF model was quantified using the Infinitesimal Jackknife method⁹⁷, which is particularly useful for complex machine-learning models (such as the RF model) where traditional resampling-based methods may be computationally prohibitive. When applied to the RF model, the Infinitesimal Jackknife works by calculating the infinitesimal perturbations of each individual tree with respect to the data on which it was trained. These perturbations are then aggregated to provide an overall estimate of the standard error of the model's predictions (see Figure 6 for the standard error when mapping Db and L over the global seafloor).

To illustrate the relative contributions of each predictor variable to the predicted Db or L values, we calculated the SHAP value for each variable based on the final model. SHAP (“SHapley Additive exPlanations”) values represent the importance of each individual variable for tree ensembles in a consistent and locally accurate manner⁴⁷. The fundamental principle of SHAP is based on random permutation. However, unlike the simple permutation method (i.e., permuting one column and comparing the prediction generated from this permuted dataset with the original

prediction), the SHAP method calculates the contributions of each parameter by averaging the differences in predictions across every possible parameter ordering. This approach considers interactions between parameters and compares the contributions of parameters fairly. In particular, the interpretation of the SHAP value for parameter x in one sample represents the contribution of parameter x to the prediction of this specific sample compared to the average prediction for the entire dataset. A positive SHAP value for a specific x indicates that the current x value contributes positively to the difference between the prediction of the specific focal sample and the average of the entire dataset. The correlation between the SHAP value of x and the x value itself signifies the relationship between the x variable and the target variable. A positive trend indicates a positive correlation, while a negative trend indicates a negative correlation. The mean $|\text{SHAP}|$ value of parameter x for all samples represents the overall absolute contribution of x to the prediction, which is treated as the feature importance for x (Figure 4).

To further test the impact of multicollinearity among predictor variables on the model's ability to rank feature importance, we implemented a scenario with an even smaller set of predictor variables than our baseline. In this refined scenario, we removed bottom-water primary productivity and distance to shore, each of which exhibits high correlation with water depth, from the parameter space. Negligible improvement in VIF metrics was observed for this smaller dataset compared with the baseline scenario. The most important predictors for Db and L that emerged from this analysis align closely with our baseline results, reinforcing the validity of the feature importance ranking of our baseline scenario.

Effectiveness of marine protected areas

We compiled marine protected area (MPA) information from the shapefile in the World Database on Protected Areas⁵². MPAs were selected by choosing the rows that had “designated” values for the “STATUS” field and “100% marine PA” for the “MARINE” field in the shapefile. We also unified MPA rows that had the same MPA name, resulting in 5056 MPA multi-polygons. To generate an area-equivalent adjacent non-protected region for each MPA, we first calculated the centroid for each MPA, then gradually enlarged a buffer polygon surrounding the centroid until the difference in area between the buffer and the MPA polygon was equivalent to the area of the MPA. After subtracting each MPA polygon from its buffer polygon, we obtained the area-equivalent adjacent polygon (Figure 7A). The area ratio of the non-protected region to the MPA region ranges from 0.9 to 1.1, supporting a direct comparison of the Db and L values between each pair of MPA regions and the non-MPA regions.

We overlaid each MPA region and its adjacent non-protected region on the predicted global Db and L maps and extracted the mean Db and L values. Given the pixel resolution of the predicted global Db and L values at 5 minutes by 5 minutes, certain small MPA regions and non-MPA regions may fall within the same pixel. This situation can bias comparisons, as the granularity of a single pixel is insufficient to discern meaningful differences between MPAs and non-MPA regions. To mitigate this potential bias, we applied a filter to include only MPA regions comprising at least three pixels, yielding a refined dataset of 1,534 MPA regions for further analysis. The distribution of Db and L values show similar magnitudes between the screened MPA regions and their adjacent regions (Figure 7B, C). To further compare the Db and L values in MPA regions versus adjacent non-protected regions, we calculated the Log Response Ratio (LRR):

$$LRR = \log \left(\frac{\text{Mean Db or L in MPA}}{\text{Mean Db or L in adjacent region}} \right)$$

In a final step, we calculated the relative percentage change of Db or L ($\% \text{ Change} = 100\% \times (e^{LRR} - 1)$); Figure 7B, C).

QUANTIFICATION AND STATISTICAL ANALYSIS

We conducted the Mann-Whitney U test for the Db and L values between shelf and open-ocean settings. We calculated Moran's I statistics for the Db and L values. We future conducted Wilcoxon Signed-Rank test for the relative change of Db and L values from areas immediately surrounding each MPA to the MPA.

Supplementary Dataset files

Data S1: Raw Db and L data used for analysis in this study.

Data S2: Geographically binned Db data (5-min resolution).

Data S3: Geographically binned L data (5-min resolution).

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TABLE FOR AUTHOR TO COMPLETE

Please do not add custom subheadings. If you wish to make an entry that does not fall into one of the subheadings below, please contact your handling editor or add it under the “other” subheading. **Any subheadings not relevant to your study can be skipped.** (NOTE: references should be in numbered style, e.g., Smith et al.¹)

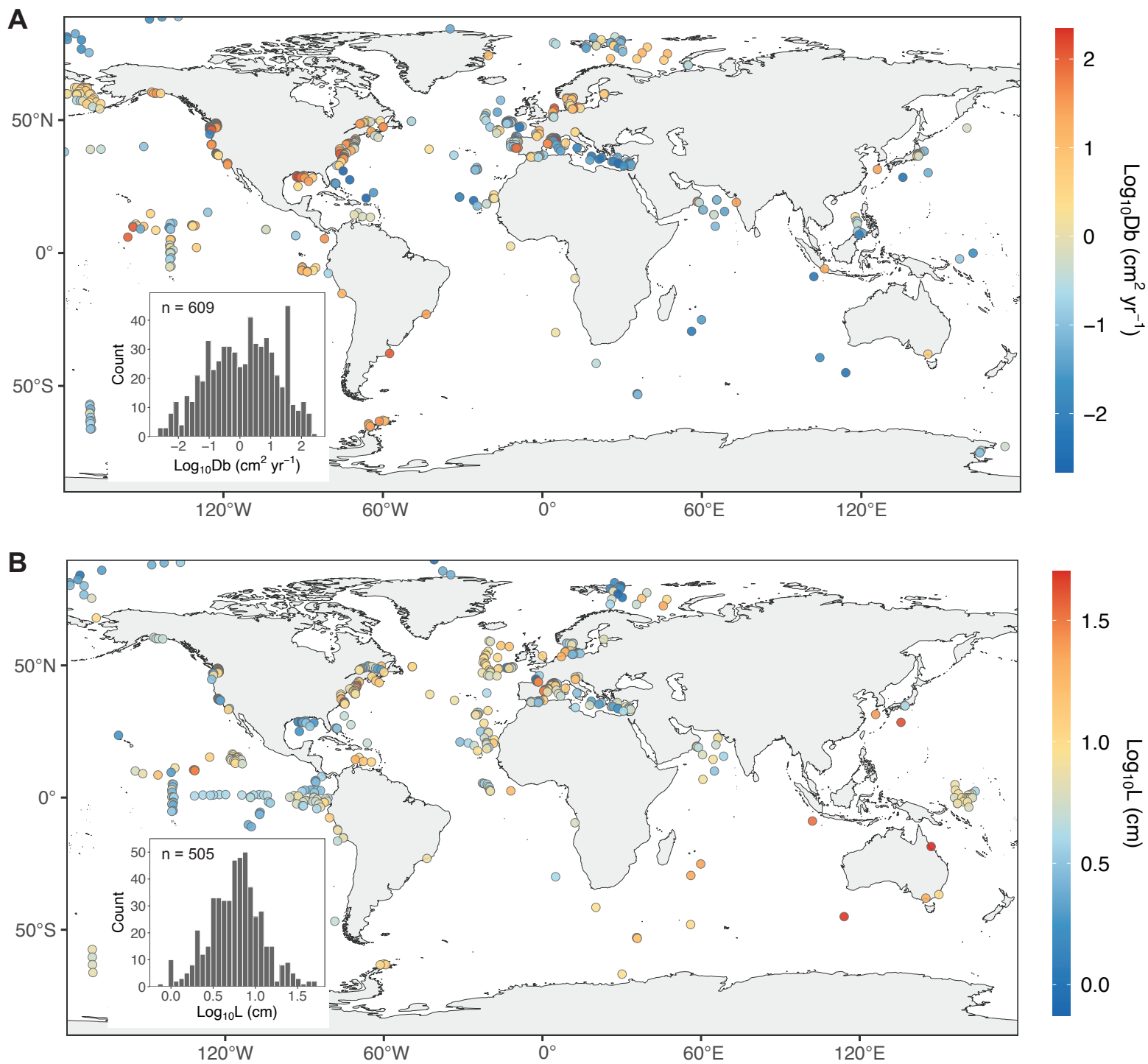
Key resources table

REAGENT or RESOURCE	SOURCE	IDENTIFIER
Antibodies		
Bacterial and virus strains		
Biological samples		
Chemicals, peptides, and recombinant proteins		
Critical commercial assays		
Deposited data		
Raw Db and L data	Solan et al. ³⁰	https://dataverse.harvard.edu/dataset.xhtml?persistentId=doi:10.7910/DVN/GBELFW
Spatially binned Db and L data	This paper	N/A
Mean surface-water temperature	Assis et al. ³⁹	https://www.bio-oracle.org/downloads-to-email-v2.php?version=2_1

