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**Article:**

Dai, X, Song, H, Wignall, PB et al. (5 more authors) (2018) Rapid biotic rebound during the late Griesbachian indicates heterogeneous recovery patterns after the Permian-Triassic mass extinction. *GSA Bulletin*, 130 (11-12). pp. 2015-2030. ISSN 0016-7606

<https://doi.org/10.1130/B31969.1>

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1        Rapid biotic rebound during the late Griesbachian  
2        indicates heterogeneous recovery patterns after the  
3                                  Permian-Triassic mass extinction  
4

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12    **ABSTRACT**

13        New fossil data of two Early Triassic (Griesbachian to Dienerian) sections from  
14        South China show unusually high levels of both benthic and nektonic taxonomic  
15        richness, occurring in the late Griesbachian. A total of 68 species (including 26 species  
16        of Triassic-type species) representing mollusks, brachiopods, foraminifers, conodonts,  
17        ostracods, and echinoderms occur in the late Griesbachian, indicating well-established  
18        and relatively complex marine communities. Furthermore, the nekton shows higher  
19        origination rates than the benthos. Analyses of sedimentary facies and size distribution  
20        of pyrite framboids show that this high-diversity interval is associated with well-  
21        oxygenated environments. In contrast to the previously suggested scenario that  
22        persistently harsh environmental conditions impeded the biotic recovery during the

23 Early Triassic, our new findings, combined with recent works, indicate a fitful regional  
24 recovery pattern after the Permian-Triassic crisis resulting in three main diversity highs:  
25 late Griesbachian-early Dienerian, early-middle Smithian and Spathian. The transient  
26 rebound episodes are therefore influenced by both extrinsic local (e.g. redox condition,  
27 temperature) and intrinsic (e.g. biological tolerances, origination rate) parameters.

## 28 **INTRODUCTION**

29 The Permian-Triassic mass extinction (PTME) was the largest biotic catastrophe  
30 of the Phanerozoic, which eliminated around 80% to 90% of marine species (Raup,  
31 1979; Song et al., 2013; Stanley, 2016) and heralded the development of modern  
32 ecosystems (Sepkoski, 1981; Brayard et al., 2017). After this mass extinction,  
33 depauperate faunas were prevailed throughout the Early Triassic, and not until the  
34 Middle Triassic did diversity rebound (Erwin and Pan, 1996; Nützel, 2005; Payne et al.,  
35 2006; Tong et al., 2007). The delay has been attributed to the magnitude of the PTME  
36 and the evolution to replace the losses (e.g. Erwin, 2001). Alternatively, persistent  
37 environmental disturbances in the Early Triassic have been held responsible such as  
38 marine anoxia (Algeo et al., 2007, 2008; Bond and Wignall, 2010; Song et al., 2012;  
39 Tian et al., 2014; Clarkson et al., 2016; Lau et al., 2016; Wignall et al., 2016), high  
40 temperatures (Sun et al., 2012; Romano et al., 2013), elevated atmospheric CO<sub>2</sub> (Fraiser  
41 and Bottjer, 2007), and abnormal productivity (Algeo et al., 2011; Meyer et al., 2011;  
42 Grasby et al., 2016). Additionally, the ‘Lilliput effect’ (dwarfism of surviving organisms)  
43 was also pervasive among many clades during the Early Triassic, e.g. foraminifers  
44 (Payne et al., 2011; Song et al., 2011), ostracods (Chu et al., 2015), bivalves (Twitchett,

45 2007), gastropods (Payne, 2005), and brachiopods (He et al., 2015). However, the  
46 spatiotemporal extent and significance of this phenomenon are debated (see e.g.,  
47 Brayard et al., 2010, 2015; Forel and Crasquin, 2015).

