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Title: The latitudinal diversity gradient of tetrapods across the Permo-Triassic mass extinction and recovery interval

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Abstract:

The decline in species richness from the equator to the poles is referred to as the latitudinal diversity gradient (LDG). Higher equatorial diversity has been recognised for over 200 years, but the consistency of this pattern in deep time remains uncertain. Examination of spatial biodiversity patterns in the past across different global climate regimes and continental configurations can reveal how LDGs have varied over Earth history and potentially differentiate between suggested causal mechanisms. The Late Permian-Middle Triassic represents an ideal time interval for study, because it is characterised by large-scale volcanic episodes, extreme greenhouse temperatures, and the most severe mass extinction event in Earth history. We examined terrestrial and marine tetrapod spatial biodiversity patterns using a database of global tetrapod occurrences. Terrestrial tetrapods exhibit a bimodal richness distribution throughout the Late Permian-Middle Triassic, with peaks in the northern low latitudes and southern mid latitudes around 20-40% and 60%, respectively. Marine reptile fossils are known almost exclusively from the Northern Hemisphere in the Early and Middle Triassic, with highest diversity around 20%. Reconstructed LDGs contrast strongly with the generally unimodal terrestrial gradients of today, potentially reflecting high global temperatures and prevailing Pangean super-monsoonal climate system during the Permo-Triassic.

Keywords: climate change, greenhouse, biodiversity, sampling bias, Tetrapoda, mass extinction

1 Background

2 The latitudinal diversity gradient (LDG) is one of the largest-scale and longest-known patterns in ecology [e.g.1-7]. Modern biodiversity (i.e., richness) gradients are broadly 3 4 defined as unimodal, with high biodiversity near the equator and low biodiversity at the poles 5 [4,6]. The specifics of these gradients differ among taxonomic and ecological groups [2-4,6], and research over the last decade has revealed greater variation in modern LDGs than 6 7 previously recognised. For example, benthic marine species richness appears to peak at 10-8 20°N, whereas pelagic species richness appears bimodal, with peaks at 10-40° on either side of the equator [8-11]. 9

10 Study of the fossil record suggests the shape of LDGs has also changed through time 11 [5,12]. Dinosaur diversity may have been greatest at temperate latitudes throughout the 12 Mesozoic [13], North American mammal diversity may have been distributed evenly across 13 latitudes for much of the Cenozoic [14,15], and peaks in marine animal diversity may have 14 drifted from the Southern to Northern Hemisphere over the course of the Phanerozoic 15 [16,17].

Numerous drivers of LDGs have been proposed [e.g.2,3,18-22]. Interaction between key processes, the complexity of feedback cycles, and the covariance of many environmental variables with latitude complicate efforts to isolate causal mechanisms [2-4,6,23]. Climate and landmass distribution, however, have been put forward consistently as potential explanatory variables. Climate, particularly temperature and water availability, has long been considered a key control on spatial patterns of terrestrial biodiversity [5,12,14,15,17,24-26] because the distributions of species are limited by climatic tolerance [22,24,27-30].

23 LDG studies in deep time have suggested palaeoclimate regime is a major control on the 24 shape and slope of LDGs. Icehouse periods have been associated with a sharp, unimodal 25 equatorial richness peak, and greenhouse periods with a bimodal distribution, characterised 26 by shallow peaks at mid-latitudes of both the Northern and Southern Hemispheres [5,17,31]. 27 This contrast has been attributed to the strength of the pole-to-equator temperature gradient between the two climate regimes [5,13,17], but may also reflect spatio-temporal variations in 28 precipitation [14,22,25]. The distribution of continental landmasses may also structure global 29 30 patterns of biodiversity by controlling the spatial distribution of relevant habitats, particularly shallow continental shelf area in marine ecosystems [32-35]. Although studies of modern 31 LDGs continue to provide insight into potential generative mechanisms [e.g.36-39], 32 33 examination of LDGs during intervals when climate and landmass distribution were considerably different to today may provide new perspectives on global biodiversity patterns 34 and their associated processes [5,13,31]. 35