Mean surface current velocity	Assis et al. ³⁹	https://www.bio-oracle.org/downloads-to-email-v2.php?version=2_1
Mean bottom-water temperature	Assis et al. ³⁹	https://www.bio-oracle.org/downloads-to-email-v2.php?version=2_1
Mean bottom current velocity	Assis et al. ³⁹	https://www.bio-oracle.org/downloads-to-email-v2.php?version=2_1
Mean bottom-water salinity	Assis et al. ³⁹	https://www.bio-oracle.org/downloads-to-email-v2.php?version=2_1
Mean bottom-water dissolved oxygen concentration	Assis et al. ³⁹	https://www.bio-oracle.org/downloads-to-email-v2.php?version=2_1
Mean surface-water primary productivity	Assis et al. ³⁹	https://www.bio-oracle.org/downloads-to-email-v2.php?version=2_1
Mean bottom-water primary productivity	Assis et al. ³⁹	https://www.bio-oracle.org/downloads-to-email-v2.php?version=2_1
Water depth	Assis et al. ³⁹	https://www.bio-oracle.org/downloads-to-email-v2.php?version=2_1
Seafloor slope	Assis et al. ³⁹	https://www.bio-oracle.org/downloads-to-email-v2.php?version=2_1
Minimum distance to shoreline	Assis et al. ³⁹	https://www.bio-oracle.org/downloads-to-email-v2.php?version=2_1
Ocean ecoregion	Bailey ⁴⁰	https://link.springer.com/book/10.1007/978-1-4939-0524-9
Seafloor sediment type	Dutkiewicz et al. ⁴¹	http://portal.gplates.org/portal/seafloor/
Seafloor sediment TOC content	Lee et al. ⁴²	https://zenodo.org/records/1471639
Seafloor sediment porosity	Martin et al. ⁴³	https://agupubs.onlinelibrary.wiley.com/doi/full/10.1002/2015GL065279
Ocean benthic biomass	Wei et al. ⁴⁴	https://journals.plos.org/plosone/article?id=10.1371/journal.pone.0015323

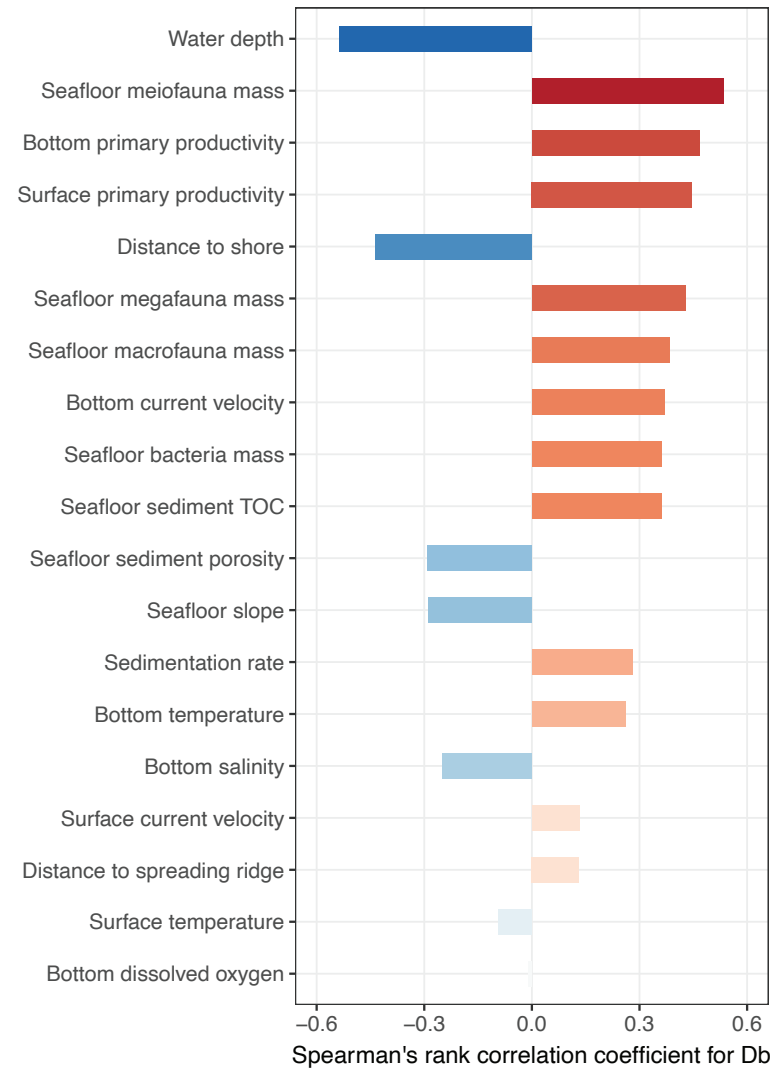
Sedimentation rate	Halevy et al. ⁴⁵	https://www.science.org/doi/10.1126/science.adh1215
Experimental models: Cell lines		
Experimental models: Organisms/strains		
Oligonucleotides		
Recombinant DNA		
Software and algorithms		
GRASS GIS v7.8	GRASS Development Team ⁹⁰	https://grass.osgeo.org
R v4.3.3	R Core Team ⁹¹	https://www.r-project.org/
FactoMineR v2.8	Lê et al. ⁹²	https://cran.r-project.org/web/packages/FactoMineR
Ranger v0.15.1	Wright and Ziegler ⁹⁶	https://cran.r-project.org/web/packages/ranger
Other		