48 Contrary to the prolonged recovery scenario, recent studies on conodonts,  
49 ammonoids and foraminifers reveal a fitful recovery pattern that shows diversification  
50 was not underway until the Smithian (Orchard, 2007; Brayard et al., 2009; Song et al.,  
51 2011), less than ~1.5 million years after the PTME. Recovery was then setback by a  
52 severe crisis in the late Smithian (Brayard et al., 2006; Stanley, 2009). Data from Oman  
53 (Krystyn et al., 2003; Twitchett et al., 2004), Italy (Hofmann et al., 2011, 2015; Foster  
54 et al., 2017), the Northern Indian Margin (Brühwiler et al., 2010; Wasmer et al., 2012;  
55 Kaim et al., 2013; Ware et al., 2015;), South Primorye (Shigeta et al., 2009), western  
56 USA (Brayard et al., 2013, 2015, 2017; Hautmann et al., 2013; Hofmann et al., 2013,  
57 2014), South China (Chen et al., 2007; Brayard and Bucher, 2008; Kaim et al., 2010;  
58 Hautmann et al., 2011; 2015; Wang et al., 2017), and Svalbard (Foster et al., 2016) also  
59 documented spatiotemporally variable diversification and occurrences of rather  
60 complex communities. However, few of these works provide exhaustive correlation  
61 between paleontological and environmental data to decipher the recovery course and  
62 its controlling factors. In many regions, marine environments were apparently  
63 depauperate during the Griesbachian-Dienerian with local exceptions (e.g. Twitchett et  
64 al., 2004). Such occurrences are considered to have become more common in the  
65 Spathian (Pietsch and Bottjer, 2014). Thus, the overall spatiotemporal pattern and the  
66 clade variability of the biotic recovery following the PTME are still unclear and

67 therefore controversial, as well its environmental drivers.

68       Sampling efforts and preservation biases influence our knowledge of the biotic  
69 recovery. For instance, Early Triassic silicified fossil assemblages are quite rare, but  
70 show relatively high diversity when present (Foster et al., 2016). Extensive sampling  
71 efforts and new findings might change our understanding of biotic recovery model (e.g.  
72 Brayard et al., 2015, 2017). So far, no hypothesis has addressed the question of why the  
73 re-diversification of a few clades (e.g. ammonoids and conodonts) were more rapid,  
74 while others were rather slow (e.g. brachiopods). Recent studies showed that  
75 environmental conditions (e.g. temperature, oxygen concentration) strongly fluctuated  
76 in the Early Triassic (Algeo et al., 2011; Song et al, 2012; Sun et al, 2012; Grasby et al.,  
77 2013; Tian et al., 2014), but only rarely have studies tried to link such environmental  
78 fluctuations to observed recovery patterns amongst clades (e.g. Pietsch et al. 2014).  
79 South China provides one of the best marine fossil records of the Permian-Triassic  
80 transition, and often serves as an example of the prolonged recovery scenario (e.g.,  
81 Payne et al., 2006; Tong et al., 2007). However, recent reports of unusually diverse  
82 earliest Triassic faunas from Guangxi (Kaim et al. 2010; Hautmann et al. 2011) question  
83 this model. Here we provide new paleontological and paleoenvironmental data from  
84 two Griesbachian-Dienerian sections in Guizhou and Hubei provinces, South China,  
85 that record relatively high diversity levels among several groups and their associated  
86 paleoenvironmental indicators as potential biotic change drivers.

## 87 **GEOLOGICAL SETTING**

88       During the Early Triassic, the South China Block was located in equatorial

89 latitudes, at the interface between Panthalassa and Tethys (Fig. 1A). Marine Lower  
90 Triassic sediments are widespread especially the carbonates of the Yangtze Platform.  
91 This platform was located centrally in the South China Block, and adjacent to the  
92 Nanpanjiang Basin (Fig. 1B). The platform successions are divided into the Daye  
93 Formation, dominated by limestones, and the overlying Jialingjiang Formation,  
94 composed of dolomites (Feng et al., 1997). Lower Triassic strata of the Nanpanjiang  
95 Basin consists of basinal clastic and carbonate rocks plus shallower carbonates of  
96 limited area which formed on isolated platforms (Feng et al., 1997; Bagherpour et al.,  
97 2017). The two studied sections (Gujiao and Jianzishan) were located at the southern  
98 and northern sides of the Yangtze Platform margin, respectively (Fig. 1B).