36 The Late Permian to Middle Triassic (~260-239 Ma) represents a period in Earth history 37 that contrasts considerably with the present day. Large-scale volcanism associated with the 38 Siberian Traps drove extreme climate change, which was exacerbated by amalgamation of most landmasses into the supercontinent Pangea [40-42]. This drove environmental 39 40 perturbations that resulted in the most catastrophic mass extinction event of all time at the end of the Permian, around 252 Ma [43]. A prolonged interval of extremely high 41 temperatures, which peaked in the Olenekian (late Early Triassic) [44], along with ocean 42 anoxia and acidification have been identified as key extinction mechanisms [43,45], with 43 44 persistence of these conditions extending well into the Middle Triassic, delaying full 45 structural recovery of marine ecosystems for as long as 50 million years [46]. On land, high temperatures and seasonal precipitation in central Pangaea resulted in drought [40,47-49], 46 while purported ozone depletion, brought about by halogen gas release from the Siberian 47 48 Traps, resulted in high UV-B levels that caused plant sterilisation and extinction [e.g.50,51]. Early Triassic temperatures at low latitudes are considered to have been beyond the 49 tolerable long-term threshold for both plants and animals, driving extinction and poleward 50 51 migration [44,52]. The climate of the Middle Triassic has received less attention, but is 52 thought to have been characterised by continued aridity in lower latitudes, with cyclical 53 temperature fluctuations overprinting a general trend of steady cooling after the final 54 eruptions of the Siberian Traps in the Olenekian [41,42].

55 Tetrapods were profoundly affected by the Permo-Triassic mass extinction (PTME). In 56 the immediate aftermath, Early Triassic tetrapod communities were composed almost 57 entirely of 'disaster fauna' such as Lystrosaurus, a herbivorous burrowing synapsid [47,53-58 55]. Following recovery from the PTME, archosauromorphs (Sauria), a group that appeared 59 in the Middle Permian, became the dominant terrestrial animals [55]. The first marine reptile 60 fossils are known from the Olenekian and were highly diverse by the Anisian, including basal 51 sauropterygians and ichthyosaurs [56].

62 Two previous studies have offered perspectives on the distribution of tetrapods across the Permo-Triassic boundary. Sun et al. [44] used oxygen isotopes in conodont apatite to 63 64 examine sea surface temperature (SST) change across the Late Permian and Early Triassic, recovering remarkably high SSTs throughout the interval but particularly during the 65 Smithian-Spathian Thermal Maximum (~248 Ma), when equatorial SSTs may have 66 67 approached 40°C. Their qualitative analysis of tetrapod occurrences revealed an equator ial 68 'tetrapod gap' in the Early Triassic, hypothesised to have occurred due to the extreme warm 69 temperatures that may have approached or exceeded the thermal tolerances of both 70 terrestrial and marine vertebrates (around 42°C [57]). Bernardi et al. [52] also exa mined the 71 distribution of individual tetrapod skeletal and footprint occurrences through the extinction

and recovery interval, finding evidence for a poleward shift in tetrapod abundance in the
Northern Hemisphere, but only in the Induan (earliest Triassic). This biogeographic pattern is
congruent with a study of tetrapods immediately prior to the PTME, which found higher
tetrapod diversity at temperate than equatorial latitudes during the Middle – Late Permian
[58].

Here, we explore further the terrestrial and marine Permo-Triassic fossil tetrapod record by comparing species-level tetrapod biodiversity across latitudinal bins. We apply coveragebased interpolation and squares extrapolation to reconstruct LDGs from the Late Permian (before the PTME), Early Triassic (in the aftermath of the PTME) and Middle Triassic (during recovery). These LDGs are then assessed in light of the hypothesis that higher diversity will be found in the cooler refugia of the mid to high latitudes during extreme greenhouse conditions, such as during the Late Permian to Middle Triassic.