Figure

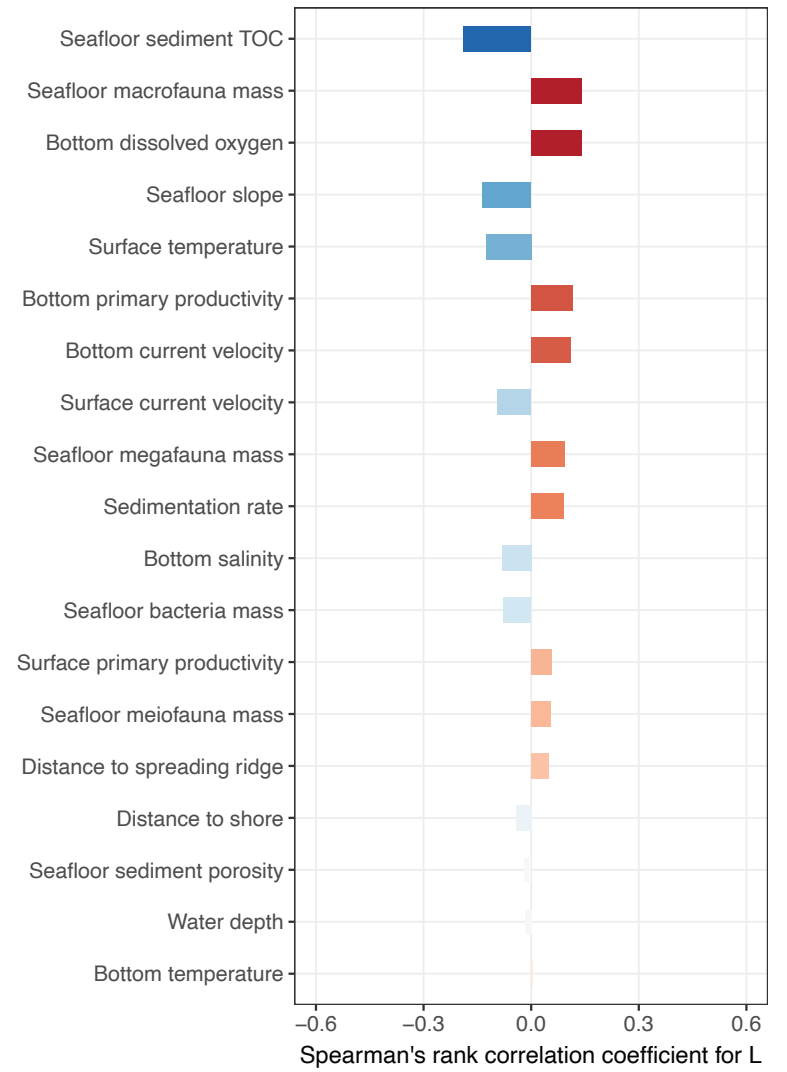


Figure

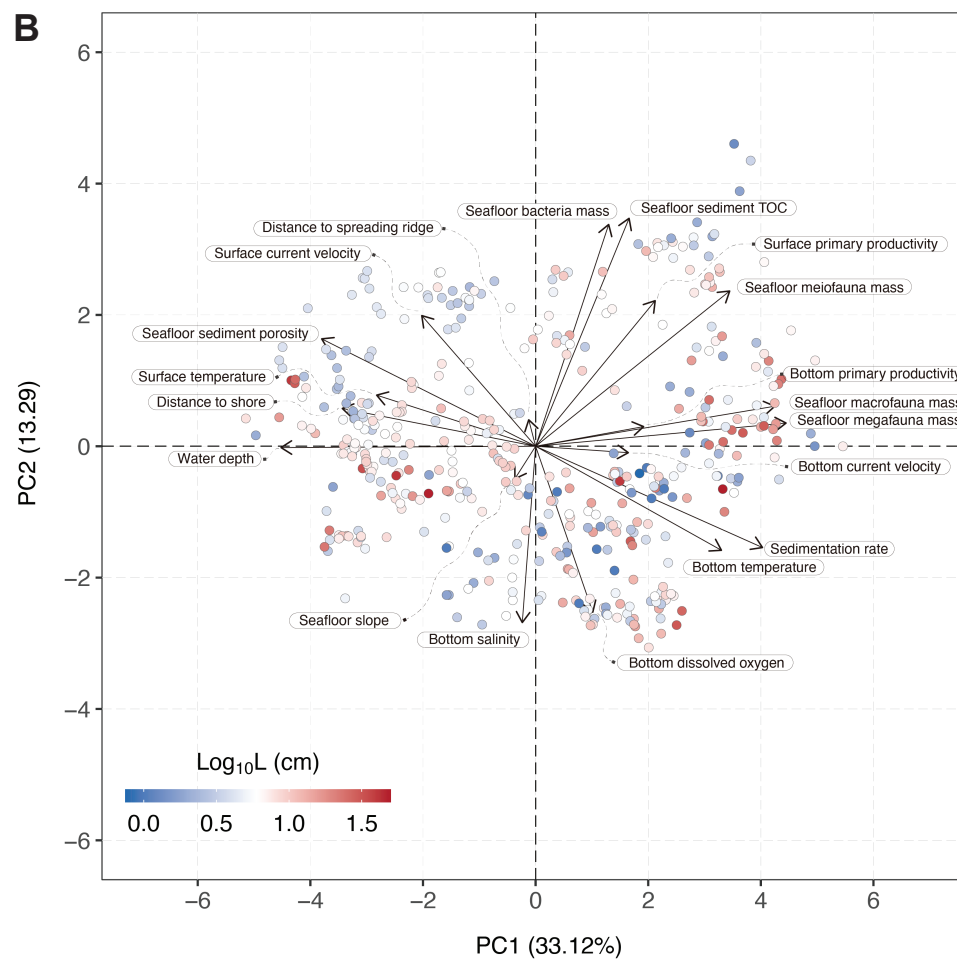
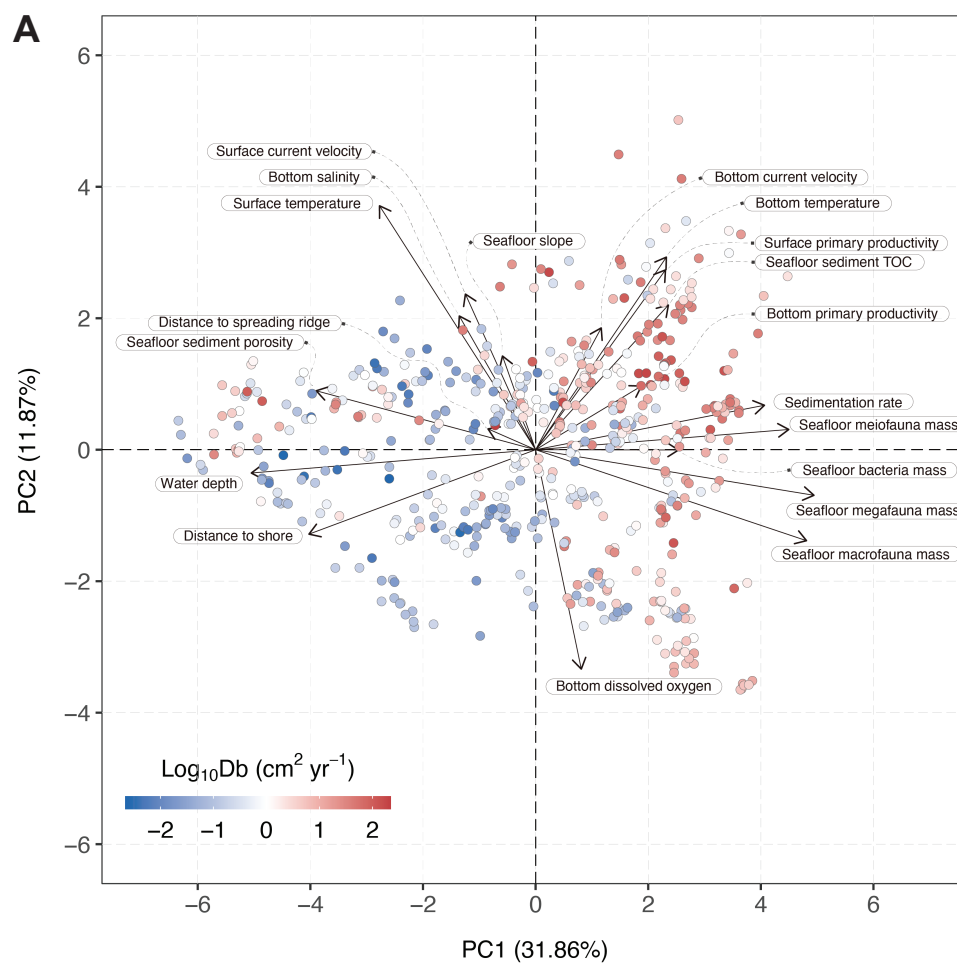
A



