1Unifying latitudinal gradients in range size and richness across marine and 2terrestrial systems

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8Supplementary online material

9Supplementary methods

Data. Species range limits for marine bivalves and terrestrial birds are 11available at Data Dryad at dx.doi.org/10.5061/dryad.p0q25. Temperature gradients in 12Figure 3 A-C represent the annual mean daily sea-surface and air temperature (the 13daily mean temperature averaged over a full year) in 1° latitude bands. Sea-surface 14temperature (SSTs) at a 9*9 km² resolution was obtained from the Moderate 15Resolution Imaging Spectroradiometer (MODIS) and rescaled to a 1°*1° resolution. 16Air temperatures were obtained from CliMond database (Kriticos et al. 2012). The 17tropical-temperate boundaries in the Eastern Pacific are at 6°S and 23.5°N and in the 18Western Atlantic at 23°S and 28°N (Spalding et al. 2007). In the global-scale analyses 19of bivalves, the boundaries are set at 25°S and 30°N. In analyses of birds, the 20boundaries are set at 30°S and 30°N.

21 *Relation between latitude and median range size.* We use generalized least-22squares to compare the relation between latitude and median latitudinal range size 23between hemispheres and between birds and bivalves. This method minimizes the 24correlation among the residuals due to the spatial structure of the data (Fig. S2), and is 25implemented in the function gls in the R package nlme (Pinheiro et al. 2013). We fit

26 five alternative models accounting for the relationship between the variance in the 27 residuals and latitudinal distance among 1° bands in semi-variograms (linear, 28 Gaussian, spherical, exponential, and rational quadratic functions; Begueria and 29 Pueyo 2009), and selected the best-fitting model based on the Akaike information 30 criterion (AIC) for each of the comparisons (Table S1).

31 *Contribution of nestedness to dissimilarity.* We subsample 20 bands (without 32replacement) for 1,000 times from all latitudinal 1° bands within each hemisphere. For 33each of the 1,000 generated datasets, we then compute species-level and genus-level 34proportional contribution of nestedness to total dissimilarity (Baselga 2012). We use 35the function beta.sample in the R package betapart (Balselga and Orme 2012). The 36proportion of cases in which the species-level contribution to nestedness is higher 37than the genus-level contribution thus represents the probability that the increase in 38nestedness from species to genus levels occurs by chance.

39 *Relation between per-genus species range size and genus range size.* We use a 40bootstrapping test to account for the dependence of genus range size on the median 41latitudinal range size of its species, as well as the mean distance among latitudinal 42midpoints of its species, because species range size and among-species distance must 43be less than or equal to genus range size (Novack-Gottshall and Miller 2003). Species 44range size and genus range size was resampled (with replacement) independently for 45each genus for 1,000 times, and resulted species range sizes (or among-species 46distances) were required to be equal to or greater than the corresponding genus range 47size in each iteration. Using these simulated data, we calculated the rank-order 48correlations as a null expectation, and compared the 1,000 null correlations with the 49observed rank correlation.

50Supplementary results and discussion

51 Species range-size gradients. Marine temperatures vary only slightly over a 52wide latitudinal range around the equator especially in the Western Atlantic (and 53Western Pacific) because the Earth's rotation drives warm tropical waters toward the 54poles along the western oceanic margins. In contrast, cold polar water are driven 55toward the equator in the Eastern Pacific, where the tropics are further truncated by 56cold-water upwelling at 6° S. Therefore, tropical zones with more or less constant sea-57surface temperature (25-27°C) are ~5,000-6,000 km broad along the western margins 58but only ~2,500-4,000 km broad along the eastern margins. Therefore, tropical waters 59in the Western Atlantic permit broad geographic ranges even in the absence of large 60thermal tolerances. In contrast, the cool temperatures just south of the equator in the 61eastern Pacific restrict the latitudinal span of tropical waters and reduce tropical 62species range sizes by ~2,000 km relative to the Western Atlantic (Fig. 3D-E). The 63actual peak in range size in the Eastern Pacific is then located at 10°S, and is 64generated by few broad-ranging species inhabiting tropical and temperate latitudes 65(see also Tomasovych et al. 2015).

