

1 Changes in the Latitudinal Diversity Gradient during the  
2 Great Ordovician Biodiversification Event

3 **Björn Kröger\***

4 *Finnish Museum of Natural History, PO Box 44, 00014 Helsinki, Finland*

5 \*E-mail: bjorn.kroger@helsinki.fi

6 **ABSTRACT**

7 Near equatorial peak diversities are a prominent first-order feature of today's  
8 latitudinal diversity gradient (LDG), but were not a persistent pattern throughout  
9 geological time. In an analysis of Ordovician (485–445 Ma) fossil occurrences, an  
10 equator-ward shift of the latitudinal diversity peak can be detected. A modern-type LDG  
11 and out-of-the-tropics range shift pattern were synchronously established during  
12 emerging icehouse conditions at the climax of the Great Ordovician Biodiversity Event  
13 (GOBE). The changes in the LDG pattern and range shift trends can be best explained as  
14 a consequence of global cooling during the Middle Ordovician and of diversification in  
15 the tropical realm following a greenhouse period with temperatures too hot to support  
16 diverse tropical marine life. These results substantiate a fundamental role of temperature  
17 changes in establishing global first order diversity patterns.

18 **INTRODUCTION**

19 Life on earth today is most diverse at or near the equator and least diverse near the  
20 poles (Pianka, 1966; Currie, 1991; Hillebrand, 2004; Mittelbach et al., 2007; Jablonski et  
21 al. 2016). This latitudinal diversity gradient (LDG) is apparent in most groups of marine  
22 and terrestrial organisms, but was not a persistent pattern throughout geological time

23 (e.g., Mannion et al., 2014). Strictly tropical peak diversities might be a feature of  
24 icehouse climate regimes, whereas warmer greenhouse regimes show generally shallow  
25 LDGs (e.g., Crame, 2002; Marcot et al., 2016; Jablonski et al., 2016)

26         These expectations can be tested against the fossil record from a time period  
27 during which one of the most dramatic and sustained Phanerozoic diversifications, the  
28 Great Ordovician Biodiversification Event (GOBE), took place. The GOBE was a ca. 40  
29 Ma diversification interval during the Ordovician Period (485–444 m.y. ago) with a  
30 massive expansion in diversity at lower taxonomic ranks (species, genus, family level)  
31 (Fig. 1) and profound paleoecological changes within the marine biota (Servais et al.,  
32 2010). The diversification occurred against a background of changes in the  
33 physicochemical environment. In particular, it has been suggested that global cooling was  
34 one of the major driving factors, because this brought temperatures to levels more  
35 compatible with metazoan metabolism and skeletal growth (e.g., Trotter et al., 2008;  
36 Rasmussen et al., 2016). Hence, before cooling, large parts of the Ordovician seas,  
37 especially in the lower latitudes, would have been too hot to sustain highly diverse  
38 metazoan communities such as metazoan reefs (Webby, 2002). The GOBE, in this  
39 scenario, would represent a predominantly tropical diversification.

40         Ordovician diversity dynamics were previously analyzed mainly on a local,  
41 paleocontinent, or global scale (see e.g., Webby et al. 2004; Servais and Harper, 2013;  
42 Trubovitz and Stigall, 2016) and changes in LDG patterns can only indirectly be inferred  
43 from these data. Here, for the first time Ordovician marine diversity is analyzed across  
44 paleolatitudinal zones based on Cambrian–Silurian fossil occurrences documented in the  
45 Paleobiology Database (PaleoBioDB, <https://paleobiodb.org>). I reconstruct Ordovician

46 LDGs and their changes and ask whether the observed trends can be explained by  
47 changes in global temperatures, paleogeographical shifts, or changes in benthic habitats.