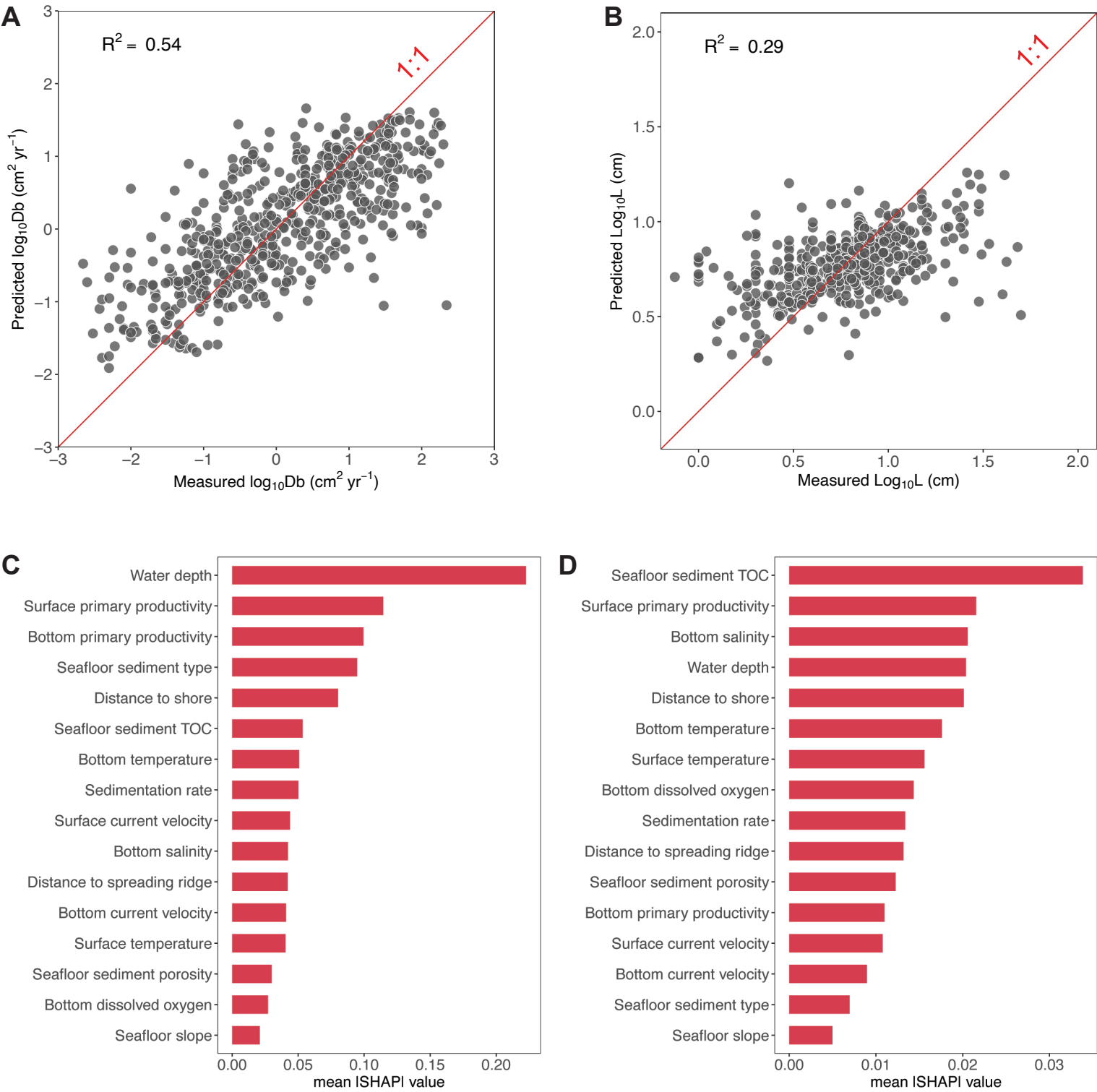
B



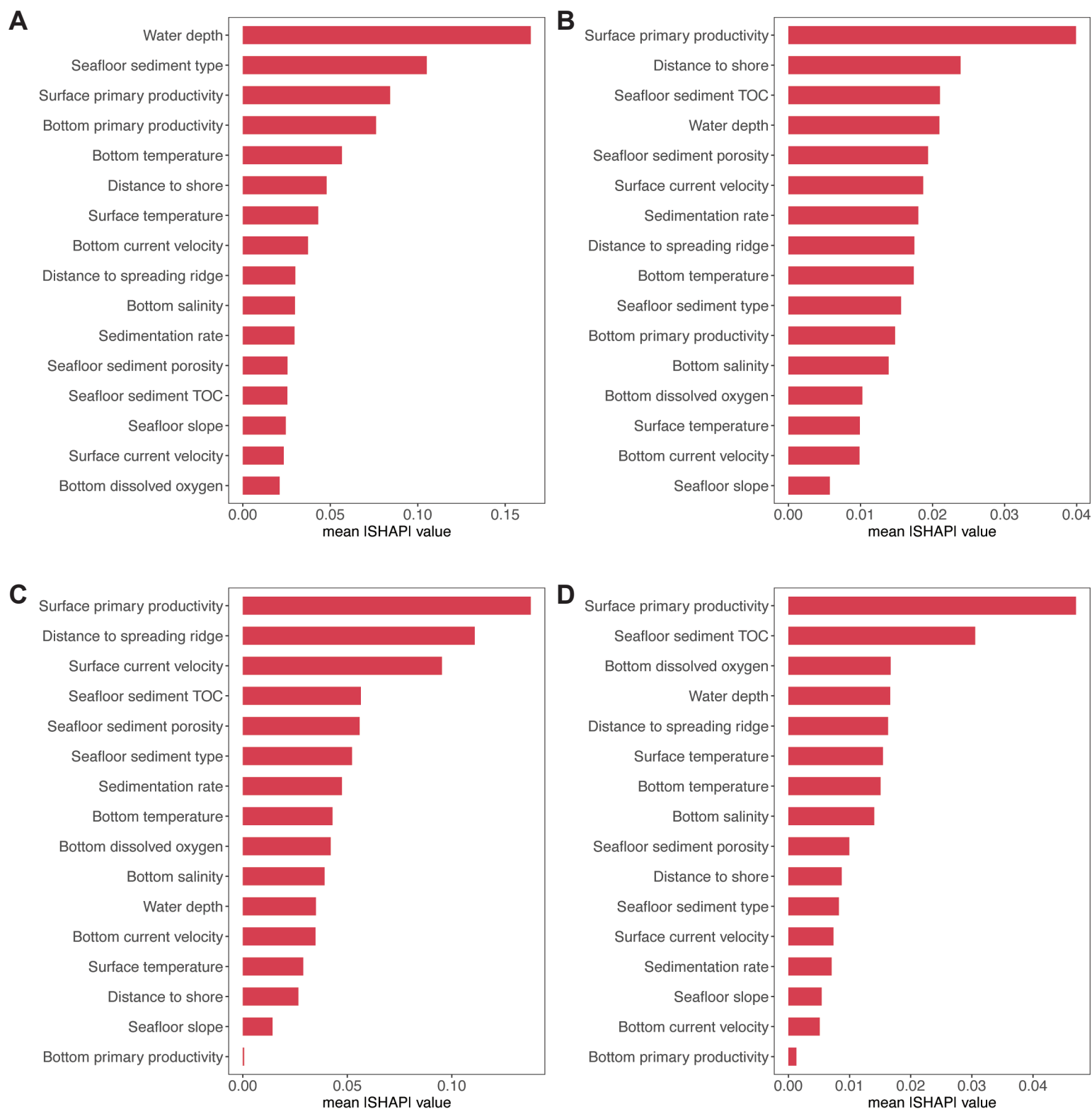
Figure



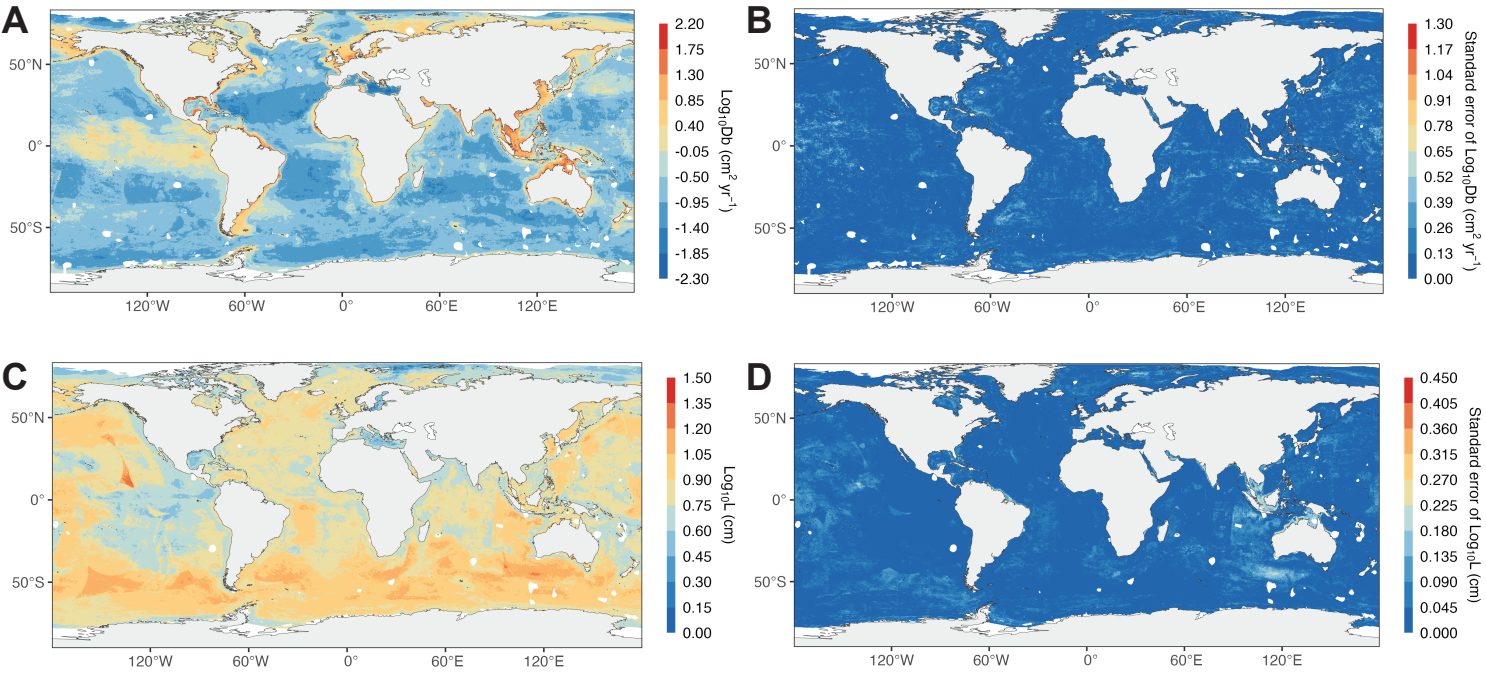
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Figure

