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Global variation in elevational diversity patterns

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While horizontal gradients of biodiversity have been examined extensively in the past, vertical diversity gradients (elevation, water depth) are attracting increasing attention. We compiled data from 443 elevational gradients involving diverse organisms worldwide to investigate how elevational diversity patterns may vary between the Northern and Southern hemispheres and across latitudes. Our results show that most elevational diversity curves are positively skewed (maximum diversity below the middle of the gradient) and the elevation of the peak in diversity increases with the elevation of lower sampling limits and to a lesser extent with upper limit. Mountains with greater elevational extents, and taxonomic groups that are more inclusive, show proportionally more unimodal patterns whereas other ranges and taxa show highly variable gradients. The two hemispheres share some interesting similarities but also remarkable differences, likely reflecting differences in landmass and mountain configurations. Different taxonomic groups exhibit diversity peaks at different elevations, probably reflecting both physical and physiological constraints.

Mountane regions harbor more than half of the world's biodiversity hotspots and recent research on biodiversity patterns has been notable for an increase in research on elevational patterns on mountains¹. Mountains provide unique opportunities as 'natural experiments' for testing ecological theories and in particular for studying the effects of climate change because they present gradients in key abiotic features such as temperature and available moisture. Recent efforts have generated interesting and sometimes conflicting results, and debates on the generality of the frequently observed unimodal ("hump-shaped"²) curves and the underlying mechanisms have not been fully resolved. Indeed, despite increased research in this area in recent years, employing markedly improved techniques and greater sampling intensity, much inconsistency and debate remains both in pattern description and interpretation. For example, surprisingly little effort has targeted how elevational diversity patterns might vary in the Northern and Southern hemispheres and across latitudinal zones³⁻⁷. To tackle these problems, detailed comparisons are needed over distinct (replicate) elevational gradients across the globe.

The upper elevational limit for phanerogams varies as a function of latitude and generally reflects limits to physiologic tolerance¹. Moreover, similar elevational ranges in different regions are likely to exhibit different underlying gradients⁸, reflecting regional climate and geography; cold-temperate mountains lack the warm climate characteristic of lower elevations at lower latitudes, and temperatures at a tropical treeline might reflect those at the base of cold-temperate mountains. Additionally, the effect of aspect is greatly reduced on tropical mountains relative to temperate ones. As a result of these latitudinal differences, structurally identical mountains located in different latitudinal zones would be expected to show different diversity patterns.

Sampling issues have rightly been a major focus in biogeographical research^{2,9-13}. Relatively few studies span the full extent of available elevational gradients¹⁰, and many studies on latitudinal gradients are constrained to cover only a portion of the full gradient (e.g., a subset of 0–90° N or S). Constraints include habitat alteration at lower elevations (e.g., agriculture, urbanization), challenges of accessing higher elevations, and the uneven distribution of landmass across latitude (and across the equator). Resulting constraints on the latitudinal or elevational extent sampled may influence ecological insights; additionally, patterns of elevational diversity gradients also depend on the specific hierarchical species groups (e.g., all seed plants vs. subsets such as trees) and their sizes (number of species). Finally, geographic comparisons assume comparable sampling intensity across all sites; this is rarely achievable due to different research objectives and funding constraints, but it is incumbent on geographers to



Figure 1 | The distribution of mountains/regions for which diversity data from 443 elevational gradients are examined in this study (made using Google Maps).

assure that spatial interpretations are biologically real and not inconsequential results of spatially varying sampling effort.

We compiled a comprehensive dataset on biota across elevational gradients on many of the major global mountain ranges (Fig. 1), and we use this to advance understanding of two fundamental questions. First, how do elevational diversity gradients/patterns vary across the globe, especially between the two hemispheres and across latitudes and among taxonomic groups? Second, in addition to climate and human impacts, what regional factors and sampling issues might explain variation among elevational patterns studied to date? We emphasize recent studies as these incorporate improved sampling procedures (e.g., intensive as well as extensive sampling) thereby minimizing concerns over differential effort, but we emphasize that further progress in this field calls for attention to standardization of sampling efforts. Our ultimate goal is to improve understanding of how elevational patterns vary across the globe; more refined evaluations to identify and discuss specific mechanisms will require high-resolution climate data and geospatial analyses to accurately account for unique features of each mountain/region involved.