### 99 **Gujiao section**

100 The Gujiao section (26°30'49.22"N, 106°52'15.17"E) is located ~20 km east from  
101 Guiyang and was situated in the transitional zone between the Nanpanjiang Basin and  
102 the Yangtze Platform during the Early Triassic (Fig. 1B). A new and well-exposed  
103 outcrop was found in 2015 along a newly-built road, close to the Gujiao County,  
104 showing a continuous Permian-Triassic succession (Fig. 2A). The Upper Permian  
105 Changxing Formation is dominated by light gray, bioclastic limestones, and yields  
106 abundant and diverse Permian organisms including brachiopods, gastropods, corals,  
107 dasycladacean algae, and foraminifers. The overlying Dalong Formation consists of  
108 cherty mudstones and black shales, with occasionally volcanic ash beds, representing a  
109 deeper basinal environment. The Dalong Formation contains abundant radiolarian  
110 (Feng and Algeo, 2014; Xiao et al., 2017) and ammonoid (Zheng, 1981) faunas that

111 indicate a late Changhsingian age. The Lower Triassic Daye Formation conformably  
112 overlies the Dalong Formation, and is dominated by marlstones alternating with shales  
113 (Fig. 2B, C). The Permian-Triassic boundary is roughly correlated with the boundary  
114 between the Dalong and Daye formations. Ammonoids are very abundant in the lower  
115 part of Daye Formation. Dai et al. (submitted) identified three ammonoid beds in this  
116 interval, the late Griesbachian *Ophiceras medium* and *Jieshaniceras guizhouense* beds,  
117 and the middle Dienerian *Ambites radiatus* bed. The exact position of the  
118 Griesbachian/Dienerian boundary is not well defined at Gujiao, but is likely just above  
119 the *Jieshaniceras guizhouense* beds.

#### 120 **Jianzishan section**

121 The Jianzishan (30°9'58.08"N, 109°0'27.5"E) section found ~20 km south of  
122 Lichuan in Hubei Province, was located on the north margin of the Yangtze Platform  
123 (Fig. 1B). The Late Permian and Early Triassic are represented by the Changxing and  
124 Daye formations respectively. The former consists of thick-bedded bioclastic  
125 limestones and calcareous sponge reefs with associated shallow-marine fossils (e.g.  
126 corals, calcareous algae and fusulinids). The diverse fauna includes nautiloids,  
127 foraminifers, dasycladacean algae, sponges, corals, and brachiopods (Liu et al., 2017).  
128 The lowermost part of the Daye Formation is a 2.4 m thick microbialite (Fig. 2D), a  
129 typical representative of the extensive microbial deposits found in the lowermost  
130 Triassic strata of South China (e.g. Lehrman, 1999; Yang et al., 2011; Bagherpour et al.,  
131 2017). The overlying strata are dominated by alternations of thin-bedded marlstones  
132 and shales (Fig. 2E) that yield abundant bivalves, brachiopods (Wang et al., 2017) and

133 ammonoids, suggestive of an outer platform facies. Two ammonoid beds, the late  
134 Griesbachian *Ophiceras* sp. indet. and *Jieshaniceras guizhouense* beds are known (Bai  
135 et al., 2017). Four conodont zones, the late Changhsingian *Clarkina changxingensis*  
136 Zone and *Clarkina yini* Zone, the Griesbachian *Hindeodus parvus* Zone and the  
137 *Hindeodus postparvus* Zone, were found in the Jianzishan section (Bai et al., 2017).  
138 Strata between the *H. parvus* and the *H. postparvus* zones are not well dated by  
139 conodonts or ammonoids. Consequently, we here use the presence of abundant  
140 *Sinolingularia* to denote the assemblage at this level.

#### 141 **MATERIAL AND METHODS**

142 Macrofossils (e.g., ammonoids and bivalves) were collected by mechanically  
143 breaking up decimeter-sized blocks. Mechanical techniques, including air and electrical  
144 scribes, were later performed in the laboratory to excavate morphologic details and  
145 thereby facilitate species-level identification. Specimens were photographed using a  
146 Canon 70D camera with a Micro lens EF 100mm f/2.8.

147 A total of 39 and 38 thin sections have been made to study microfossils from the  
148 Gujiao and Jianzishan sections, respectively. In addition, 16 and 18 samples were  
149 respectively collected from the Gujiao and Jianzishan sections for analyses of the size  
150 distribution of pyrite framboids. A polished slab (~1 × 1 cm) of each sample was  
151 examined for size measurements of pyrite framboids using a Scanning Electronic  
152 Microscope (SEM; Hitachi SU8000) at the State Key Laboratory of Biogeology and  
153 Environmental Geology in Wuhan, China. Paleoenvironmental factors have been  
154 investigated including local redox conditions determined using pyrite framboid size

155 analysis, following the approach of Bond and Wignall (2010). A minimum of 100 pyrite  
156 framboids were measured when possible, according to the procedure of Huang et al.,  
157 (2017).