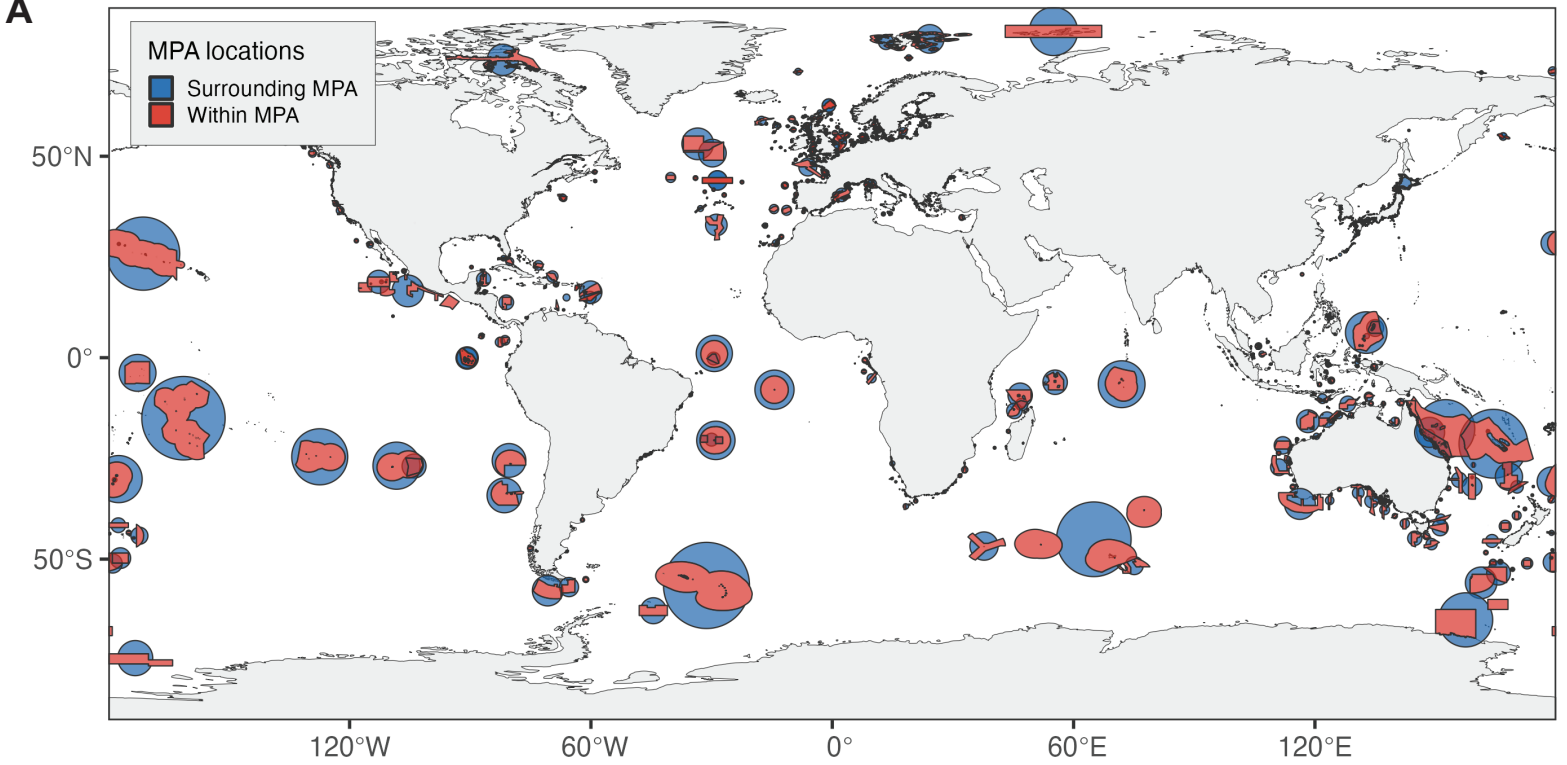


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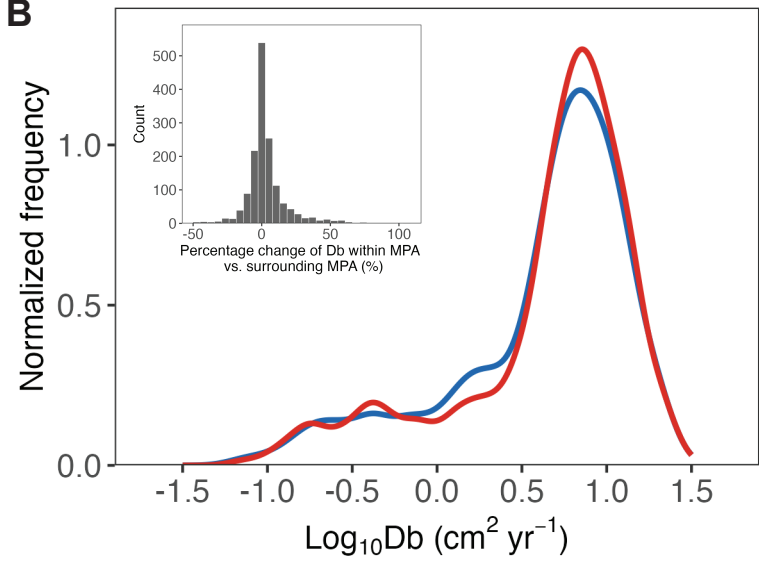


