

1 **Title: Functional diversity of marine ecosystems after the Late Permian mass**
2 **extinction event**

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11 **Introductory paragraph:** The Late Permian mass extinction event was the most severe such
12 crisis of the past 500 million years and occurred during an episode of global warming. It is
13 assumed to have had significant ecological impact, but its effects on marine ecosystem
14 functioning are unknown and the patterns of marine recovery are debated. We analysed the fossil
15 occurrences of all known Permian-Triassic benthic marine genera and assigned each to a
16 functional group based on their inferred life habit. We show that despite the selective extinction
17 of 62-74% of marine genera there was no significant loss of functional diversity at the global
18 scale, and only one novel mode of life originated in the extinction aftermath. Early Triassic
19 marine ecosystems were not as ecologically depauperate as widely assumed, which explains the
20 absence of a Cambrian-style Triassic radiation in higher taxa. Functional diversity was, however,
21 significantly reduced in particular regions and habitats, such as tropical reefs, and at these scales
22 recovery varied spatially and temporally, probably driven by migration of surviving groups.
23 Marine ecosystems did not return to their pre-extinction state, however, and radiation of
24 previously subordinate groups such as motile, epifaunal grazers led to greater functional evenness
25 by the Middle Triassic.

26 **Main text**

27 The Late Permian extinction event, which occurred $252.28 \pm 0.08\text{Ma}$ (1), during a
28 period of climate warming (2-3), was the largest such crisis of the Phanerozoic in terms of
29 both diversity loss (4-6), and long term ecological impact (7). Local-scale, field-based studies
30 have suggested, however, that extinction magnitude and rate of subsequent recovery varied
31 with depositional setting (8-12), palaeolatitude (10, 13-14) and region (15-16). The ecological
32 impact of past extinctions is, however, difficult to quantify and is arguably poorly understood
33 (17).

34 Here we use the ecospace model of Bambach et al. (18) to provide the first
35 quantitative analysis of the autecology of all known Permian-Triassic benthic marine genera.
36 This approach has been successfully applied to analysis of the Cambrian radiation, and to
37 compare the functional diversity of Palaeozoic and modern ecosystems (e.g. 19-20), but has
38 not been used previously to quantify ecological change during extinction events. Our
39 analyses show that the Late Permian extinction event caused almost no loss of overall
40 functional diversity at the global scale, but did result in significant ecological shifts in benthic
41 marine communities from particular depositional settings, palaeolatitudes and regions.

42 Two complementary databases were constructed and analysed (available from
43 <http://www.pangaea.de/>, doi to be confirmed). Firstly, a database of all known marine
44 invertebrate genus-level occurrences from the Late Permian to the Middle Triassic, compiled
45 from the Paleobiology Database <www.pbdb.org> (PaleoDB), supplemented by additional
46 published data, along with locational, stratigraphic, lithological and environmental data. This
47 database comprises 22,263 recorded occurrences of 1420 genera. Occurrence data do not
48 capture Lazarus taxa, however, especially those that range-through, but do not occur within,
49 the study interval, and therefore underestimate the total number of genera known to have
50 existed in any particular time interval. Therefore, a second database comprising range-

51 through data of 1770 genera was compiled by ranging-through the occurrence database, and
52 supplementing these data with additional data from the PaleoDB and Sepkoski (21) [see
53 Supplementary Information].

54 The occurrences were binned into 12 unequal-duration time intervals equivalent to
55 Permian ages and Triassic sub-ages. Vetting of the data followed (22) [see Supplementary
56 Information]. Modes of life were inferred by using data from extant relatives, previous
57 publications and functional morphology, and each genus was assigned to a bin in the
58 ecospace model of (18) based on its tiering, motility and feeding habits [see Supplementary
59 Information]. Occurrences with sufficient geological information were assigned one of five
60 broad depositional settings along an onshore-offshore gradient: Inner Shelf (i.e. above fair
61 weather wave base), Middle Shelf (between fair and storm weather wave base), Outer Shelf
62 (between storm weather wave base and the shelf edge), Slope and Basin (beyond the shelf
63 edge), and Reefs (including build ups and bioherms). All occurrences were assigned to one of
64 the following palaeolatitudinal zones: Tropics (within 30° of the palaeoequator), northern
65 (>30° north) and southern (> 30° south). Finally, all occurrences were assigned to one of four
66 regions: Panthalassa, Boreal Ocean, Palaeotethys and Neotethys.