Results

Globally, 279 of 443 elevational diversity gradients examined (63%) were unimodal. This was followed in the Northern Hemisphere by negative, positive, none, either unimodal or negative (“unimodal/negative” hereafter), and bimodal or polymodal (“bimodal/polymodal” hereafter), and in the Southern Hemisphere by negative, none, positive, and unimodal/negative (Fig. 2). The proportions of the gradients corresponding to distinct patterns was similar between plants and animals ($X^2 = 3.76$, $P = 0.71$), and the proportion of gradients exhibiting unimodal patterns was similar for plant functional groups (shrubs, 9/12; herbs, 13/22; ferns, 29/48; trees, 13/25; $X^2 = 0.43$, $P = 0.93$). Among animals, 42 of 76 (55%) invertebrates and 73 of 115 (64%) vertebrates showed unimodal patterns ($X^2 = 0.97$, $P = 0.57$). Among vertebrates, non-flying mammals had the highest proportion of unimodal pattern (76%), followed by bats (50%) and birds (40%).

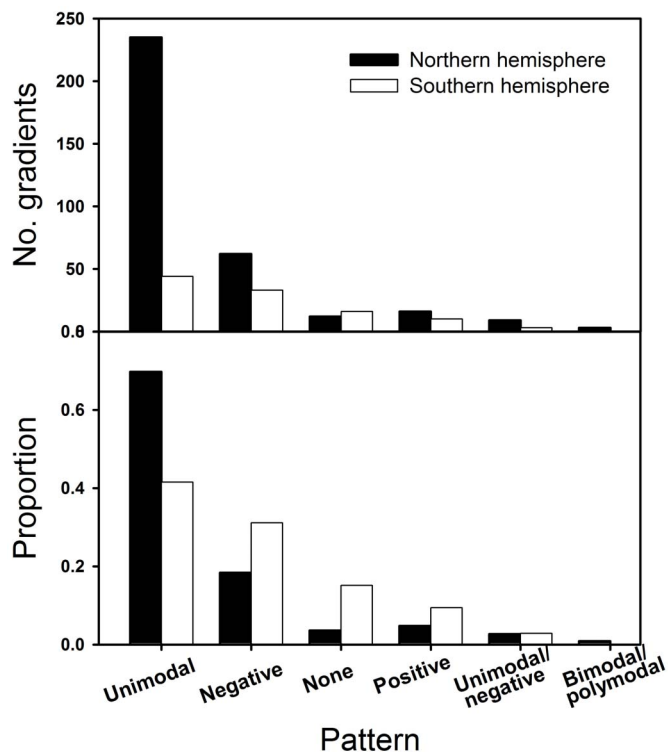


Figure 2 | Top: Number of gradients (out of 443) in each pattern category in Northern and Southern hemispheres. Bottom: Proportion (0–1) of each of the six patterns among the studies gradients in Northern vs. Southern hemisphere. See Methods for detailed pattern classification and descriptions (symbols).

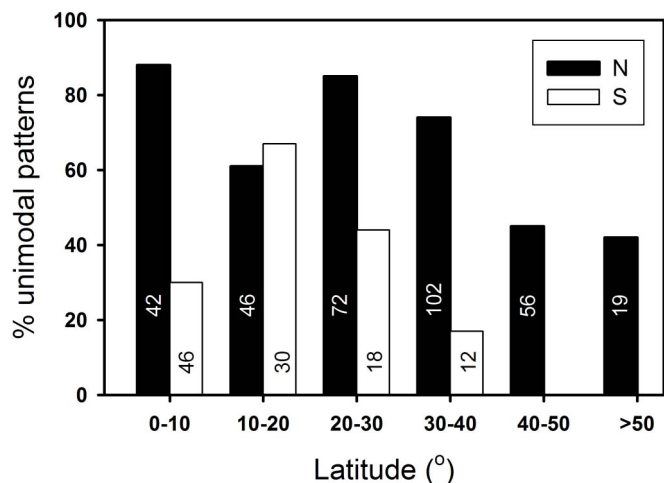


Figure 3 | Distribution of examined diversity gradients with unimodal pattern across latitudes in both Northern and Southern hemispheres. The number in each bar represents the no. of gradients in each category. Except for the “1–10°N” category in Northern Hemisphere, mid-latitudes had more gradients with unimodal pattern.