Figure

A

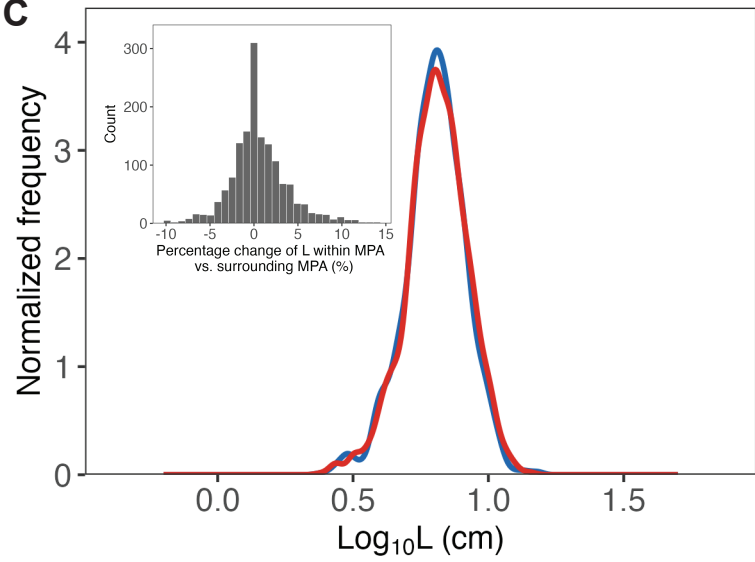


B



MPA locations: ■ Surrounding MPA ■ Within MPA

C



MPA locations: ■ Surrounding MPA ■ Within MPA

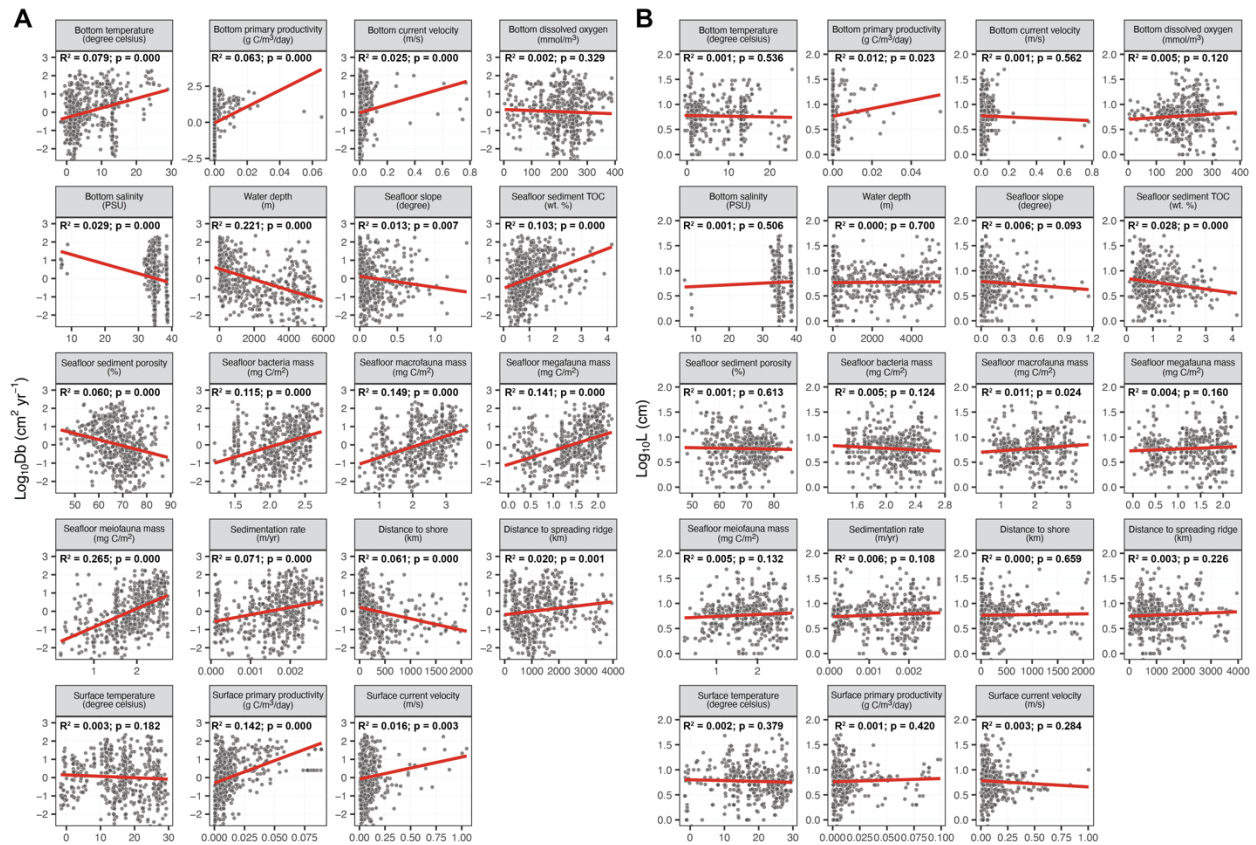


Figure S1. The relationship between bioturbation and spatially correlative environmental and ecological parameters. Related to Table S1 and Figure 2. (A) Bioturbation intensity (Db) (B) Mixed layer depth (L). In each panel, the red lines represent the linear regression between Db and the environmental and ecological parameters (shown solely for purposes of visualizing data trends). The associated R² value and p-value for each linear regression are noted in each panel. The source of each parameter is listed in Table S1.

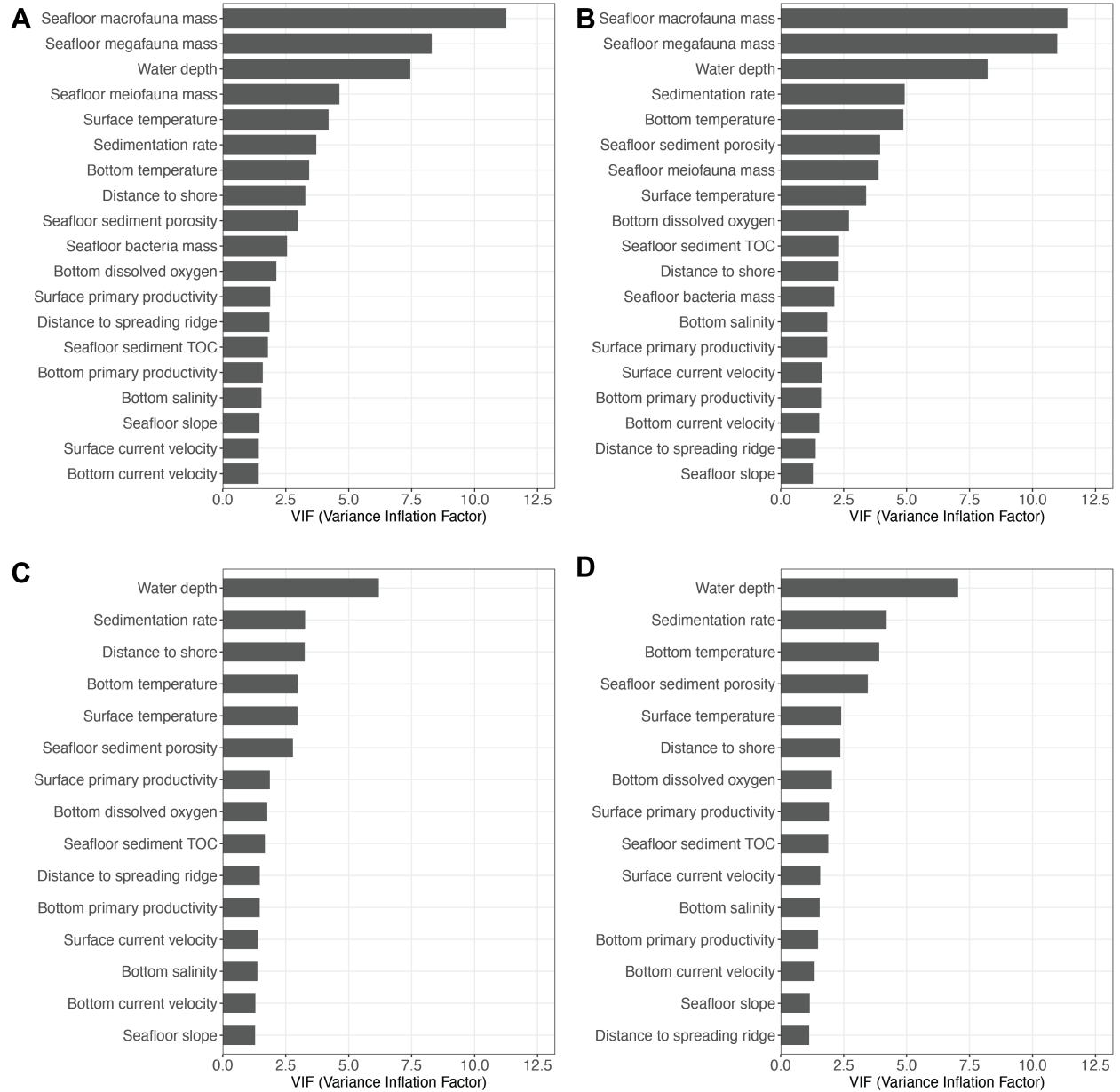


Figure S2. The Variance Inflation Factor (VIF) for predictor variables in the (A, C) Db and (B, D) L datasets. Related to Figure 2. The predictor variables include seafloor biomass parameters in (A) and (B) but exclude those parameters in (C) and (D).

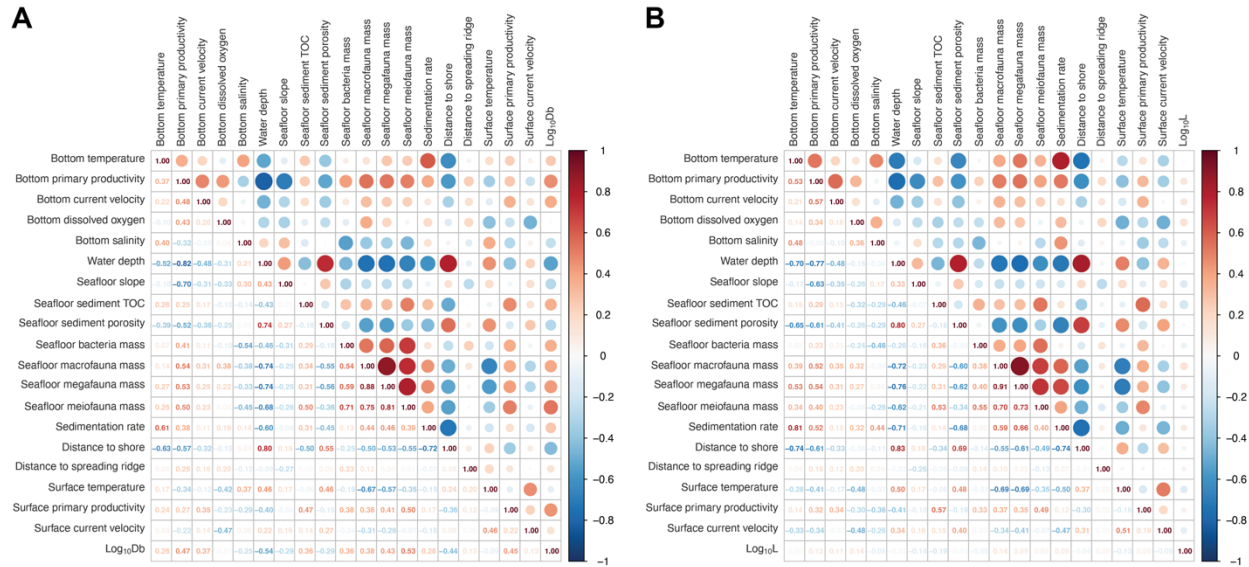


Figure S3. Pairwise spearman's rank correlation matrix of all spatially correlative environmental and ecological parameters and bioturbation. Related to Figure 2. We present the correlation matrix for (A) bioturbation intensity (Db) and (B) sediment mixed layer depth (L). Positive correlations are displayed in red and negative correlations in blue. Color intensity and circle diameter are proportional to the size of the correlation coefficients.