Genus range-size gradients. Globally and along the New World coasts, Genus range sizes of bivalves significantly increase from within the tropics to e8at the highest latitudes. In birds, median genus range size is $\leq \sim$ 4,700 km in the 99tropics and increases significantly to the highest southern latitudes; in the Northern 70Hemisphere, it peaks at ~8000 km near 55°N but then declines towards higher 71latitudes. In the New World, the temperate Northern Hemisphere median again 72exceeds tropical values, but the maximum occurs at ~30°N, with a northerly decline. 73Nonetheless, narrow-ranging genera (which we define as encompassing < 60° of 74latitude, i.e., more than ~6,600 km) are concentrated in the tropics for New World 75birds and bivalves, by a factor of 3-5 relative to 45°N (Fig. S7). For New World birds

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76and bivalves, narrow-ranging genera (which we define as encompassing < 60° of 77latitude, i.e., less than ~6,600 km) are concentrated in the tropics by a factor of 3-5 78relative to 45°N (Fig. S7). Broad-ranging genera (> 60° of latitude) show shallower 79richness gradients in the Eastern Pacific bivalves and New World birds. In contrast, a 80greater number of broad-ranging bivalve genera occur in the tropical Western 81Atlantic, likely due to the shallower temperature gradients, resulting in tropical 82climatic conditions that are latitudinally broader. The increase in median genus range 83size of New World birds from 0° to 30°N (grey line in Fig. 3F) is produced by the 84declining richness of broad-ranging genera briefly exceeding the declining richness of 85narrow-ranging genera, which cluster around the equator. Overall, there is a steep 86decline in the number of narrow-ranging genera at around 25°N to 30°N both on land 87and in the sea (Fig. S7). The deserts of Mexico, Argentina, and Chile on land, and 88strong thermal gradients in the sea apparently restrict many genera to the tropics (Fig. 893A-C).

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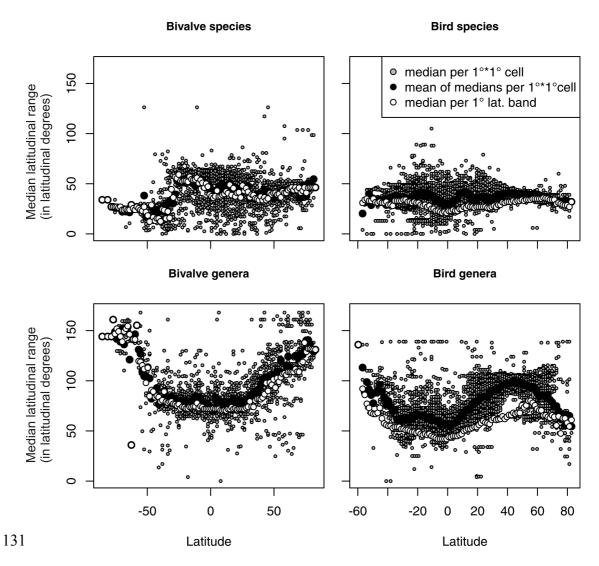
118Supplementary tables and figures

119Table S1 - The generalized least-square estimates of slope of the median latitudinal
120range size over latitude at the species and genus level, and the selected correlation
121structures that minimize the spatial correlation among the residuals. The slopes are
122significantly negative in bivalve species at regional scales and in the Southern
123Hemisphere at the global scale. The slopes become mostly positive in both groups at
124the genus level.