## 158 **PALEONTOLOGICAL RESULTS**

159 Over 3500 specimens were collected from the Gujiao and Jianzishan sections (Figs  
160 3, 4, Tables 1, 2), including ammonoids, nautiloids, bivalves, gastropods, brachiopods,  
161 ostracods, echinoids, foraminifers, and conodonts. Paleozoic holdover and new  
162 originated genus and species (species that belongs to new originated genus) are used  
163 to qualitatively assess the composition of the biotas.

### 164 **Ammonoids**

165 Thirteen ammonoid species occur in the lower Daye Formation at the Gujiao  
166 section, and seven ammonoid species occur in the lower Daye Formation from the  
167 Jianzishan section (Bai et al., 2017). All ammonoid taxa from the two sections are new  
168 originated. At Gujiao, five species (*Ophiceras medium*, Ophiceratidae gen. indet.,  
169 Gyronitidae gen. indet., *Vishnuites pralambha* and ?Mullericeratidae gen. indet.) were  
170 found in the *Ophiceras medium* beds. Five species (*Vishnuites pralambha*, *Ophiceras*  
171 sp. indet., *Mullericeras* sp. nov., *Jieshaniceras guizhouense* and *Proptychites* sp. indet.)  
172 were identified in the *Jieshaniceras guizhouense* beds. Four species (*Ambites radiatus*,  
173 *Pseudoproptychites* cf. *hiemalis*, ?Gyronitidae gen et sp. nov and ?*Ussuridiscus* cf.  
174 *varaha*) occur in the *Ambites radiatus* bed. At Jianzishan, only one taxon was found in  
175 the *Ophiceras* sp. indet. beds. Six species (*Vishnuites pralambha*, *Ussuridiscus varaha*,  
176 *Jieshaniceras guizhouense*, ?Gyronitidae gen. et sp. nov, *Hubeitoceras yanjiaensis* and

177 *Shangganites* sp. indet.) occur in the *Jieshaniceras guizhouense* beds.

178 Two successive ammonoid diversity levels are thus obvious: 1) a relatively high  
179 species richness during the late Griesbachian, and 2) a relatively low species richness  
180 level in the middle Dienerian, with a minimum richness above that. At Jianzishan, no  
181 ammonoids were found in strata overlying the *Jieshaniceras guizhouense* beds.

## 182 **Nautiloids**

183 Nautiloids show a weak diversity level during the late Griesbachian (Fig. 5). Two  
184 new originated nautiloids (*Xiaohenautilus huananensis* and *X. sinensis*) were identified  
185 in the *Ophiceras medium* and *Jieshaniceras guizhouense* beds at the Jianzishan section.  
186 Except for these two species, an unidentified nautiloid was also found in the  
187 *Jieshaniceras guizhouense* beds at the Gujiao section. It mainly differs from  
188 *Xiaohenautilus* by marked ribs on its flanks (Fig. 5). *X. huananensis* and *X. sinensis*  
189 also occur in the *Jieshaniceras guizhouense* beds at Jianzishan.

## 190 **Bivalves**

191 At Gujiao, only three beds yield well preserved bivalves. A bed ~ 20 cm below the  
192 middle Dienerian *Ambites radiatus* bed shows abundant bivalves identified as *Claraia*  
193 *stachei*. The *Ambites radiatus* bed contains moderately abundant *Claraia radialis*, and  
194 the overlying beds have abundant *Claraia aurita* (Fig. 6). These three *Claraia* species  
195 belong to Paleozoic holdovers.

196 Seven bivalve species were found at Jianzishan (Figs. 6, 7): three species (*Claraia*  
197 sp. indet., *Claraia wangi* and *Eumorphotis venetiana*) in the *Sinolingularia* beds, six  
198 species (*C. wangi*, *C. zhenanica*, *E. venetiana*, *Eumorphotis* sp. indet., *Pteria ussurica*

199 *variabilis*, and *Scythentolium scutigerulus*) in the *Ophiceras* sp. indet. beds and two  
200 species (*C.* sp. indet. and *C. zhenanica*) in the *Jieshaniceras guizhouense* beds. Three  
201 of the species are new originated: *Eumorphotis venetiana*, *Eumorphotis* sp. indet. and  
202 *Scythentolium scutigerulus*. The *Ophiceras* sp. indet. beds contain the most diverse and  
203 abundant bivalves.