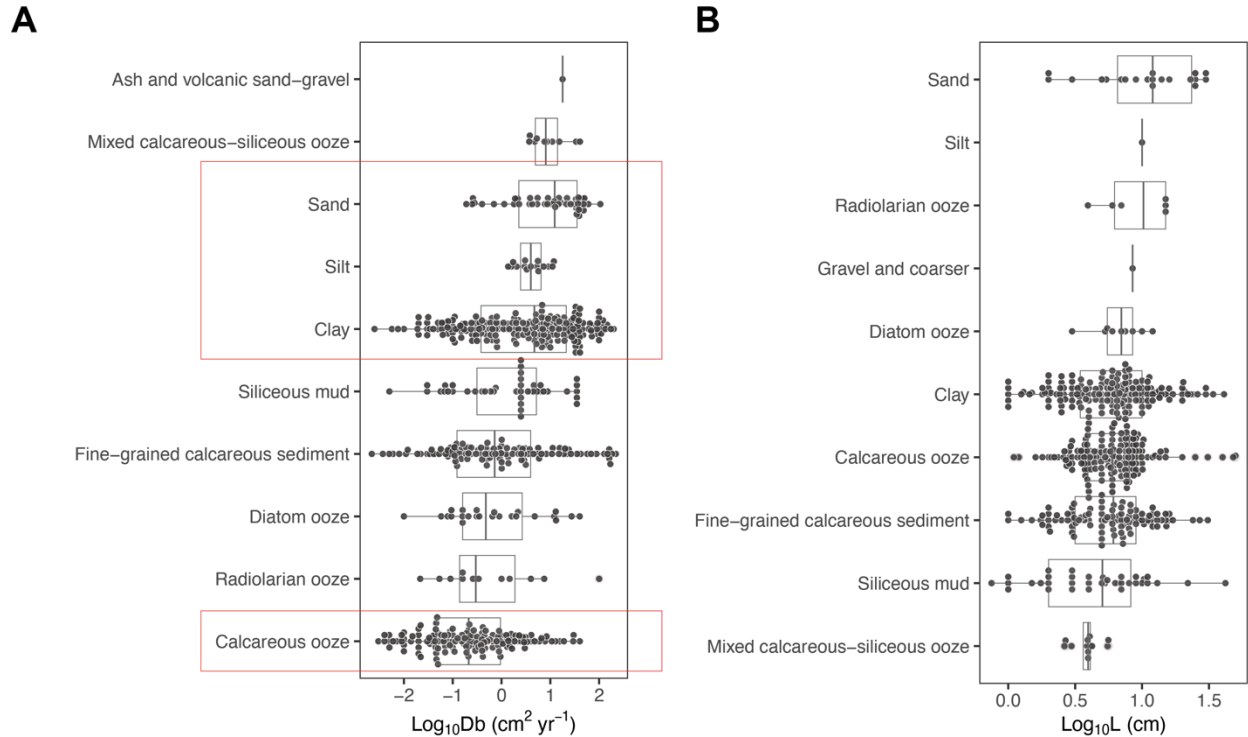


Figure S4. The relationship between seafloor sediment type and bioturbation (Db and L).

Related to Figure 3. In each case, the median is indicated at the midpoint, the upper and lower quartiles are indicated by the box margins, horizontal lines represent the spread of the data outside the quartiles, and closed circles indicate individual observations. In panel (A), the upper red rectangle highlights siliciclastic sands, silts, and clays, which are associated with higher bioturbation intensities. Conversely, the lower red rectangle highlights biogenic calcareous ooze, associated with the lowest bioturbation intensities. In panel (B), no rectangles are displayed as no discernible relationship could be identified between sediment mixed layer depth and sediment type.

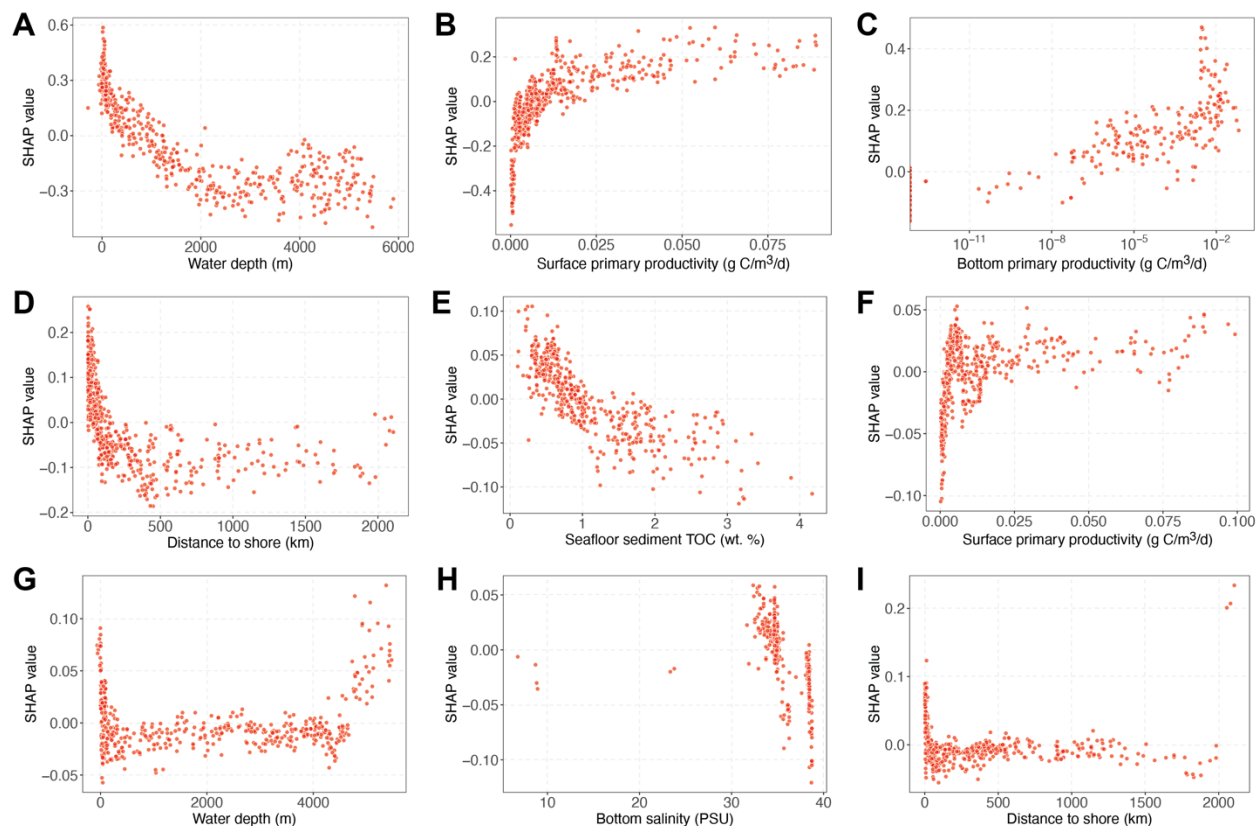


Figure S5. Relationship between SHAP values and the most important non-categorical predictor variables identified by machine learning. Related to Figure 4. (A–D) Relationship for Db. (E–I) Relationship for L. Features with positive SHAP values positively impact the prediction, while those with negative values have a negative impact. A positive correlation between a feature's values and its SHAP values can result in an increase in the prediction outcome as the feature value increases, while a negative correlation between a feature's values and its SHAP values can result in a decrease in the prediction outcome as the feature value increases.