While some general trends were similar between the two hemispheres (e.g., elevational peaks of diversity declined with latitude), there was also remarkable *hemispheric asymmetry* in certain other aspects. For example, there were proportionally more unimodal patterns among studies in the Northern Hemisphere relative to the Southern Hemisphere (235/337 or 70% vs. 44/106 or 42%; $\chi^2 = 35.56$, $P < 0.0001$; this excludes mountains $> 45^\circ$ since these are not available in any Southern Hemisphere data; Fig. 3). Also, the diversity peaks were positioned at significantly higher elevations in the Northern than in the Southern Hemisphere ($\bar{x} = 1508$ m vs. 1141 m; $t = 3.9$, $P < 0.001$). With the exception of the lowest latitude bracket (0–10°N) in the Northern Hemisphere, mid-latitudes (20–40°N and 20–30°S) had more gradients with unimodal pattern than either lower or higher latitudes (Fig. 3). On average, the highest elevations of species diversity peaks occurred at a much higher latitude in the Northern hemisphere (30°N) than in the Southern hemisphere (5°S). In the Northern Hemisphere, the lowest and highest sampling limits occurred around 25–30° and the greatest elevational extents of the studied mountains at about 30°. In contrast, the corresponding values were lower (around 5–15°) in the Southern

Hemisphere. The low, mid, and high elevations were 668, 1911, and 3154 m in the Northern Hemisphere vs. 469, 1623, and 2778 m in the Southern Hemisphere; and all three were significantly different between the two hemispheres ($t = 2.68$, 2.96, and 3.81, $P = 0.008$, 0.003, and 0.0002, respectively; see also Figs. 4, 6, S1); in contrast, the elevational extents sampled do not differ (2505 vs. 2353 m; $t = 1.09$, $P > 0.05$; Fig. 4, Fig. S1).

In general, mountains presenting greater elevational extent were more likely to display unimodal patterns. All elevational-extent categories > 1000 m had a high percentage of unimodal patterns; ranges exceeding 4000 m in extent presented the greatest proportion of unimodal patterns, while all lesser extents (e.g., 1000–2000, 2000–3000, or 3000–4000 m extent) had very similar proportions of unimodal patterns (Fig. 5, S1).

About 75% (208 out of 279) of the unimodal gradients in our data set were positively skewed (e.g., peak diversity below the elevational midpoint; Fig. 6). The elevations of diversity peaks increased with the elevations of both lowest and highest sampled limits although the effects of the latter were weaker (Fig. 6).

Elevational diversity patterns varied by taxonomic group. The elevations of diversity peaks for plants were higher than those for animals ($\bar{x} = 1530$ vs. 1274 m; $t = -3.27$, $P < 0.001$), although these two groups did not differ in the lowest sampling limits (634 vs. 577 m for plants vs. animals; $t = 0.91$, $P = 0.36$), the highest sampling limits (3135 vs. 3007 m; $t = 1.16$, $P = 0.25$), and the sampled elevational extents (2508 vs. 2427 m; $t = 0.68$, $P = 0.5$). Among animals, the diversity peaks for vertebrates were at significantly higher elevations than those for invertebrates ($\bar{x} = 1422$ vs. 1047 m; $t = -3.36$, $P < 0.001$) although there was no difference in either the sampled elevational extents or the lowest/highest sampled elevations between the two groups. The elevation of diversity peaks for non-flying mammals ($\bar{x} = 1906$ m) was much higher than that for either bats ($\bar{x} = 924$ m) or birds ($\bar{x} = 980$ m; $t = -5.96$ and -4.52 , respectively; $P < 0.0001$) but there was no difference between the latter two groups (Fig. 7). Among plants, the diversity peaks of herbs were higher than those for trees ($\bar{x} = 1880$ vs. 1413 m; $t = 2.41$, $P < 0.05$) but there was no difference between trees and shrubs (1730 m) or between shrubs and herbs (sample sizes for epiphytes precluded analysis here). There was no difference in the low/high sampling limits and extents among these three groups ($F = 0.03$, $P = 0.97$; $F = 0.03$, $P = 0.97$, and $F = 0.01$, $P = 0.99$, respectively). Also, there was no difference between ferns and seed plants, although the diversity peaks for ferns tended to be higher in elevation than trees ($t = -1.45$, $P = 0.08$) (Fig. 7).