## 48 **METHODS**

49 The analysis was based on a download of the complete set of global, genus-level,  
50 Ordovician fossil occurrences from the Paleobiology Database on 26 November 2016.  
51 All accepted genus name entries in the PaleoBioDB were included in the analysis,  
52 without any additional filters set. Additionally, the complete set of 4081 genera from 773  
53 fossil collections from the Cambrian Trempealeau Regional Stage and the Silurian  
54 Rhuddanian Stage were downloaded. The stratigraphic data associated with each of the  
55 fossil occurrences were matched with stage level binned stratigraphic units in the  
56 RNames Database (<http://rnames.luomus.fi/>), resulting in 90,471 latest Cambrian to  
57 Rhuddanian occurrences (for time binning see Kröger and Lintulaakso, 2017). The genus  
58 occurrences are analyzed at stage- and stage-slice-level (Bergström et al., 2009) time  
59 resolution (average length 6.2 m.y. and 2.2 m.y., respectively).

60 PaleoBioDB collections are assigned to lithological and environmental classes.  
61 Based on these data I further classified all collections with sufficient lithology and  
62 environmental information into carbonate/siliciclastic and shallow/deep depositional  
63 environments (total = 13085 and 5739, respectively) and calculated their relative  
64 abundance (see the GSA Data Repository<sup>1</sup>). Additionally, I used the PaleoBioDB data on  
65 lithology and environment for calculation of environmental heterogeneity using the HRel  
66 statistics of Wilcox (1973), (see the Data Repository).

67 All PaleoBioDB collections are explicitly attributed to paleoplates and are linked  
68 to paleogeographic latitudinal/longitudinal coordinates based on GPlates

69 (<https://www.gplates.org/>) rotations (Wright et al., 2013). The collections were binned  
70 into 15° and 30° paleolatitudinal zones without distinguishing between northern and  
71 southern paleo-hemispheres, because in most cases sample size for the northern paleo-  
72 hemisphere was not sufficient for analysis.

73 I estimated genus range shifts across paleolatitudinal zones and distinguished  
74 between “active” and “tectonic” shifts. The observed shift of a genus from one  
75 paleolatitudinal zone into another can be either a result of a tectonic shift of the genus  
76 location through time or can be caused by a active migrations or a range expansion of the  
77 genus, or both. Active shifts can be neutralized or reinforced by tectonic shifts of  
78 occurrences.

79 For distinction between tectonic and active range shift, I binned all genus  
80 occurrences into one-degree latitude/longitude quadrants and tracked their  
81 paleogeographical shift. Occurrences with identical quadrants in one stage ( $t_i$ ) and its  
82 previous stage ( $t_{i-1}$ ) and with different paleolatitudinal zones are interpreted as  
83 tectonically shifted. Occurrences with different latitude/longitude quadrants and different  
84 paleolatitudinal zones are interpreted as active range shifts.

85 Tectonic and active range shifts from one paleolatitudinal zone to another for  
86 times  $t_{i-1}$  to  $t_i$ , were counted with a unit length of  $\pm 1$  depending on the direction, such that  
87 positive values indicate pole-ward directions and negative values indicate equator-ward  
88 directions. In order to eliminate boundary effects, I compared the distance from the mean  
89 value of each latitudinal zone.

90 The diversities were estimated using Shareholder Quorum Subsampling (Alroy,  
91 2010) (herein  $D_{SQS}$ ), Shannon Entropy Hill number (Chao et al., 2014) (herein  $D_{Chao}$ ), and

92 the capture-mark-recapture (CMR) method (Liow and Nichols, 2010) (herein  $D_{CMR}$ ).  
93 These three methods were chosen because they represent independent approaches to  
94 account for sampling and preservation bias, which in their entirety give an estimate on  
95 the robustness of the estimated diversity pattern (Liow and Nichols, 2010; Chao et al.  
96 2014, see the Data Repository). Because all three diversity measures,  $D_{SQS}$ ,  $D_{Chao}$ , and  
97  $D_{CMR}$ , exhibit a robust trend (Fig. DR1 in the Data Repository), only the  $D_{CMR}$  trends are  
98 described in detail in the main article, which additionally produced the most realistic (=   
99 largest) 95% confidence intervals. The downloaded data from the PaleoBioDB and the R  
100 code of the analysis are available at <https://doi.org/10.5281/zenodo.197057>.