84

85 Methods

86 We conducted an in-depth literature review to maximise the completeness and robustness of our Late Permian to Late Triassic dataset for tetrapods. All tetrapod fossils from the 87 Wuchiapingian (early Late Permian) through Carnian (early Late Triassic) were downloaded 88 from the Paleobiology Database (PBDB). Genus names from this download were used to 89 90 conduct a systematic literature search in Google Scholar, and any new taxa and occurrences were added to the PBDB. Once completed, the same criteria were used to 91 92 download the enlarged dataset (in October 2018) [59]. We reviewed each 'collection', 93 representing fossils from a particular locality and considered to be of a similar age, to 94 increase temporal resolution. A literature search for formation names was conducted, with 95 publications that listed the ages of specific beds or members further refining the geological 96 date of collections, where possible [59]. We streamlined the mode of preservation and taxon 97 habitat categories, reduced to either 'body' or 'trace', and 'marine' or 'terrestrial', respectively. Finally, the modern latitude and longitude of fossil localities were rotated to 98 their palaeo-position at the time of deposition by filtering occurrences on a stage-by-stage 99 basis then using the PALEOMAP Global Plate Model [60], implemented in G-Plates (version 100 101 2.1.0) [61]. The final dataset constituted 3563 unique tetrapod occurrences assigned to stage level, with our search efforts contributing 490 of these occurrences (13.8%). 102

All subsequent data manipulation and plotting was carried out in R [62] utilising the
 'tidyverse' suite of packages [63]. The final dataset was filtered to include only records
 representing unique species, comprising those identified to species level and those identified
 to coarser taxonomic levels but representing a clade not already within their spatiotemporal

bin. Since abundance data in the Paleobiology Database are relatively incomplete and
 inconsistently applied, the presence of a species within any given collection was treated as a
 single occurrence. Fossil occurrences dated to a single geological stage were used to
 produce raw sampled-in-bin tetrapod richness curves.

To compare tetrapod richness patterns across space between the Late Permian, Early Triassic, and Middle Triassic, stage-level occurrences were binned using 20° latitudinal bands, from 90°S to 90°N (the central bin includes the equator, from 10°N t o 10°S). Terrestrial and marine body fossils were analysed separately, with 'marine tetrapods' referring to species whose morphology indicates life in marine habitats. This informal group is polyphyletic and includes basal ichthyosaurs, sauropterygians, tanystropheids and thalattosaurs.

118 When reconstructing historical spatial biodiversity patterns, allowances must be made for 119 the spatiotemporal unevenness of the fossil record [5,56,64-70]. LDGs in deep time can be 120 estimated if sampling rates in the clade of interest are relatively high, and consideration is 121 given to partitioning variation in richness likely attributable to sampling biases versus that 122 likely attributable to biological patterns [71]. Furthermore, subsampling and extrapolation 123 methods can help alleviate issues of sampling heterogeneity. Coverage-based approaches are currently the most effective approach for mitigating the effects of fossil record bias in 124 large-scale biodiversity analyses [72,73]. 125

We applied two analytical approaches to account for spatiotemporal sampling biases in occurrence data: coverage-based interpolation [74,75] and squares [76]. Both were applied to collections within latitudinal bins for the Late Permian, Early Triassic, and Middle Triassic time intervals (analyses were repeated for individual stages, Figure S2). Only body fossils were used for these analyses, due to the biological non-equivalence of trace fossil and body fossil species; one animal can produce multiple trace fossils, and traces are not easily allied to individual body fossil species.

Richness estimates were generated using coverage-based interpolation following the 133 134 approach of Dunne et al. [77] using the R package iNEXT [75]. This approach conducts coverage-based rarefaction using the equations of Chao & Jost [74] (analogous to 135 shareholder quorum subsampling [64,72]) and extrapolation based on the Chao1 estimator. 136 Extrapolated estimates were discarded if more than three times the observed sample size, 137 138 as this suggests a high species-to-occurrence count ratio that indicates the bin under consideration is likely to be undersampled [75]. Bins containing fewer than three species 139 140 were incompatible with subsampling and therefore excluded from analyses (Table S1). Coverage-based rarefaction curves are also presented (Figure S3) to illustrate the 141

relationship between coverage and coverage-standardised diversity estimates in each bin[72,77].