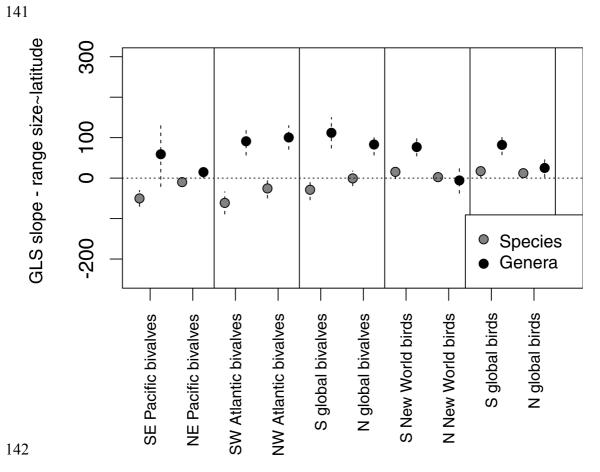
| | Southern Hemisphere range- size~latitud e slope | p | Cor. Structure | Northern Hemisphere range- size~latitude slope | p | Cor. Structure |
|---------------------------|---|---------|-------------------|---|---------|-------------------|
| Species | | | | | | |
| Eastern Pacific bivalves | -50.2 | <0.0001 | Spherical | -9.9 | <0.0001 | Spherical |
| Western Atlantic bivalves | -61.3 | <0.0001 | Spherical | -25.5 | 0.0377 | Spherical |
| Global bivalves | -28.8 | 0.0263 | Spherical | -0.7 | 0.9406 | Spherical |
| New World birds | 15.4 | 0.0824 | Spherical | 2.4 | 0.6243 | Exponentia |
| Global birds | 17.2 | 0.0315 | Spherical | 12.1 | 0.0040 | Spherical |
| Genera | | | | | | |
| Eastern Pacific bivalves | 59.0 | 0.1527 | Spherical | 14.9 | 0.0452 | Spherical |
| Western Atlantic bivalves | 91.0 | <0.0001 | Spherical | 100.4 | <0.0001 | Spherical |
| Global bivalves | 112.0 | <0.0001 | Spherical | 83.0 | <0.0001 | Spherical |
| New World birds | 76.7 | <0.0001 | Ratio | -5.7 | 0.7233 | Spherical |
| Global birds | 82.0 | <0.0001 | Ratio | 25.1 | 0.0314 | Spherical |

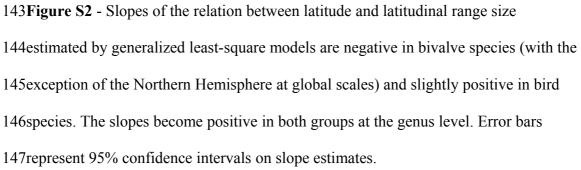
Table S2 - Wilcoxon rank-sum test comparing per-genus richness of tropical species
127belonging to bridge and tropics-only genera (Tropics) and per-genus richness of
128extratropical species belonging to bridge and extratropics-only genera (Extratropics).
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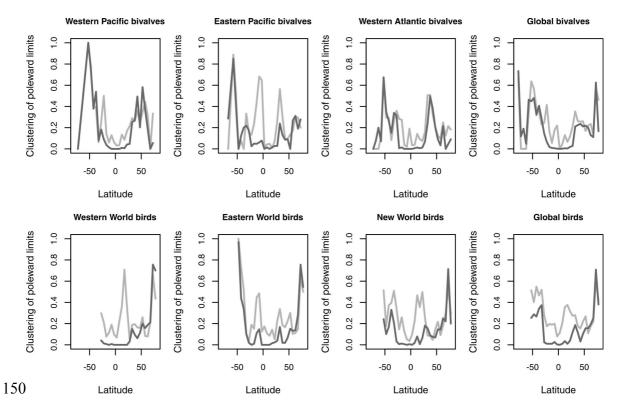
| | Mean richness of restricted genera | Mean richness of bridge genera | W stat. | p-value |
|--|---------------------------------------|-----------------------------------|---------|---------|
| Tropics-Global bivalves | 1.58 | 3.68 | 18957.5 | <0.0001 |
| Extratropics-Global bivalves | 1.67 | 2.92 | 17500 | <0.0001 |
| Tropics-Global birds | 2.59 | 6.59 | 196847 | <0.0002 |
| Extratropics-Global birds | 1.58 | 2.86 | 9487 | <0.000 |
| Tropics-Western Atlantic bivalves | 1.07 | 1.48 | 929.5 | 0.0111 |
| Extratropics-Western Atlantic bivalves | 1.34 | 2.19 | 2140 | 0.0045 |
| Tropics-Eastern Pacific bivalves | 1.18 | 1.50 | 989 | 0.0221 |
| Extratropics-Eastern Pacific bivalves | 1.54 | 2.16 | 3705 | 0.0038 |
| Tropics-New World birds | 2.78 | 5.47 | 47801 | < 0.000 |
| Extratropics-New World birds | 1.86 | 2.56 | 3120.5 | 0.0080 |