## 101 **RESULTS: CHANGING LDGs DURING THE COURSE OF THE** 102 **DIVERSIFICATION**

103 Ordovician diversity exhibits a trend with relatively low estimates for the first ~18  
104 m.y. (Tremadocian–Dapingian), a steep increase during the remaining Middle Ordovician  
105 (Darriwilian), a Late Ordovician plateau, and a drastic decrease during the terminal  
106 Ordovician (Fig. 1). When partitioned into four 15° paleolatitudinal zones it becomes  
107 apparent that the bulk of the Ordovician diversity increase took place within the tropical  
108 realm (0–30°, Fig. 2). In temperate latitudes the GOBE is hardly visible.

109 This predominance of the tropical diversification can be interpreted as real  
110 because it is apparent in all diversity estimates explored herein (Fig. DR1), and it does  
111 not reflect changes in the relative abundance of carbonate/siliciclastic and shallow/deep  
112 depositional environments, in environmental and lithologic heterogeneity, or in  
113 paleogeographical shift (Fig. 3; Fig. DR5). However, although the pattern is not restricted

114 to a specific organismal clade because it is apparent in e.g., brachiopods, mollusks, and  
115 trilobites, it is less pronounced or absent in planktic graptolites (Figs DR2 and DR3).

116       When comparing the per-time-bin diversity estimates against their paleolatitudinal  
117 zone, an Ordovician LDG trend becomes apparent, the most important features of which  
118 are an Early to Middle Ordovician (Tremadocian–Darriwilian) peak at the 15–45° zone, a  
119 Late Ordovician (Sandbian–Katian) change toward higher tropical diversities with a peak  
120 at 0–30°, and again a 15–30°-zone peak during the terminal Ordovician (Hirnantian) (Fig.  
121 2; Fig. DR4). Notably, the Darriwilian marks not only the climax of the diversification,  
122 but also a major pattern shift in Ordovician LDGs toward maximum values in the tropics.

123       LDG trends are an effect of changes in evolutionary rates and dispersal (see  
124 Jablonski et al. 2016). Dispersal dynamics are driven by differential rates of active and  
125 tectonic range shifts (see Methods). These dispersal factors are not correlated in the  
126 Ordovician data, and tectonic shift appears to play a minor role in the overall picture (Fig.  
127 3). The relative range shift pattern exhibits considerable variation during the Ordovician  
128 with different trends near the equator and in temperate paleolatitudes (Fig. 2). In the high  
129 tropics (0–15°), pole-ward shifts were relatively rare during the Early Ordovician but  
130 became abruptly more common during the late Middle and Late Ordovician. In contrast,  
131 in low tropical and temperate paleolatitudes, the trend exhibits initial (Tremadocian) high  
132 rates of pole-ward shifts, relatively high rates of equator-ward shifts during the Floian-  
133 Dapingian and an abrupt change toward moderate rates of pole-ward and equator-ward  
134 shifts during the Darriwilian.

135       These trends reveal an intricate change in the general migration dynamics of the  
136 Ordovician world with shifting diversity sinks and sources (Fig. 4). One prominent

137 feature of this pattern is the change of the high tropics from a sink during the Early  
138 Ordovician toward a source during the latter part of that period. This pattern is consistent  
139 with a change toward a modern-type “out of the tropics” scenario (Jablonski et al., 2006)  
140 during the Darriwilian concurrent with the main diversification pulse of the GOBE.