In addition to coverage-based interpolation, richness estimates were generated using
the squares method [76]. Squares is an extrapolater based on the proportion of singletons in
a given sample, and is considered more robust to biases arising from small sample sizes
and uneven abundance distributions than other interpolation methods [73,76]. Squares
richness estimates were produced using the equation stated by Alroy [76].

Finally, we tested whether variation in sampling intensity among time bins influenced 149 richness estimates, particularly given the expected reduction in Early Triassic tetrapod 150 occurrences following the PTME. We subsampled to the same number of collections in each 151 time interval (Late Permian, Early Triassic and Middle Triassic) using a bootstrap routine. 152 153 For each time bin, we randomly sampled 250 collections for terrestrial tetrapods and 30 154 collections for marine tetrapods. Collections were allocated to their corresponding latitudinal 155 bin and species richness was quantified across collections within each bin. This process was 156 repeated 100 times. Diversity curves were produced using the mean species diversity in 157 each latitude bin across the 100 replicates, allowing for comparison of LDGs among time 158 bins given an artificially-fixed sampling intensity.



159 160 **Figure 1**

161 Tetrapod diversity by latitude in the Late Permian, Early Triassic and Middle Triassic. The 162 grey bars indicate 30-60°N and S.

a. Palaeo-rotated occurrence locations plotted over maps from Scotese [59]; maps representthe Lopingian, Induan-Olenekian, and Ladinian.

- b. Raw occurrences within 20° latitude bins (e.g. central bin is 10°N-10°S).
- 166 c. Squares diversity by latitudinal bin for terrestrial (green) and marine (blue) tetrapods.
- d. Interpolated diversity by latitudinal bin for terrestrial tetrapods. Bins with < 3 species have
- 168 been plotted as '0', while missing points indicate an estimated diversity of more than three
- times the observed value. Error bars indicate 95% confidence intervals.
- e. Interpolated diversity by latitudinal bin for marine tetrapods. Bins with < 3 species have
- been plotted as '0', while missing points indicate an estimated diversity of more than three
- 172 times the observed value. Error bars indicate 95% confidence intervals. The oldest marine
- tetrapod fossils are Olenekian (late Early Triassic; 251-247Ma) in age.

174 Results

175 Sampling

176 Raw richness, squares and interpolation estimates produced similar diversity-through-time

177 curves (Figure S1). The number of collections with terrestrial tetrapod body fossils was

relatively consistent through time (Late Permian, 291; Early Triassic, 307; Middle Triassic,

179 354), while the number of collections containing marine tetrapods increased from the Early

to Middle Triassic (Early Triassic, 32; Middle Triassic, 207). Curves of raw species richness

181 by latitude bin produced by bootstrapping to the same number of collections for each time

interval were near-identical to those using the full datatset (Figure S4).

183 Terrestrial distribution

184 Terrestrial tetrapod occurrences were broadly distributed but clustered throughout the 185 studied interval (Figure 1a). Both squares and interpolation analyses of terrestrial tetrapods by latitude (Figure 1c) show a consistent bimodal richness distribution throughout the Late 186 Permian to Middle Triassic, with a persistent dip in diversity in the low southern latitudes. In 187 188 the Northern Hemisphere, diversity peaked at 40% in the Late Permian. By the Early Triassic, the peak in species diversity had shifted to the 20[°]N bin (Figure 1b), with stage-189 level analyses indicating this occurred in the Olenekian (Figure S2b). In the Middle Triassic, 190 the Northern Hemisphere peak returned to 40%. The gradient in the Southern Hemisphere 191 192 remained relatively unchanged throughout the Late Permian to Middle Triassic, characterised by a consistent 60°S diversity peak. 193

194 Marine distribution

195 Marine tetrapods were generally restricted to the Northern Hemisphere during the Early 196 and Middle Triassic, despite having a relatively broad longitudinal distribution (Figure 1a). Early Triassic marine tetrapods were most diverse in the 20% bin, with the only other 197 occurrences found in the 40% bin (Figure 1d). The 20% peak in biodiversity persisted into 198 199 the Middle Triassic, but with new occupation of the equatorial and 20°S bins. The stage-level analyses generally show comparable trends to those seen in the epoch-level time bins, but 200 often with fewer bins occupied, producing patchier and less constrained gradients (Figure 201 202 S2).