67 Aggregation of data within these bins was quantified using Green's coefficient of
68 dispersion (23). High Green coefficient values indicate dominance by one or two categories
69 within a particular stratigraphic bin, with values >0.5 considered to represent a significant bias
70 (24). In order to take into account variations in sampling intensity within and between the
71 different time bins and categories we applied the subsampling routine of Miller and Foote (25)
72 using a modified R script of Lloyd et al. (26) to standardise the functional richness based on the
73 number of occurrences. Subsampling quotas were set at 250 occurrences per bin for the global
74 analysis and 100 occurrences for the smaller scale analyses of palaeolatitude, region or
75 depositional setting. In each case, time bins with less than the specified quota of occurrences were
76 excluded from subsampling. This compromise avoids the degradation that may occur if the

77 quality of the poorest bin is used as the subsampling quota (26), but we recognise that higher
78 subsampling levels could reveal additional variability.

79 **Global-scale changes in functional diversity**

80 Global genus-level extinction is estimated at 62% (using range-through data) or 74%
81 (using occurrence data). There are often considerable differences between the range-through
82 and occurrence data, with the former providing lower extinction estimates for most modes of
83 life (Fig. 1). This difference is an unsurprising consequence of the poor quality Early Triassic
84 fossil record, which is due to a variety of factors including rock record biases, incomplete
85 sampling and poor preservation. Thus, while occurrence data indicate that as many as nine
86 modes of life completely disappeared across the P/Tr boundary (Fig.1), six of these are
87 recorded by the range-through data, implying that genera exhibiting those particular modes of
88 life have either been overlooked, were present but at abundances that were too low to be
89 recorded by the fossil record, or existed at locations other than those that have been sampled
90 to date. Thus, at most, only three modes of life disappeared across the P/Tr boundary:
91 surficial, fast moving predators; semi-infaunal, stationary, unattached deposit feeders; and
92 deep infaunal, facultatively motile, unattached suspension feeders (Fig. 1).

93 Range-through data at the taxonomic resolution of families and superfamilies, which
94 have longer ranges than their constituent genera and may span an interval of time from which
95 no constituent genera have been recorded, show, however, that even this is an over-estimate
96 (Fig. 2a). The boundary-crossing family Erymidae demonstrates that at least one genus of
97 surficial, fast moving, predator must have been present somewhere. Likewise, deep infaunal,
98 facultatively motile, unattached, suspension feeders must have survived the extinction event,
99 as demonstrated by the boundary-crossing superfamily Pholadomyoidea. Thus, it would
100 appear that the greatest mass extinction event of the past 500 million years led to the loss of
101 just a single mode of life – that of semi-infaunal, stationary, unattached deposit feeders; a

102 consequence of the extinction of the rostroconchs. Subsampling to standardise the global
103 occurrence data also supports a lack of change in functional diversity across the P/Tr
104 boundary, despite the significant drop in generic richness (Fig. 2b). Therefore, overly
105 simplistic models suggesting that Early Triassic marine ecosystems functioned for millions of
106 years with primary producers only and no higher trophic levels (e.g. 27) can be rejected.

107 The resistance of functional groups to global extinction, even those with few
108 constituent genera, is further demonstrated by the upward trajectory of functional diversity
109 since the start of the Phanerozoic (18-20; 28). Even though functional diversity was reduced
110 locally and regionally, for example in the tropics, because the Late Permian extinction did not
111 reduce functional diversity at the global scale an ecologically diverse group of marine
112 metazoan survivors were able to refill this vacant ecospace once conditions ameliorated (29-
113 30). This explains why relatively few classes, orders, or phyla originated in the early
114 Mesozoic, despite a return to levels of taxonomic diversity that were similar to the early
115 Palaeozoic (29). This may also explain why just one novel mode of life originated in the
116 Early Triassic; namely the erect, facultatively motile, attached suspension feeders, which
117 appeared with the evolution of motile crinoids in the Smithian (31-32; Fig. 2a). Globally, the
118 Early Triassic benthic ecosystem functioned much like a ship manned by a skeleton crew;
119 each post was occupied, but by only a few individual taxa.