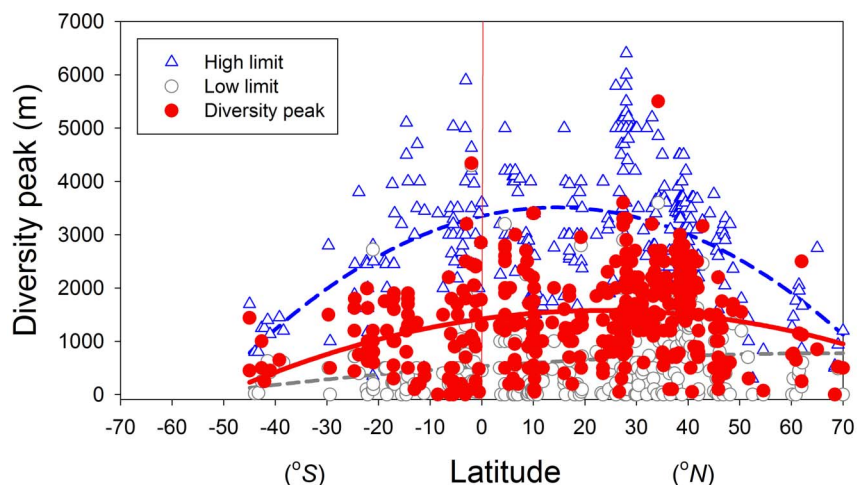


Figure 4 | The relationships among latitude, elevations of diversity peaks, and related sampling limits based on the 443 gradients. All relationships are significant ($p < 0.0001$) but the ones for Northern Hemisphere are better described by second-order regression and those for the Southern Hemisphere are described by a linear regression.

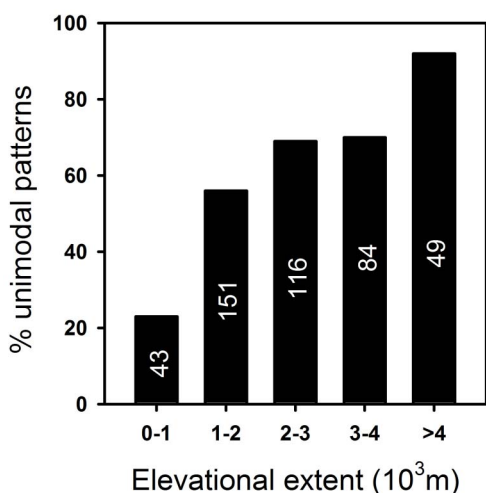


Figure 5 | Distribution of unimodal diversity gradients in relation to elevational extents measured from low to high sampling limits based on the 443 gradients in both Northern and Southern hemispheres. The number in each bar represents the no. of gradients in each category. Relationships examined separately for Northern and Southern hemispheres yield similar results (not shown).

Qualitative description of elevational diversity patterns is partly a function of the diversity of the groups assessed, and within our dataset the probability of a hump-shaped pattern increased with the richness of a taxonomic group, while patterns for constituent subgroups were highly variable. For example, the proportion of unimodal patterns across all seed plants was higher (70%) than for constituent groups (trees, shrubs, herbaceous species, or ferns alone; see corresponding values above). For all mammals, the proportion of ranges that were unimodal was $22/24 = 92\%$ but the average for immediate subgroups (e.g., bats, birds, rodents, and small mammals) was $34/63 = 54\%$. For insects, the proportion of unimodal patterns was $3/4 = 75\%$, but the average for its immediate subgroups (e.g., ants, bees, butterflies, beetles, termites) was only $26/47 = 55\%$. When these three major groups and their subgroups are pooled, the larger groups were more likely to display unimodal patterns than were their subgroups ($t = 3.33$, $P = 0.039$).