#### 141 **DISCUSSION: CHANGING LDGs AND GLOBAL COOLING**

142 Previous analysis of Ordovician diversity trends below global level have focused  
143 on paleocontinents or paleo-oceanic regions that range across paleolatitudinal zones and  
144 often revealed heterogenous diversity pattern that are difficult to interpret (Miller, 1997;  
145 Webby et al., 2004). Only recently have synchronous Darriwilian diversification peaks  
146 within low paleolatitude regions of Baltica and Laurentia been detected in brachiopods;  
147 these results were used to infer global drivers, such as climatic cooling, intensified sea-  
148 water circulation, greater oxygenation and increased primary production (Trubovitz and  
149 Stigall, 2016). Here, I show that, although the Darriwilian peak diversification affected  
150 all latitudinal zones, it had by far the most profound impact and highest amplitude within  
151 the high tropics (Fig. 2). Combined with escalating rates of ex-tropical range shifts during  
152 the Middle Ordovician, this tropical diversification appears to be primarily caused by in  
153 situ origination, and in turn this tropical in situ origination must have been the  
154 quantitatively most important driver of the GOBE.

155 If this pattern is realistic, then tropical in situ origination is crucial to  
156 understanding the GOBE and hence the question of how this high tropical diversity was  
157 produced comes into focus. In a recent study and review this question was addressed with  
158 respect to the GOBE in general with oscillations between events of immigration and  
159 geographic differentiation suggested as one effective “speciation dynamo” (Stigall,

160 2017). Therein, geographic differentiation, conceptualized as tectonically induced  
161 regional isolation (e.g., Harper, 2010) and habitat heterogeneity (e.g., Miller and Mao,  
162 1995) were considered to be key factors in promoting origination (Stigall, 2017).  
163 However, the habitat heterogeneity trends estimated herein are not related to the diversity  
164 pattern and generally the heterogeneity is not higher in the tropics (Fig. DR5). Because it  
165 is currently not known if regional differentiation was more intense in the tropics during  
166 the GOBE, these two factors, although not in conflict, cannot explain the LDG pattern  
167 detected herein.

168         Alternatively, geographic differentiation can be more generally conceptualized as  
169 ecological niche differentiation, including biologically-induced niche differentiation  
170 (*sensu* Sepkoski, 1988). With this more general conceptualization it is possible to set the  
171 GOBE into context with the temperature and evolutionary speed (TES) hypotheses of  
172 e.g., Allen et al. (2002), Clarke and Gaston (2006), and Brown (2014), which link  
173 ecological and evolutionary rates with temperature and predict highest diversities in the  
174 tropics. Generally, the Ordovician diversity trend is contrary to naive expectations from  
175 TES hypotheses, because they would predict decreasing ecological and evolutionary  
176 rates, and diversities and flattened LDG amplitudes under climatic cooling (Tittensor et  
177 al., 2010; Brown, 2014; Tittensor and Worm, 2016). This mismatch, however, can be best  
178 explained by the existing range limits of the published diversity-temperature  
179 dependencies. All published TES hypotheses are based on empirically observed positive  
180 diversity-temperature dependencies. These dependencies are constrained by Recent sea  
181 surface temperatures with a maximum of ~30 °C (e.g., Brown, 2014). Estimated Early  
182 Ordovician tropical sea surface temperatures are well beyond today's maximum values



183 (Trotter et al., 2008, Veizer and Prokoph, 2015). It is not known what diversity-  
184 temperature dependencies would be like in a hypothetical expanded temperature range,  
185 but thermal tolerances of complex multicellular organisms are limited, and today an  
186 upper limit for metazoans exists at ~45–47° (Ravaux et al., 2013). These limitations  
187 suggest that past diversity-temperature curves beyond modern temperature ranges had a  
188 parabolic shape with a diversity maximum in moderate temperatures and decreasing  
189 trajectories toward extreme temperatures. Past LDG's under hotter-than-today global  
190 regimes theoretically should reflect these expected trajectories with diversity peaks in  
191 temperate latitudes rather than near the equator. Data from Mesozoic and Cenozoic  
192 greenhouse intervals support this assumption (e.g., Markwick, 1998; Marcot et al., 2016)  
193 and the new data from the Ordovician are consistent with earlier assessments that tropical  
194 peak diversities are an exclusive feature of LDGs of icehouse periods (Mannion et al.,  
195 2014). Additionally, the Ordovician shift of peak LDG values toward the equator  
196 coincides with rapid initial expansion and diversification of metazoan reefs after a long  
197 period of dominance of reefs built by calcimicrobes, suggesting a (re)colonization of the  
198 low latitudes after a prolonged period of too-hot temperatures in the tropics (Webby,  
199 2002; Kröger et al. 2017).