120 The Late Permian extinction event was highly selective (Fig. 3; 29), with the erect and
121 surficial, stationary, attached suspension feeders comprising 64% of the genera that
122 disappeared. Selection against these formally dominant groups led to increases in the relative
123 abundance at the global scale of most other groups during the Induan (Fig. 3). Despite
124 slightly increasing in relative abundance in the Olenekian, sessile epifaunal suspension
125 feeders never returned to the same level of dominance that characterised the Lopingian.
126 Instead, the appearance of motile crinoids (31) and the radiation of the ‘modern biota’,

127 including motile and infaunal bivalves (34), predatory arthropods (35), and grazing
128 gastropods (36), increasingly filled a diverse suite of lifestyles during the early Mesozoic (Fig.
129 3). By the Middle Triassic, functional evenness at the global scale had increased, due
130 especially to a radiation of the mobile grazing epifauna. Beginning even before the Late
131 Permian extinction event, surficial, slow moving grazers underwent a gradual, uninterrupted
132 increase in relative abundance through the entire study interval (Fig. 3). Apart from those
133 groups that originated after the extinction, this represents the most important change to the
134 functional landscape of benthic marine ecosystems during the earliest stages of the Mesozoic
135 Marine Revolution.

136 **Latitudinal trends and regional biases**

137 Although the Late Permian extinction event eliminated just a single mode of life at the
138 global scale, at smaller scales functional diversity loss was more pronounced. Prior to the
139 extinction event, equatorial palaeolatitudes, as expected, housed the greatest ecological and
140 functional diversity (Fig. 4a). Following extinction, however, raw and subsampled data
141 record similar levels of functional diversity in tropical and northern palaeolatitudes during the
142 Induan, and especially in the Dienerian (Fig. 4a). This low gradient of functional diversity
143 between the tropics and northern belt during the Induan may reflect the cosmopolitan nature
144 of Induan benthic faunas (5) and implies relatively greater ecological impact in the tropics.
145 Greater ecological impact in the tropics may be a consequence of climate warming, including
146 concomitant factors such as expansion of hypoxic dead zones, reduced circulation, reduced
147 nutrient cycling, and temperature rise itself, as well as the loss of reef ecosystems.

148 By the Anisian, raw and subsampled data show that tropical functional diversity had
149 exceeded that recorded at low palaeolatitudes during the Lopingian, and a steep functional
150 gradient between the tropics and higher latitudes had been re-established (Fig. 4a). Greater
151 tropical functional diversity during the Middle Triassic is probably due to migration from

152 higher latitudes of taxa with lifestyles that were not previously recorded in the tropics. In
153 many time bins, however, occurrences are not distributed evenly between palaeolatitudes, but
154 are biased towards the tropics (Green coefficients > 0.5 ; Fig. 4a). The Spathian and Anisian
155 record the greatest such bias, being dominated by well-studied sections from Europe and
156 South China (37-39), and the ‘global’ record of Spathian-Anisian recovery is therefore really
157 a ‘tropical’ record.

158 Raw functional diversity decreases from the Changhsingian to the Griesbachian in all
159 regions except the Boreal Ocean, which records an earlier decline, with minima being
160 recorded at different times within the Early Triassic (Fig. 4b). Results of subsampling suggest,
161 however, that except for Panthalassa these trends may be an artefact of sampling biases (Fig.
162 4b). The timing and patterns of recovery of functional diversity also apparently vary between
163 regions, although there are too few post-Griesbachian occurrences in Neotethys and the
164 Boreal Ocean for robust subsampling. A return to greater functional diversity occurred sooner
165 in Panthalassa than in Palaeotethys, although relative differences during the Middle Triassic
166 may be due to sampling problems because the regional occurrence data are strongly biased
167 towards Palaeotethys during the Anisian (Green coefficients >0.5 ; Fig. 4b). Indeed, the
168 widely held view that post-extinction recovery did not occur until the Anisian (e.g. 27) is
169 likely due to this regional bias and the ‘global’ signal is evidently just a Palaeotethyan signal.