Fisher's exact test showed that the sampling using both mixed sources of data (type-D; see Methods) and regional floras/faunas

(type-C) revealed proportionally more unimodal elevational patterns than sampling using field transects (type-A; note that samples based on specimens, or type-B, was not included in this analysis due to small sample size, $n = 6$); but there was no difference between types C and D in this regard (Fig. S2). However, we failed to detect significant association between the proportions of type-C sampling and the unimodal pattern across latitudes (Northern Hemisphere: $F = 2.49$, $P = 0.19$; Southern Hemisphere: $F = 0.01$, $P = 0.92$). Similarly, we detected no difference in sampled taxonomic group size among the four sampling categories. For example, when we evaluated "all" plants, mammals, and insects, respectively, type-C sampling (based on regional floras/faunas) yielded very similar proportions of unimodal gradients (16/52 or 31%, 9/16 or 56%, 31/46 or 67%) to those based on type-A sampling (e.g., field transects; 50/185 or 27%, 23/38 or 61%, 26/43 or 61%).

Discussion

Elevational patterns differ both across hemispheres and across latitudes¹⁴. Many of the differences between the two hemispheres and even the locations of many global biodiversity hotspots likely reflect differences in landmass and mountain-range configuration, and relatively higher temperature and climatic stability in the Southern Hemisphere¹⁵. For example, the higher elevations of diversity peaks in the Northern Hemisphere may simply reflect the higher elevations of mountains and consequently a higher range of elevations sampled (Fig. 4). In addition, the latitudinal extent examined is shorter ($0-45^\circ$) in the Southern vs. Northern Hemisphere ($0-70^\circ$)¹⁶.

Interestingly, in both hemispheres, a greater proportion of unimodal patterns occurs at lower mid-latitudes (e.g., temperate zones; Fig. 3). Three factors might be responsible for this trend; (1) greater elevational extents (or gradients) associated with higher mountains in this latitudinal belt, (2) greater anthropogenic disturbance at lower elevations due to human activity¹⁷, and (3) shorter climatic gradients at higher latitude (cold temperate, boreal) zones (e.g., tree line occurs at lower elevations) that may limit elevational bulges in diversity. Further work to elucidate the underlying cause of this pattern would be worthwhile.

The elevational positions of the lower/upper sampling limits and extents are clearly linked to the observed diversity patterns and elevations of diversity peaks which generally are higher at mid-latitudes for major taxonomic groups (Figs. 4–6). This could reflect a tendency by researchers to study higher mountains with strong or more extensive elevational gradients and more vegetation zones for

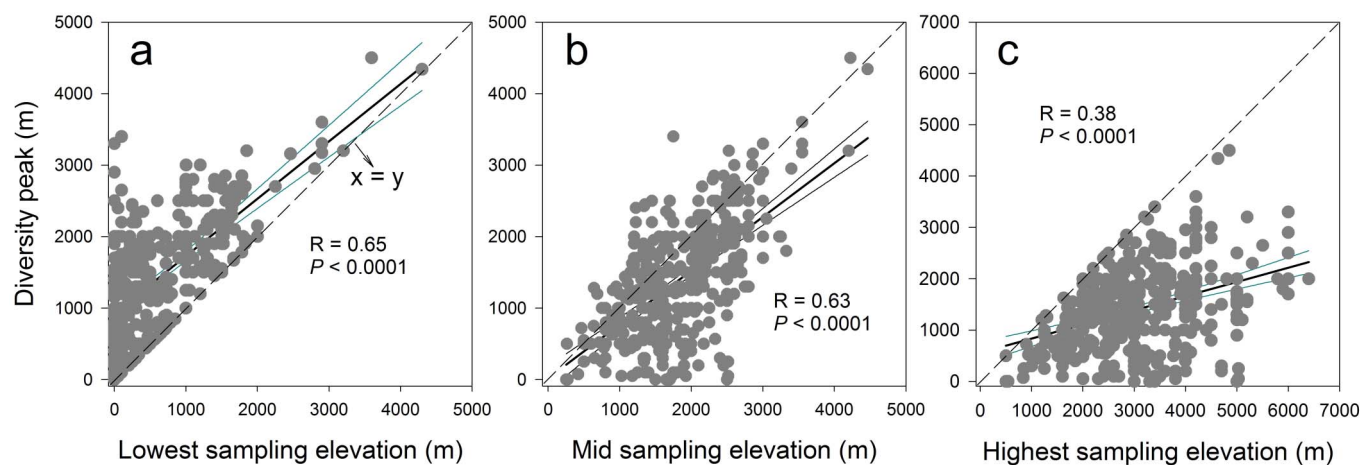


Figure 6 | Effects of the lowest, mid, and highest sampling limits (green) on the elevational positions of diversity peaks (linear regressions with 95 confidence intervals). Most of the diversity peaks are located lower than the mid-elevation of the sampled gradients (i.e., closer to the lowest limits than to the highest limits or $a < b$), leading to the positive skews in diversity curves. The lowest sampling limits show stronger relationships with the elevations of diversity peaks than highest limits ($R = 0.65$ vs. 0.38).