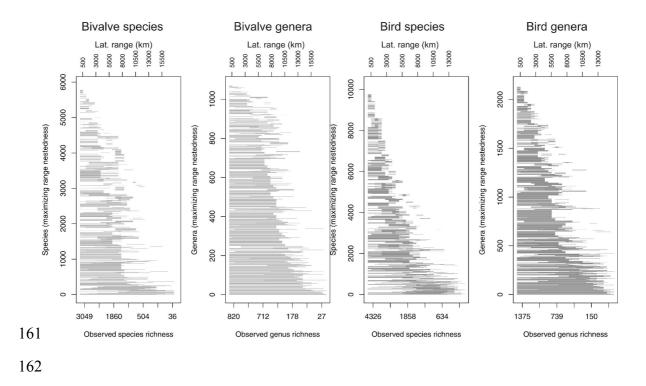
132Figure S1 - The effects of spatial resolution on latitudinal gradients in range size of 133species and genera in bivalves and birds. Median range size per 1° latitudinal band 134(white points) tends to be lower than the per-band average (black points) of median 135range size per equal-area ~1° cells (25,000 km²) because narrow-ranging species 136contribute less to individual 1° cells than to full 1° bands. This offset is larger in birds 137than in bivalves. Gray points show individual median range size for all 1° cells. In 138contrast to the main body of the paper where gaps in species distributions were 139ignored, i.e., species occurrences were counted in all latitudinal bands that are situated 140within the latitudinal range limits of the species, these plots show raw data.



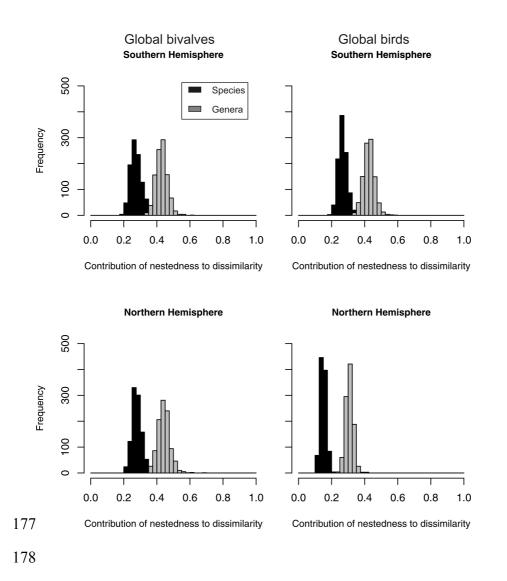




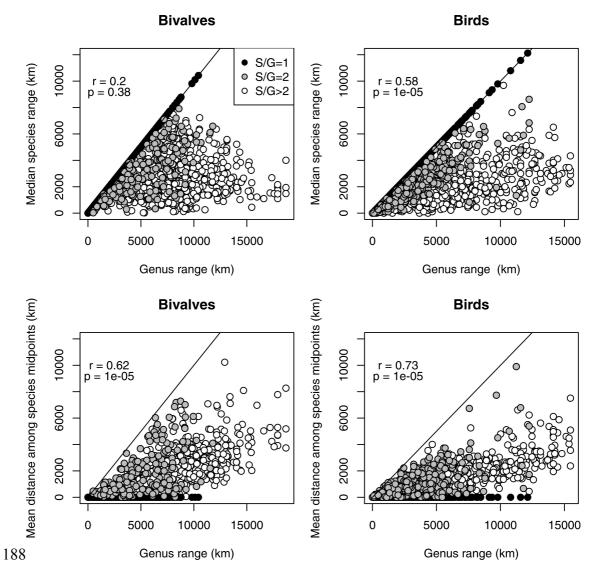
151**Figure S3** Clustering of poleward range limits of genera at the resolution of 5° bands 152(i.e., the number of genera with poleward range limits relative to all genera in a given 153band). In bivalves, genus range limits are rare within the tropics, in contrast to some 154major clustering of range limits of bird genera. Here three additional regions are 155included beyond those in the main text, the Western Pacific for bivalves, and the 156Western Old World (including the western Palearctic, Africa, and Saharo-Arabian 157areas) and Eastern Old World (including the eastern Palearctic, Oriental, Indo-Pacific, 158Australia and New Zealand) for birds to separate the tropical areas of Africa and the 159Indo-Pacific according to the biogeographic divisions suggested by Holt et al. (2013).