170 **Functional diversity in reefs and shelf settings**

171 Although most Early Triassic occurrences are from the inner shelf, there is no major
172 sampling bias between depositional settings through the study interval (Green coefficients
173 <0.5 ; Fig. 4c). It is well known that reefs disappeared during the earliest Triassic (e.g. 40),
174 but our raw and subsampled data demonstrate for the first time that the major loss of
175 functional diversity in reef ecosystems occurred significantly before the late Changhsingian
176 mass extinction horizon (Fig. 4c). During the Wuchiapingian, metazoan reefs were globally

177 the most important habitat in terms of functional richness, recording 19 of the 23 known
178 modes of life (i.e. 83%), from 74 localities in UK, Russia, Germany, Greece, China,
179 Greenland, Lithuania and Pakistan. Only four modes of life are, however, recorded in
180 Changhsingian reefs, from a total of 36 localities from China, Russia, Tajikistan, Greece and
181 Thailand. These data imply that reef ecosystems underwent precipitous collapse, and/or a
182 major contraction, prior to climate warming in the latest Changhsingian. This may have been
183 due to Lopingian sea-level fall or to a currently unknown episode of climatic or
184 oceanographic change, possibly related to the earliest phases of Siberian Trap volcanism (41).
185 The subsequent rebuilding of reefs began locally in the Smithian (42) and Spathian, involving
186 organisms such as sponges (42), cementing bivalves (43) and *Tubiphytes* (44). Metazoan
187 reefs of the Anisian were as functionally rich as their Wuchiapingian predecessors (Fig. 4c),
188 although reefs were not common until after the Illyrian following the appearance and
189 radiation of the Scleractinia.

190 During the Early Triassic 'reef gap', basin and slope settings gradually lost functional
191 richness, whereas shelf seas, especially the inner and outer shelf, were functionally the most
192 diverse settings (Fig. 4c). By the Middle Triassic, inner shelf settings housed greater
193 functional richness than the re-emerging reefs. The 'habitable zone' hypothesis (10) predicts
194 that nearshore, wave-aerated settings should house greater diversity in the earliest Triassic
195 and although our inner shelf data support this hypothesis, the outer shelf data imply that even
196 if deeper settings were taxonomically depauperate functional diversity was still maintained, at
197 least in the Griesbachian (Fig. 4c).

198 Even though Early Triassic shelf settings were functionally diverse, the loss of erect
199 and deep infaunal taxa led to a reduction in occupied tiers. This restriction of benthic animals
200 to within a few centimetres above and below the sediment-water interface is the most
201 dramatic such shift of the entire Phanerozoic (45), and characterises the immediate extinction

202 aftermath worldwide (14; 46). It would have significantly impacted nutrient cycling and
203 secondary production, with bioturbation depths in most ecosystems returning to levels not
204 recorded globally since the Cambrian. Genera identified as being ‘deep infaunal’ reappear
205 first in the Smithian of eastern Panthalassa, but are represented by small-sized taxa such as
206 *Sinbadiella pygmaea* (47), which probably could not burrow that deeply. Full ecosystem
207 function would not have been restored until larger bioturbators reappeared. The earliest
208 reappearances of the erect tier are also represented by relatively small-sized animals: in the
209 inner shelf by bryozoans in the Dienerian (48); and in slope and basin settings by the crinoid
210 *Holocrinus* in the Smithian (49).

211 Despite the significant worldwide loss of species richness, the Late Permian
212 extinction event did not significantly reduce functional diversity of benthic marine
213 ecosystems at the global scale. As a consequence, few higher taxa and only one new mode of
214 life originated in the extinction aftermath. At smaller scales, however, significant changes in
215 functional diversity did occur in particular regions and depositional settings. Ecological
216 changes during this critical time in Earth history are more complex than simple models have
217 hitherto suggested, and supposed ‘global’ patterns may reflect biases in regional data,
218 especially in the post-extinction recovery interval.