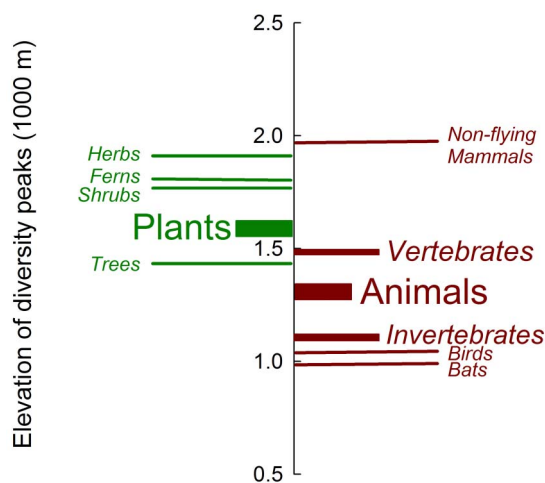


Figure 7 | Comparison of mean elevations of diversity peaks among hierarchical taxonomic groups.

examining diversity patterns. As such, many smaller (or lower) mountains that are not represented in Fig. 1 have yet to be investigated; however, this study underscores the fact that smaller ranges, even with more intensive sampling, may be more likely to exhibit non-unimodal patterns, likely reflecting the limited geographical range over which such patterns could be expressed.

The weaker correlation between the elevation of peak diversity and the upper sampling limit could be due to the difference between the highest elevation sampled and the position of treeline, snowline, or particularly vegetation line. Related to this, the unimodal distribution of diversity peaks across latitude (Fig. 4) may be an artifact of the fact that mid-latitude gradients have higher starting points; the 20–45°N latitude range includes the highest mountains on earth, but these also occur on the largest landmass and as such the elevational gradients begin at higher elevations than many that were studied at higher and lower latitudes. As noted above, monotonic elevation-diversity relationships may be observed if the elevational gradient is limited that it lacks sufficient variation in biotic conditions; this could occur even if the start position is low when working in high latitudes^{18–20} or if it were high in lower latitudes (e.g., the high base of the elevational gradient).

These results beg the question of what drives the commonly observed unimodal elevational diversity pattern? The interactive effects of temperature and precipitation have been widely invoked to explain this; however, while temperature generally declines with elevation, patterns of precipitation vary greatly depending on mountain height, aspect, and latitude⁶. This could in part explain why patterns across elevation do not often mirror the well-documented pattern across latitude (over which both temperature and precipitation generally are greatest in equatorial regions)¹¹. Reduced diversity at higher elevations is often related to decreasing temperature, area, soil nutrition, and increased isolation⁴ but other factors such as spatial constraints (e.g., the mid-domain effect)^{21,22} and human impacts^{23–26} also are important. It is beyond the scope of this work to specifically address the role of climate in structuring elevation diversity gradients, and sufficient data are lacking from transects (or even slopes with different aspect on the same transect) on the individual mountains where diversity has been surveyed. In any case, all these putative explanations fail to predict the right-skewed elevational pattern in diversity. One explanation that seems likely is that spatial constraints at higher elevations preclude accumulation of higher species richness¹, either through reduced diversification and immigration or through increased risk of extirpation/extinction.

Some deviations from the common unimodal pattern likely are a reflection of the particular taxa under consideration. For example,

within tropical mountain ranges, temperate taxa generally increase in diversity with elevation but then decline at the highest elevations, whereas tropical taxa generally decrease in diversity with elevation²¹ (unless the lower elevations are highly disturbed by human activities). Also, differences in elevational diversity peaks among plants and animals and among their subgroups (Fig. 7) probably reflect differences both in physiological tolerance and niche partitioning among species groups (e.g., trees vs. shrubs vs. herbs; bats vs. birds vs. non-flying mammals). In general, less inclusive groups (i.e., those with fewer species, such as genera vs. families, or families vs. orders) are less likely to present unimodal patterns because their constituent species usually have narrower distribution/elevational ranges and more specialized habitat preferences (e.g., total niche space of any given species is by definition less than that of all species within that particular genus). Therefore, we would expect smaller groups to demonstrate varied elevational patterns, as has been demonstrated across latitudinal gradients²⁷.