163**Figure S4**. Visualization of range configuration and Rapoport patterns: species show 164a pattern of turnover, with broad-ranging bivalve species at low latitudes replaced by 165narrow-ranging bivalve species at high latitudes (inverse Rapoport's pattern), and 166broad-ranging bird species at higher latitudes replaced by narrow-ranging bird species 167at low latitudes (Rapoport's pattern). However, bivalve and bird genera show 168predominantly nested ranges - genera with broad ranges are found at most latitudes 169whereas narrow-ranging genera are much more common in the tropics. The columns 170are 5° latitudinal bands (ordered by increasing richness). The rows represent species 171ordered such that broad-ranging taxa are at the bottom and taxa with narrower ranges 172are more frequent towards the top (i.e., they are nested within broad-ranging species if 173they occupy latitudes within their range). This ordering is based on an iterative 174procedure, attempting to maximize both the gradient in richness and in nestedness of 175ranges.



179**Figure S5** - At species level, turnover (Simpson dissimilarity) contributes with 75% 180(Southern Hemisphere) and 85% (Northern H.) to total (Sorenson) dissimilarity in 181birds, and with 73% (Southern H.) and 72% (Northern H.) to total dissimilarity in 182bivalves. At genus level, turnover contributes with 57% (Southern H.) and 69% 183(Northern H.) to total dissimilarity in birds, and with 57% (Southern H.) and 55% 184(Northern H.) to total dissimilarity in bivalves. The mean contribution of nestedness 185to total dissimilarity is significantly larger at genus than at species level in both 186hemispheres and in both groups (all p < 0.001). In both groups, turnover rather than 187nestedness thus accounts for species range-size gradients.



189**Figure S6** *Top row:* Spearman rank correlation (*r*) between genus latitudinal range 190size and per-genus median species latitudinal range size for bivalves (left) and birds 191(right). *Bottom row:* Per-genus mean latitudinal distance among species centroids 192against genus latitudinal range size for bivalves (left) and birds (right). Similar results 193apply to three regional datasets. Genera are subdiviced according to their per-genus 194species richness (S/G) into three categories, including monospecific genera (where 195species and genus range sizes are the same by default, black points), genera with two 196species (gray points), and genera with more than two species (white points).

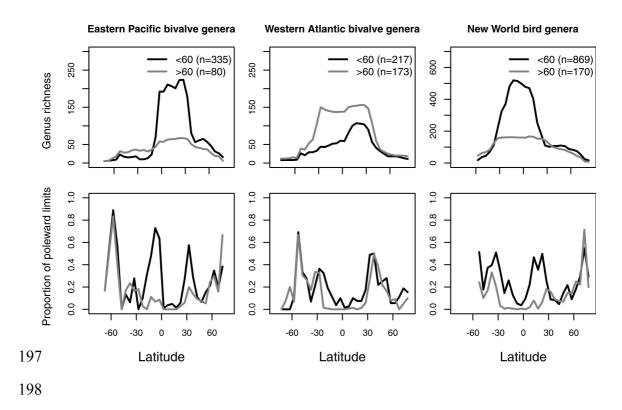


Figure S7 - Latitudinal gradients in genus richness and the proportion of poleward 200range limits (i.e., the number of genera with poleward range limits relative to all 201genera in a given band) of two groups of genera delimited by having a range size 202smaller or larger than 60° (here, at the resolution of 5° bands).