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355

356 WJF and RJT are equally responsible for the project design, interpretation, and writing. WJF
357 compiled the databases and undertook the analyses.

358

359 **Fig 1:** The percentage of genera that became extinct within the Changhsingian in each
360 individual mode of life, for (a) range-through and (b) occurrence data. The numbers of
361 genera recorded in the Changhsingian (bold) and surviving into the Griesbachian (grey) are
362 reported above each respective bar. Error bars represent one standard error in each direction
363 [supplementary information]. 1 = erect, stationary, attached, suspension feeder; 2 = surficial,
364 fast moving, deposit feeder; 3 = surficial, fast moving, predator; 4 = surficial, slow moving,
365 suspension feeder; 5 = surficial, slow moving, deposit feeder; 6 = surficial, slow moving,
366 grazer; 7 = surficial, slow moving, predator; 8 = surficial, facultatively motile, unattached,
367 suspension feeder; 9 = surficial, facultatively motile, attached, suspension feeder; 10 =
368 surficial, facultatively motile, attached, deposit feeder; 11 = surficial, stationary, unattached,
369 suspension feeder; 12 = surficial, stationary, attached, suspension feeder; 13 = semi-infaunal,
370 slow moving, suspension feeder; 14 = semi-infaunal, facultatively motile, attached,
371 suspension feeder; 15 = semi-infaunal, stationary, unattached, deposit feeder; 16 = semi-

372 infaunal, stationary, attached, suspension feeder; 17 = shallow infaunal, slow moving,
373 suspension feeder; 18 = shallow infaunal, slow moving, miner; 19 = shallow infaunal, slow
374 moving, predator; 20 = shallow infaunal, facultatively motile, unattached, suspension feeder;
375 21 = shallow infaunal, facultatively motile, unattached, miner; 22 = shallow infaunal,
376 facultatively motile, attached, suspension feeder; 23 = shallow infaunal, facultatively motile,
377 attached, other; 24 = deep infaunal, facultatively motile, unattached, suspension feeder; 25 =
378 deep infaunal, facultatively motile, attached, other. Images of fauna are not to scale.

379

380 **Fig 2:** Diversity curves for generic and functional richness across the studied interval. **(a)**
381 Generic richness of raw range-through data (filled squares) and occurrence data (filled
382 circles); and number of modes of life occupied using generic range-through (open triangles),
383 generic occurrence data (open squares), and range-through data for higher taxonomic levels
384 (open circles). **(b)** Subsampled generic and functional richness using occurrence data. Median
385 generic richness (open circles) and median number of modes of life occupied (filled squares).
386 Error bars represent 5th and 95th percentiles. The Lopingian and Middle Triassic epochs are
387 shaded, bold vertical line shows the Permian/Triassic boundary.