203 Comparison with modern LDGs

The Early Triassic terrestrial LDG produced by interpolation was compared to LDGs of modern birds, mammals and amphibians (Figure 2; modern data derived from

BiodiversityMapping.org, as used by Saupe et al. [22]). The modern curves have unimodal

- distributions that peak at low latitudes (maximum diversity at 9.5°S for birds and amphibians,
- 208 2.5[°]N for mammals), whereas the Early Triassic terrestrial curve peaks at higher latitudes,
- with a clear bimodal distribution (maximum diversity at 32.5°N and 62.5°S).



211 Figure 2

210

- 212 Smoothed latitudinal gradients for species of modern birds (a), mammals (b), and
- amphibians (c), compared with Early Triassic terrestrial tetrapods (as an example) based on
- 214 interpolation analyses (d). Modern gradients derive from data obtained from
- 215 BiodiversityMapping.org. Silhouettes are from Phylopic.org.

216 Discussion

217 In contrast to gradients for modern terrestrial tetrapods, the Permo-Triassic terrestrial tetrapod gradient was likely bimodal with reduced diversity at low latitudes (10°N-30°S) 218 219 (Figure 1) [4,6]. The general shape of the terrestrial tetrapod richness gradient, particularly 220 its bimodality, remained relatively constant throughout the Late Permian to Middle Triassic, 221 and may reflect the prevailing climate regime (greenhouse versus icehouse) [5,17,31]. Interestingly, the shape of the gradient did not seem affected by the PTME or even higher 222 223 temperatures of the Early Triassic (equatorial SSTs increased from ~24°C in the latest Permian to $\sim 40^{\circ}$ during the late Smithian [44]). Marine tetrapods, by contrast, maintained a 224 diversity peak at low latitudes in the Northern Hemisphere from the Early to Middle Triassic 225 226 (Figure 1). The bimodal terrestrial LDG obtained here is comparable to the distribution of raw Early Triassic tetrapod occurrences from Sun et al. [44] and Bernardi et al. [52], and 227 suggests continuity of LDG shape from the Middle Permian [58]. The shape of the gradient is 228 also broadly comparable to the gradient of Mesozoic dinosaurs, which Mannion et al. [13] 229 230 attributed to the distribution of land area during the break-up of Pangaea. This congruence 231 suggests LDGs may have been bimodal for much of the Permian to mid Cenozoic, with modern LDGs only developing as global climate gradually cooled through the late Paleogene 232 233 and early Neogene [5,14,15,31].

Although latitude is a reasonable proxy for temperature in the modern, this 234 235 relationship does not hold for the Triassic [41]. The latitudinal temperature gradient today largely reflects the operation of Hadley cells, but these cells may have collapsed in the Late 236 Permian to give way to a more zonally asymmetric atmospheric system, with strong 237 238 seasonal variation in temperature and precipitation [40,41,49]. Although the Tethyan coastal regions experienced supermoonsoons, considerably less precipitation reached the 239 continental interior, resulting in high aridity, particularly in the southern low to mid latitudes 240 [40,43]. Climate model reconstructions for the latest Permian suggest large areas of central 241 Pangea were desert, with seasonal average temperatures up to 45° in the arid subtropics 242 at 20-25[®]N and S [78]. Late Permian palaeoenvironmental evidence from localities in South 243 244 Africa indicates considerable drought even at relatively high latitudes (~ 65 %) [47]. Thus, much of the supercontinent interior may have been uninhabitable in the Late Permian, which 245 246 could explain the bimodal, asymmetric tetrapod LDG reconstructed here. However, in 247 contrast to Permian climates, Triassic climates have not been well studied [41,42], and the 248 development of high-resolution climate models for the Triassic is essential for determining the key drivers of tetrapod extinction and migration during this interval. 249