#### 204 **Gastropods**

205 Gastropods specimens are poorly preserved at both sections (Fig. 8). Thus, their  
206 taxonomic assignment is often tentative. Identifications were mainly based on their  
207 shell height and width, and number of whorls. We also categorized studied specimens  
208 under the following designation ‘unidentified gastropod A, B, C, etc’. to distinguish  
209 them when necessary.

210 At least seven species are present at Jianzishan, two (Bellerophontidae gen. indet.  
211 and unidentified gastropod A) in the *Hindeodus parvus* Zone, one (Bellerophontidae  
212 gen. indet.) in the *Sinolingularia* beds, two (Bellerophontidae gen. indet.  
213 and *?Pseudomurchisonia* sp. indet) in the *Ophiceras* sp. indet. beds and three  
214 (unidentified gastropods A, B and C) in the *Jieshaniceras guizhouense* beds.

215 At Gujiao, four gastropods species were found, three (*?Naticopsis* sp. indet.,  
216 unidentified gastropods D, E) in the *Ophiceras medium* beds, four (*?Naticopsis* sp.  
217 indet., Bellerophontidae gen. indet. and unidentified gastropods D, E) in the  
218 *Jieshaniceras guizhouense* beds, and one (unidentified gastropod E) in the *Ambites*  
219 *radiatus* bed. Gastropods are absent in the overlying beds. In summary, gastropods are  
220 abundant in the late Griesbachian *Ophiceras medium* and *Jieshaniceras guizhouense*

221 beds, but seemingly display low diversity.

## 222 **Brachiopods**

223 Only one brachiopod species (*Lichuanorelloide lichuanensis*) was excavated from  
224 the *Jieshaniceras guizhouense* beds at Gujiao, whilst five brachiopods species  
225 (including three articulated brachiopods) were recognized at Jianzishan. *Sinolingularia*  
226 sp. indet. is very abundant in the *Sinolingularia* beds (Fig. 7). The *Ophiceras* sp. indet.  
227 beds contain two brachiopods species (*Crurithyris* sp. indet. and *Lingularia* sp. indet.,  
228 Fig. 7) and the *Jieshaniceras guizhouense* beds have a relatively diverse brachiopod  
229 fauna with five identified species (*Sinolingularia* sp. indet., *Lingularia* sp. indet.,  
230 *Crurithyris* sp. indet., *Lichuanorelloide lichuanensis* and *Lissorhynchia* sp. indet.).  
231 Among these five species, three (*Sinolingularia* sp. indet., *Lichuanorelloide*  
232 *lichuanensis* and *Lissorhynchia* sp. indet.) are new originated, and suggest a notable  
233 brachiopod diversification after the PTME (Wang et al., 2017).

## 234 **Foraminifers**

235 A total of 20 species were identified in the Gujiao and Jianzishan sections (Fig. 9).  
236 At Gujiao, seven species (*Nodosinelloides* sp. indet., *Gaudryina* sp. indet., *Dentalina*  
237 sp. indet., *Tolypammina* sp. indet., *Glomospira* sp. indet., *Duotaxis* sp. indet.  
238 and *?Dagmarita* sp. indet.) occur in the *Ophiceras medium* beds, and ten  
239 (*Nodosinelloides* sp. indet., *Gaudryina* sp. indet., *Dentalina* sp. indet., *Tolypammina* sp.  
240 indet., *Glomospira* sp. indet., *Duotaxis* sp. indet., *?Dagmarita* sp. indet., *Geinitzina* sp.  
241 indet., *?Vervilleina* sp. indet. and *Nodosaria* sp. indet.) in the *Jieshaniceras guizhouense*  
242 beds. Two species are new originated, *Gaudryina* sp. indet. and *Duotaxis* sp. indet.

243 Overall, foraminiferal species richness was relatively high during the late Griesbachian,  
244 whereas no foraminifer specimens were retrieved from the overlying Dienerian strata.