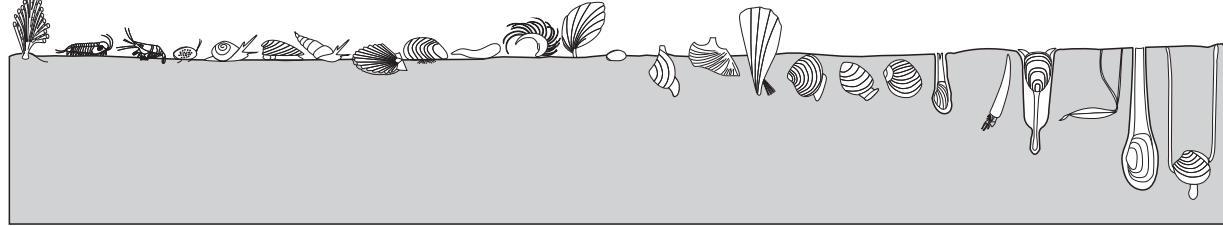
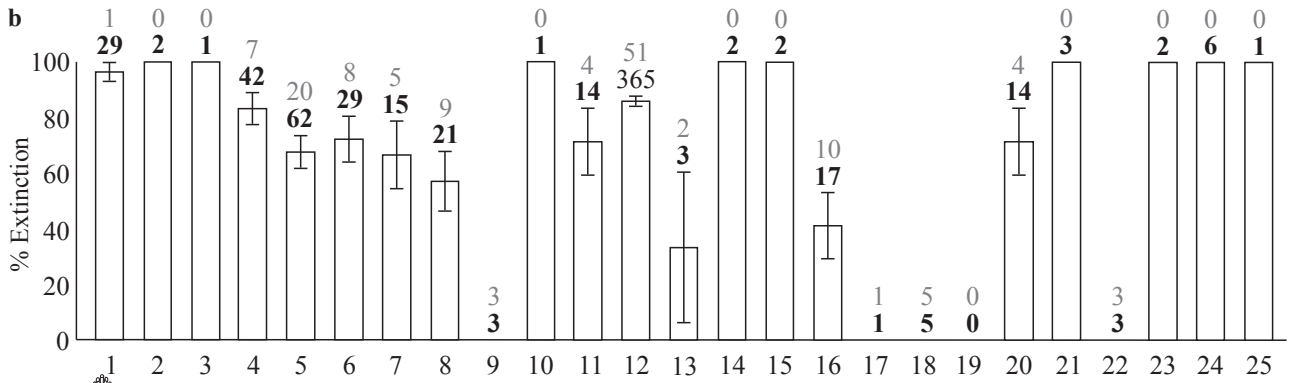
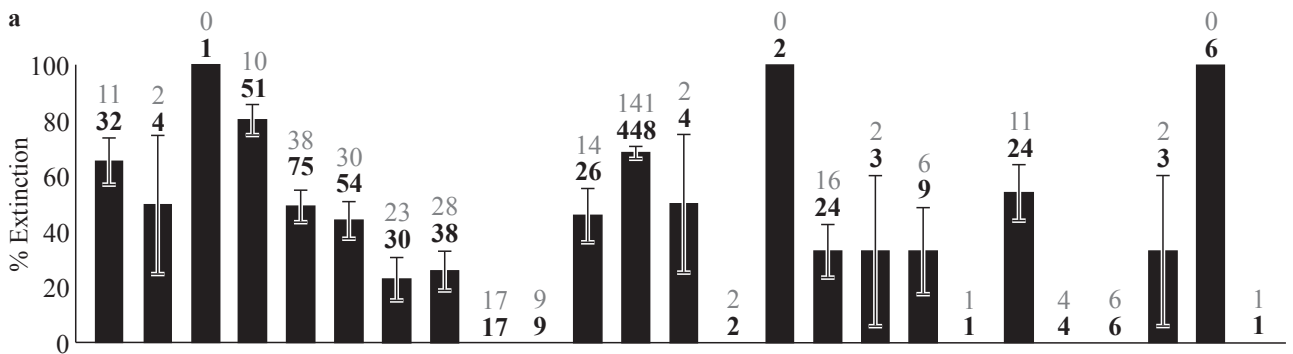
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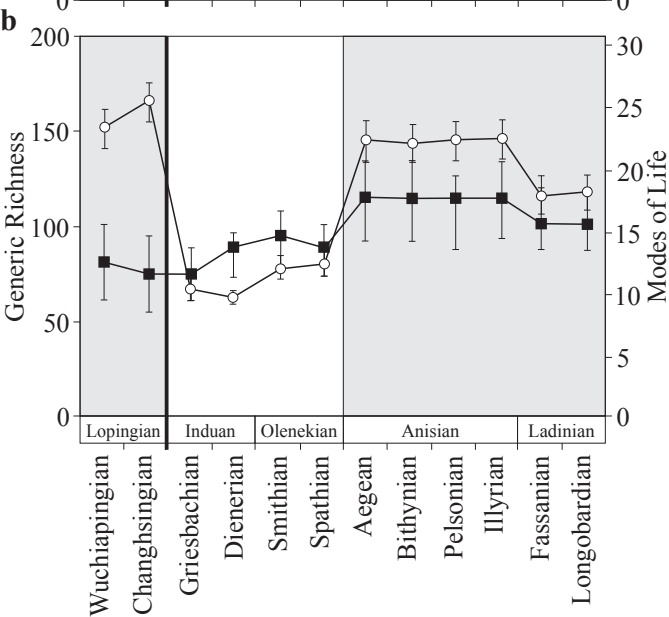
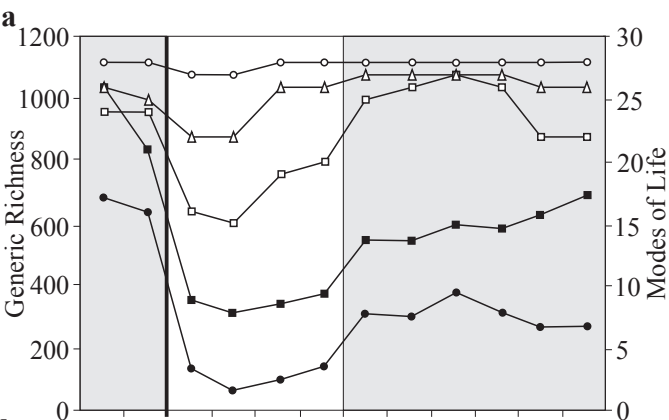
389 **Fig 3:** Relative abundance of genera in each mode of life across the studied interval. Colours
390 indicate changes in abundance of >0.1%: increases (red), decreases (blue) and no change
391 (yellow) from the previous time bin. Wuchiapingian abundances are unshaded. 1 = erect,
392 facultatively motile, attached suspension feeder. 2 = erect, stationary, attached suspension
393 feeder; 3= surficial, fast moving suspension feeder; 4 = surficial, fast moving deposit feeder;
394 5 = surficial, fast moving predator; 6 = surficial, slow moving suspension feeder; 7 = surficial,
395 slow moving deposit feeder; 8 = surficial, slow moving grazer; 9 = surficial, slow moving
396 predator; 10 = surficial, facultatively motile, unattached suspension feeder; 11 = surficial,