250 We cautiously interpret the bimodal richness distributions found here as biologically 251 meaningful, particularly given the agreement between the different sampling methodologies 252 employed. In addition, collections from southern low latitude regions are consistently of low alpha diversity throughout the entire Late Permian to Middle Triassic, in comparison with 253 254 some very high levels of alpha diversity in mid-latitude collections during the same intervals (Table S2). However, the spatial and temporal resolution of the analyses, and our certainty 255 in the observed distributions representing biological patterns, would benefit from better 256 geographic spread and higher density of samples [67-69]. New discoveries from the 257 258 southern low to mid latitudes could particularly help to distinguish between low biodiversity 259 and poor sampling, but fossiliferous outcrops of this age and palaeolatitude are uncommon, 260 particularly from terrestrial environments (Figure S5) [41,49,67]. Although extensive shallow and marginal marine deposits, such as those in Oman, are rich in invertebrate fossils 261 [e.g.79], vertebrate fossils are known only from a handful of localities, such as Gour Laoud in 262 Algeria (Jesairosaurus lehmani, Odenwaldia sp., indeterminate amphibians; palaeolatitude 263 9°S [80]) and Mariakani in Kenya (Kenyasaurus mariakaniensis; palaeolatitude 42°S [81]). 264 265 Unfortunately, the age of fossils from these localities is poorly constrained and were 266 therefore not included in our analyses.

Although broad stasis in bimodal richness gradients was observed over the ~23 267 million year interval considered here (Late Permian – Middle Triassic), smaller-scale 268 variability can be detected among time bins. Both squares and interpolation analyses 269 270 suggest a shift in peak diversity in the Northern Hemisphere towards the equator in the Early 271 Triassic, before returning to mid-latitudes in the Middle Triassic. This shift is also supported by the relatively high number of trace fossil occurrences in the equatorial and 20% bins 272 during the Early Triassic (Figure 1b). An Early Triassic equatorward shift in diversity in the 273 274 Northern Hemisphere seems surprising given that global temperatures were increasing at the time [44]. Instead, this shift may reflect differential sampling bias. Most of the 275 276 interpolation rarefaction curves are exponential in shape, but the Early Triassic 20% bin has 277 a more asymptotic curve (Figure S3), indicating sampling completeness may be substantially 278 higher in this bin relative to the others, inflating diversity estimates [72]. This peak in diversity 279 corresponds to the high density of tetrapod fossils known from the Olenekian of Eastern 280 Europe [82].

281 Conclusions

282 Our results suggest terrestrial tetrapods exhibited a bimodal richness distribution and 283 were most diverse at mid-latitudes during the Late Permian – Middle Triassic. In contrast, 284 marine tetrapods were generally restricted to northern low latitudes in the Early and Middle 285 Triassic. Tetrapods were not excluded from equatorial regions during this interval, but were reduced in diversity at low southern latitudes. The bimodal LDG for terrestrial tetrapods 286 during the Late Permian – Middle Triassic contrasts with the unimodal, equatorial diversity 287 288 peaks exhibited by most terrestrial tetrapod clades in the modern, including birds, mammals and amphibians (Figure 2). Permo-Triassic LDGs were likely shaped by the extreme climatic 289 conditions of the time, particularly high global temperatures and heterogeneous precipitation. 290 291 As is often the case regarding the vertebrate fossil record, our data are insufficient to determine conclusively whether observed patterns predominantly reflect true biological 292 signal or heterogeneous spatial sampling. Further examination of Triassic climates and 293 294 increased sampling intensity may advance our understanding of this time interval, providing 295 greater insight into the potential influence of extreme greenhouse conditions on global 296 patterns of biodiversity.

297

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308

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No ethical considerations were required for this study.

317

318 Data, code and materials

- 319 The datasets supporting this article have been uploaded as part of the supplementary
- 320 materials.
- 321

322 Competing interests

323 We have no competing interests to declare.

324

325 Authors' contributions

- 326 BJA downloaded and reviewed dataset, contributed additional Paleobiology Database
- 327 entries, conducted statistical analyses and drafted the manuscript. All authors contributed to
- 328 data interpretation and editing the manuscript. All authors also gave final approval for
- 329 publication and agree to be held accountable for the work performed therein.

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