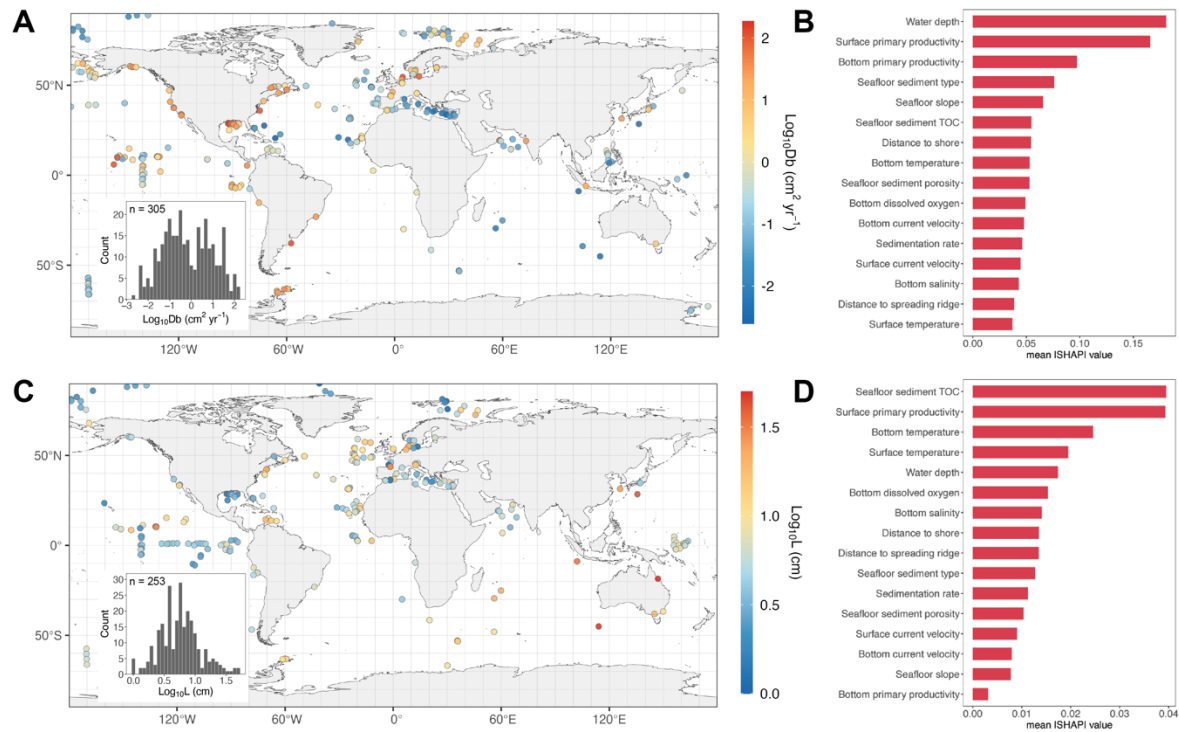


Figure S6. Distribution and machine-learning results for spatially weighted random sub-sampling of the full bioturbation dataset. Related to Figure 4. Site locations are shown for Db (A) and L (C). The mean absolute SHAP values (displayed in decreasing order, from top to bottom) associated with calculation of target variables Db (B) and L (D) from predictor variables are presented. We randomly sampled 50% of the entire Db or L dataset, with the selection probability scaled by the inverse density of samples within each $10 \times 10^\circ$ grid, across the entire global seafloor. The insets in (A) and (B) represent the histogram distributions of Db and L, with data split into 30 equal bins and the total number of observations indicated in the upper left portion of each inset.

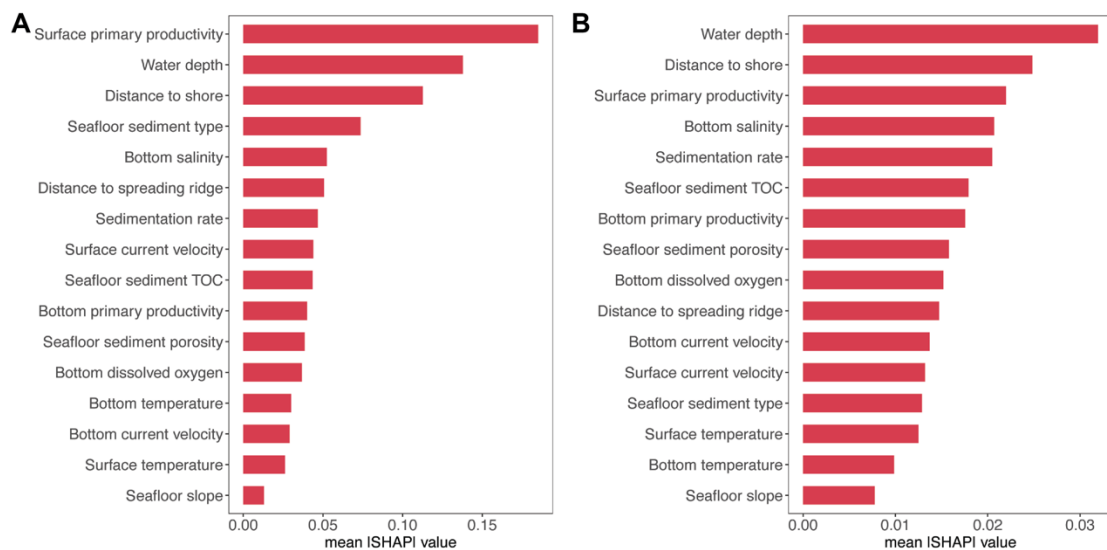


Figure S7. Machine-learning outcomes for Db and L for the screened ^{210}Pb data. Related to Figure 4. Mean absolute SHAP values (displayed in decreasing order, from top to bottom) associated with calculation of target variables Db (A) and L (B) from predictor variables are shown.

Variable Type	Database Name	Units	RMSE	R ²	Spatial Resolution	Time Span	Methods	Source
Environmental	Mean surface-water temperature	degree celsius	0.75	0.98	5 minutes	2000-2014	data re-analysis	S1
	Mean surface current velocity	m/s	-	-	5 minutes	2000-2014	data re-analysis	S1
	Mean bottom-water temperature	degree celsius	0.75	0.98	5 minutes	2000-2014	data re-analysis	S1
	Mean bottom current velocity	m/s	-	-	5 minutes	2000-2014	data re-analysis	S1
	Mean bottom-water salinity	practical salinity unit (PSU)	0.52	0.86	5 minutes	2000-2014	data re-analysis	S1
	Mean bottom-water dissolved oxygen concentration	mmol/m ³	23.5	0.92	5 minutes	2000-2014	biogeochemical model	S1
	Water depth	m	-	-	5 minutes	constant	empirical estimate	S1
	Seafloor slope	degrees	-	-	5 minutes	constant	empirical estimate	S1
	Seafloor sediment porosity	%	8.8	-	5 minutes	constant	machine learning	S2
	Seafloor sediment type	categories	-	-	6 minutes	constant	machine learning	S3
	Sedimentation rate	m/yr	-	-	1 degree	constant	empirical estimate	S4
	Minimum distance to spreading ridge	km	-	-	5 minutes	constant	empirical estimate	S2
	Minimum distance to shoreline	km	-	-	5 minutes	constant	empirical estimate	S1
Environmental/ Ecological	Mean surface-water primary productivity	g C/m ³ /day	-	-	5 minutes	2000-2014	biogeochemical model	S1
	Mean bottom-water primary productivity	g C/m ³ /day	-	-	5 minutes	2000-2014	biogeochemical model	S1
	Seafloor sediment TOC content	wt. %	-	0.61	5 minutes	constant	machine learning	S5
Ecological	Ocean ecoregion	categories	-	-	Shapefile	constant	empirical estimate	S6
	Ocean benthic biomass	mg C/m ²	-	0.8	1 degree	constant	machine learning	S7

Table S1 Summary of the source and spatial and temporal resolution of the environmental and ecological predictor variables used in data exploration and machine learning. “RMSE”

represents Root Mean Squared Error. “R²” represents the coefficient of determination.

“Methods” denotes how these parameters were generated. Data either span specific periods of time (listed in the “Time Span” column) or record a static (“constant”) snapshot. Cited references (“Source”) listed at the end of this document as well as in the main text.

Algorithms	Hyperparameters	Ranges	R ² for Db	R ² for L
Random Forest	number of trees	500-1500	0.54	0.29
	minimal node size	1-10		
Support Vector Machine	kernel	linear, polynomial, radial, sigmoid	0.53	0.24
	epsilon	0.1-0.5		
	gamma	0.1-1		
ElasticNet Regression	s	0-15	0.46	0.09
	alpha	0-1		
K-Nearest Neighbors (KNN)	number of neighbors	1-20	0.52	0.21
Gradient Boosting	number of trees	500-1500	0.47	0.14
	maximum tree depth	1-10		

Table S2 Summary of the model performance of five machine-learning algorithms for the test data using R² value as a metric of relative performance. Note: For Random Forest, the “number of trees” parameter specifies the number of trees to be ‘grown’ in the model. The “minimal node size” parameter specifies the minimum number of observations that must exist in a node for it to be split further. For Support Vector Machine, the “kernel” parameter allows specification of which kernel function to use for the regression task. The “epsilon” parameter defines the width of the ε -insensitive loss function used in the regression. The “gamma” parameter is the kernel coefficient for the Radial Basis Function (RBF) kernel. For ElasticNet regression, the “s” parameter represents the regularization term λ in the elastic net penalty. It controls the amount of shrinkage applied to the coefficients, with higher values resulting in more regularization. The “alpha” parameter controls the mixing between L1 and L2 regularization. An alpha of 1 corresponds to Lasso regression (L1 regularization), and an alpha of 0 corresponds to Ridge regression (L2 regularization). Values between 0 and 1 are associated with a mixture of each. For K-Nearest Neighbors (KNN), the “k” parameter specifies the number of nearest neighbors to consider when making a prediction. For Gradient Boosting, the “number of trees” parameter specifies the total number of boosting iterations to be carried out or, equivalently, the total number of trees to fit. The “maximum tree depth” parameter specifies the maximum depth

of each individual regression estimation tree. For each algorithm, we iterated over a series of hyperparameter combinations within their ranges to find the optimal set. We then used the optimal set of hyperparameters to train each model and validate its performance (the R^2 value) using 10-fold cross validation.

Supplemental references

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