397 facultatively motile, attached suspension feeder; 12 = surficial, facultatively motile, attached
398 deposit feeder; 13 = surficial, stationary, unattached suspension feeder; 14 = surficial,
399 stationary, attached suspension feeder; 15 = semi-infaunal, slow moving suspension feeder;
400 16 = semi-infaunal, facultatively motile, unattached predator; 17 = semi-infaunal,
401 facultatively motile, attached suspension feeder; 18 = semi-infaunal, stationary, unattached
402 deposit feeder; 19 = semi-infaunal, stationary, attached suspension feeder; 20 = shallow
403 infaunal, slow moving suspension feeder; 21 = shallow infaunal, slow moving miner; 22 =
404 shallow infaunal, slow moving predator; 23 = shallow infaunal, facultatively motile,
405 unattached suspension feeder; 24 = shallow infaunal, facultatively motile, unattached miner;
406 25 = shallow infaunal, facultatively motile, attached suspension feeder; 26 = shallow infaunal,
407 facultatively motile, attached other; 27 = deep infaunal, facultatively motile, unattached
408 suspension feeder; 28 = deep infaunal, facultatively motile, unattached suspension feeder; 29
409 = deep infaunal, facultatively motile, attached other. Images of fauna are not to scale.

410

411 **Fig. 4.** Permian-Triassic functional richness in different (a) palaeolatitudes, (b) regions, and
412 (c) environments. Raw data (open circles) and subsampled medians (filled squares). Error
413 bars represent 5th and 95th percentiles. Green coefficient values >0.5 indicate bias in the data
414 within that time bin. The Lopingian and Middle Triassic epochs are shaded. The Permian/
415 Triassic boundary is marked with a thick line.





Relative abundances

○ 0.1-1% ○ 2-5% ○ 6-10% ○ 11-20% ○ 21-50% ○ >50%