Research on elevational gradients has employed a diverse array of sampling methods (Fig. S2)^{10,28}, and we structured our analyses to provide some insight to this potential bias. Among our data, mixed sampling (type-D) produced a higher proportion of unimodal elevational pattern than the other three sampling types (i.e., field transects, specimens, and floras/faunas; see Methods), which we cautiously attribute to the more complete documentation of overall species distribution along the elevational gradients provided by combining different types of data. However, this observation was not a consequence of differential sampling across the taxonomic hierarchy (e.g., different sampling types were not preferentially used for more vs. less inclusive taxonomic groups), presumably because (1) when researchers examine elevational gradients using regional biotas (i.e., type-C), they also extract data from such databases to examine the elevational patterns of individual species groups, (2) a large proportion of studies employ multiple methods in generating the elevational patterns⁶, and (3) newer biotas are increasingly complete and comprehensive.

However, improved sampling (both in extent and intensity) over time may have contributed to the increase in the proportion of unimodal patterns. For example, less than a decade ago, 50% of 204 gradients reported by Rahbek¹⁰ displayed unimodal patterns, whereas our data suggests about 63%. This difference likely reflects greater sample size as well as more comprehensive and expansive sampling in recent years. It is also possible that earlier studies were based on lower sampling intensity and erroneously reported monotonic decline in diversity with elevation, leading to the perception that elevation patterns mirror latitudinal patterns^{11,29}.

To further improve sampling in the future, we suggest more efforts to compare aspects of the same mountain. Future comparisons among all 4 major aspects (*N*, *S*, *W*, *E*) but especially between the *N*- and *S*-slopes or between wet-(rain-receiving) and dry-slopes of the same mountains would, to some level, reduce the confounding effects of mountain height and latitude. It could also help separate the influences from one set of variables such as area, range/positions, and even the species pool that are relatively constant and that from other sources such as radiation, temperature, and moisture which may vary enormously.

A few related questions require further attention. For example, (1) why are diversity peaks generally located at lower mid-elevations rather than at the symmetrical midpoint as predicted by the mid-domain effect (Fig. 6)? (2) Why do we see qualitative differences among hierarchical taxonomic groups? (3) What factor(s) underlie different patterns on different aspects of the same mountain? (4) What evolutionary or historical processes produce observed elevational diversity patterns³⁰? (5) How might diversity peaks change with time³¹, and how will climate warming and other anthropogenic impacts (e.g., habitat loss and degradation) re-position the diversity peaks of various taxonomic groups? (6) How do exotic



species integrate into existing elevational patterns and what are the consequences for native species (e.g., insinuation vs. competitive replacement)? Answers to these questions will advance our understanding of global elevational biodiversity gradients and simultaneously inform pressing conservation efforts. For instance, if climate warming results in upward shifts of species ranges, will those species groups that exhibit diversity peaks at the highest elevation (Fig. 7) automatically experience the most immediate conservation threats, and if so, what actions could be taken to save these species?

Elevational diversity patterns are quite diverse and to some extent reflect the unique contemporary and historical contingencies of each range. The uneven distribution of mountains and of their elevational extents across the planet clearly influences latitudinal diversity patterns but this role appears to have been underappreciated. Quantifying this influence is likely to help interpreting deviations from the conventional latitudinal patterns. With more and better data available, global elevational patterns should become increasingly clear, enabling us to better understand causal mechanisms. Distinguishing between scale and gradients, among taxonomic groups, and implicitly incorporating the importance of human and climate change in future comparative studies will greatly benefit our understanding of basic ecological phenomena and how research findings may be appropriately applied in conservation.

Methods

We compiled and examined 443 elevational gradients from the literature on vertical diversity patterns from around the world (see Supplementary Information for a complete reference list). We searched Google Scholar using key words such as elevation(al), altitude/altitudinal, diversity, and richness up to Oct. 31, 2012 when final analyses were started. We selected those studies that provided (1) the location where sampling/survey was conducted, (2) latitude/longitude (when not provided, we extracted this using the latitudinal locations of the mountain peaks), (3) the organism types, (4) lower/upper sampling endpoints (thus extents), (5) diversity or richness measures, and (6) the position of diversity/richness peaks along the transects with relatively high sampling intensity. We selected those with full access to contents in peer-reviewed journals. Additionally, we searched and added the literature/data in reverse chronological order, preferentially selecting more recent studies because they generally were conducted with more complete and more intensive sampling.