200       Hence, the mid-Ordovician cooling must have been a major factor for the  
201 diversification of the tropics. But why did this tropical diversification and LDG shift have  
202 such an overwhelming effect on global net diversity? An answer to this question is not  
203 easily found with the data at hand. The tropical mid-Ordovician diversification took place  
204 in a larger area and in regions with less intense seasonality compared with the temperate  
205 locus of the diversification during the Early Ordovician. The effects of these differences

206 on global Ordovician diversity are currently unknown and offer great potential for future  
207 investigations. Furthermore, the tropical mid-Ordovician diversification did not affect all  
208 clades. Brachiopods, mollusks, and trilobites diversified mainly in low latitudes and  
209 during the Darriwilian, whereas the tropical diversification of graptolites occurred earlier  
210 and the Darriwilian pulse is visible only in temperate latitudes. This asynchronicity  
211 between clades was known before (e.g., Webby et al., 2004) and new studies potentially  
212 will be key to answering this question in the future.

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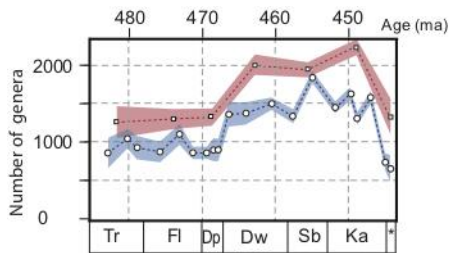
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- 334

335

336 **FIGURE CAPTIONS**



337

338 Figure 1. Global genus-level diversity trends for the Ordovician Period with stratigraphic  
339 resolution at stage level (squares) and stage slice level (circles, Bergström et al., 2009).

340 The peak diversification of the Great Ordovician Biodiversification Event (GOBE) was  
341 during the Darriwilian Stage. Data downloaded from Paleobiology Database

342 (PaleoBioDB, <https://paleobiodb.org>) on 26 November 2016. D<sub>CMR</sub>—diversity estimate  
343 from capture-mark-recapture (CMR) modeling approach with 95% confidence intervals

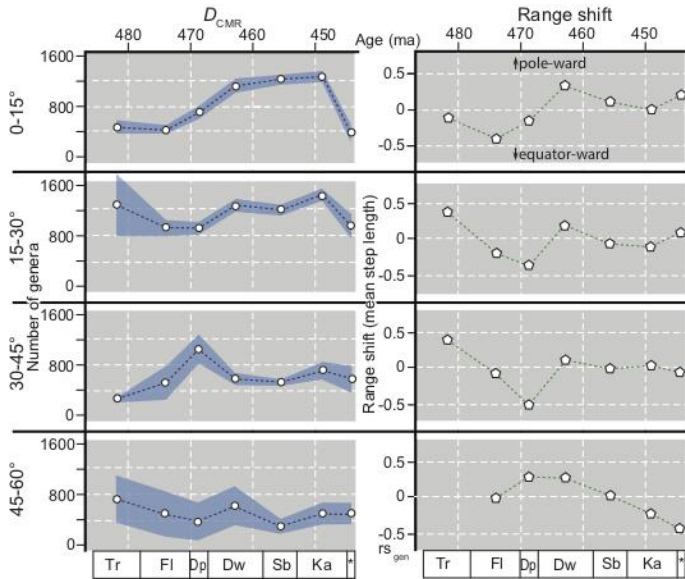
344 (shaded areas). Stages: Tr—Tremadocian; Fl—Floian; Dp—Dapingian; Dw—