The 443 gradients/transects all fell within 0–70° latitude in both Northern ($N = 337$) and Southern hemispheres ($N = 106$; Fig. 1). These studies involved diverse taxonomic groups of plants ($N = 249$), animals ($N = 192$), and microorganisms ($N = 2$) with various sizes and heterogeneities in terms of the number of species and composition. The studies on plants included those on seed plants, pteridophytes, epiphytes, and bryophytes. The studies on animals were more diverse, including all non-flying mammals, small mammals, bats, birds, insects, ants, lizards, frog, snakes, and snails. The studies on microorganisms were mostly based on soil samples. Our dataset is available upon request to the senior author and we continue to expand this to include other related aspects such as life history features, mountain-specific weather/climate data for future comparisons.

To examine how elevation diversity patterns change with latitude, we segregated studies by 10° latitudinal zones (e.g., 0–10°, 10–20°, 20–30°, 30–40°, 40–50°, >50°) to ensure sufficient sample sizes for comparative analyses. The actual lower and upper sampling limits were used as surrogates to represent the length and start/end positions of underlying gradients. In many cases, the sampling limits did not represent the actual bases or tops because different mountain aspects had different bases and sometimes treeline or vegetation line limited sampling frames. Elevational diversity patterns were classified by the relationship between elevation and taxonomic richness; as such, classifications included positive (richness increases with elevation), negative, unimodal, bimodal or polymodal (“bimodal/polymodal”), unimodal or negative (“unimodal/negative”), and “none”. The category “unimodal/negative” included gradients that showed both patterns as well as those that could be classified as either unimodal or negative depending on the statistical techniques used (i.e., first vs. second order regression³²) or individual discretion (especially when the diversity peak is very close to the lower limit of sampling). “None” included all gradients that were either very difficult to describe (and classify). Pattern classifications were based on the original authors’ descriptions with further confirmation during this study. For some gradient patterns (3 of 6), we adopted a procedure similar to that of Rahbek¹⁰ except that the patterns of “horizontal” and “other” were replaced by “bimodal or polymodal”, “unimodal or negative”, or “none”. Because the majority of these studies did not provide specific information regarding key features of the gradients sampled, we did not attempt to evaluate the influence of aspect on diversity patterns.

To examine whether sampling techniques may have biased results, we classified studies (gradients) into four categories as follows: field transects (type-A; $n = 291$), specimen-based (type-B; $n = 6$), regional floras/faunas (type-C; $n = 98$), and mixed sources (type-D; $n = 48$). The number of type-B transects was limited primarily because this method had mostly been used in combination with other methods such

as field surveys⁶ and/or local/regional floras or faunas¹³, and as such were classified as type-A, -C, or -D transects (see Fig. S2 and additional references in Supplementary Information).

While we are unable to categorically exclude the possibility that differential sampling intensity may influence our results, we believe that our approach makes this unlikely. In particular, we have selected recent studies which tend to sample both more intensively and more extensively, we grouped gradients into 10° latitudinal belts to minimize local influences, and we have compared results based on four sampling techniques (which are known to have different sampling biases). As we note in Discussion, further research in this field should emphasize standardized sampling across elevations, latitudes, and aspects to most effectively promote our understanding of geographical patterns in vertical gradients.

We used regression analyses, *t*-tests, contingency tests (Chi-square and Fisher’s exact test), and one-way ANOVA for comparative analysis among latitudinal zones and different elevational extents used in the case studies and linear regressions to examine the correlations of the elevations of diversity peaks with the elevations of sampling limits (lower, mid, and upper) and latitude. For comparative purposes, most analyses were performed separately for Northern and Southern Hemispheres.

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Author contributions

Q.G. initiated and designed the research. Q.G., Z.S., H.L., L.H., H.R. and J.W. collected data. L.H. and H.R. also revised and edited the manuscript. Q.G. and D.K. performed data analyses and wrote the paper.

Additional information

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