345 Darriwilian; Sb—Sandbian; Ka—Katian, \*—Hirnantian.

346



347

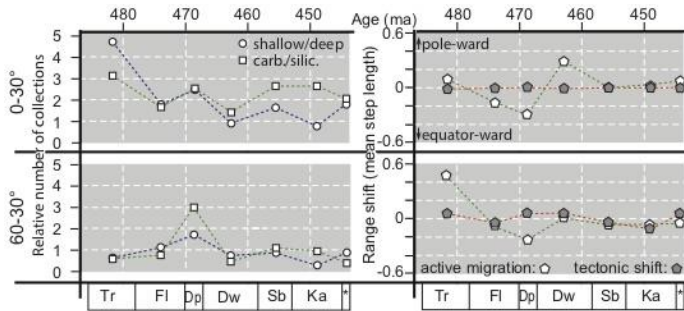


348

349 Figure 2. Ordovician diversity trends and range shift dynamics within four  
350 paleolatitudinal zones (Northern and Southern Hemispheres combined). Note the massive  
351 Darrivilian diversification pulse in low paleolatitudes.  $D_{CMR}$ , diversity estimate from  
352 capture-mark-recapture (CMR) modeling approach with 95% confidence intervals  
353 (shaded areas).  $n_{gen}$ —number of genera;  $rs_{gen}$ —mean range shift (given as distance from  
354 Ordovician mean). Positive values indicate dominance of pole-ward; negative values  
355 dominance of equator-ward shifts.

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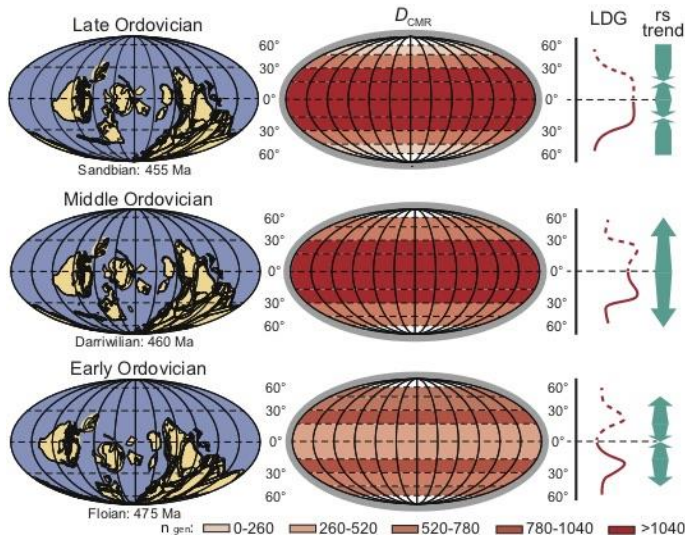


358

359 Figure 3. Trends in relative abundance of Ordovician fossil occurrences downloaded  
360 from Paleobiology Database (PaleoBioDB, <https://paleobiodb.org>) on 26 November 2016  
361 from different environments and lithologies (carb.—carbonatic; silic. —siliciclastic), and  
362 direction of active/tectonic shift within two paleolatitudinal zones (Northern and  
363 Southern Hemispheres combined). Note the differences between active/tectonic shift,  
364 environmental trends, and diversity trends (see Fig. 2).

365

366



367

368 Figure 4. Summary of global change during the Ordovician. Paleogeographic  
369 reconstructions from BugPlates (<http://www.geodynamics.no/bugs>). Diversity estimates  
370 and range shift dynamics simplified from Figure 2 (see also Fig. DR4 [see footnote 1]).  
371 Estimates are combined for northern and southern hemisphere and are symmetrically  
372 plotted across equator for simplicity.  $n_{gen}$ —number of genera.

373

374 GSA Data Repository item 2018xxx, xxxxxxxx, is available online at  
375 <http://www.geosociety.org/datarepository/2018/>, or on request from  
376 [editing@geosociety.org](mailto:editing@geosociety.org).