245 At Jianzishan, 12 species (*Earlandia* sp. indet., *Postcladella kalhori*, *Ammodiscus*  
246 sp. indet., *Dentalina* sp. indet., "*Nodosaria*" sp. indet., *Tezaquina* sp. indet.,  
247 *Heimigordius* sp. indet., *Nodosinelloides* sp. indet., ?*Duotaxis* sp. indet., *Geinitzina* sp.  
248 indet., "*Nodosaria*" *elabugae* and "*Nodosaria*" *skyphica*) were found in the *Hindeodus*  
249 *parvus* Zone, but no foraminifer specimens were found in the *Sinolingularia* beds and  
250 *Ophiceras* sp. indet. beds. Eight species (*Postcladella kalhori*, *Nodosinelloides* sp.  
251 indet., *Geinitzina* sp. indet., *Glomospira* sp. indet., *Tolypammmina* sp. indet., *Froncina*  
252 sp. indet., *Vervilleina* sp. indet. and *Nodosinelloides sagitta*) occur in the *Jieshaniceras*  
253 *guizhouense* beds. All these species belong to Paleozoic holdovers.

## 254 **PYRITE FRAMBOID RESULT AND PALEOREDOX INTERPRETATION.**

### 255 **Gujiao section**

256 At Gujiao, five of the 16 samples exhibit abundant framboids, whereas the rest  
257 contained few or no pyrite framboids (Fig. 10). Framboids in samples GJ-f and GJ-11  
258 display a small mean diameter (3.15  $\mu\text{m}$  and 4.76  $\mu\text{m}$  respectively), as well as small  
259 standard deviations (1.14 and 2.66 respectively; Fig. 11). These measurements suggest  
260 euxinic-anoxic conditions for these beds (Fig. 12), based on the approach of Bond and  
261 Wignall (2010) and Tian et al. (2014). Nine samples from the *Ophiceras medium* and  
262 *Jieshaniceras guizhouense* beds yield few or no framboids. Three of the remaining six  
263 samples from the overlying Dienerian strata show occurrences of framboids (Fig. 10).  
264 Only one sample, (GJ-40) from the *Ambites radiatus* bed, is dominated by small

265 framboids, with a small mean diameter and standard deviation (MD = 4.76  $\mu\text{m}$ , SD =  
266 1.98; Fig. 11), indicating euxinic-anoxic condition (Fig. 12). Above this bed, two  
267 samples (GJ-III-5+0.2 and GJ-III-5+0.9) yield abundant framboids, with a larger mean  
268 diameter, about 5.8  $\mu\text{m}$  (Fig. 11). These plot in the weakly dysoxic (GJ-III-5+0.2) and  
269 anoxic fields (GJ-III-5+0.9) in figure 12 which is calibrated from studies of modern  
270 environments that record a range of redox-related regimes (Bond and Wignall, 2010).

271 The paleoredox trends derived from pyrite framboid size distributions are  
272 congruent with occurrences of trace fossils and the observed succession of facies (Figs  
273 2, 13). Anoxic and dysoxic intervals, inferred from framboid data, coincide with an  
274 absence of bioturbation and correspond to dark, thin-bedded, finely laminated  
275 marlstones interbedded with black shales (Fig. 2B). The beds with rare or no pyrite  
276 framboids are dominated by light-gray, thin bedded marlstones (Fig. 2C) with abundant  
277 vertical tracers *Arenicolites* (Fig. 13A).

### 278 **Jianzishan section**

279 At Jianzishan, only two samples (JZS-2+1.1 and JZS-2+2.1) from the basal  
280 Triassic microbialites yield moderately abundant, large pyrite framboids, whereas  
281 framboids were rare or absent in other samples (Fig. 10). Both framboid populations  
282 exhibit diameters and standard deviations that plot in dysoxic fields (Fig. 12). Other  
283 samples lack pyrite framboids and likely indicate oxic conditions during deposition of  
284 the *Sinolingularia*, *Ophiceras* sp. indet. and *Jieshaniceras guizhouense* beds.  
285 Moderately abundant vertical ichnofossils (e.g. *Skolithos*) occur in the *Jieshaniceras*  
286 *guizhouense* beds (Fig. 13B), also suggesting well-oxygenated conditions.

287 **DISCUSSION**

288 **Late Griesbachian rapid recovery**

289 Previous works in South China support the claims of a delayed recovery in China  
290 caused by persisting environmental stresses up until at least the Spathian (e.g. Payne et  
291 al., 2006; Tong et al., 2007; Sun et al., 2012; Song et al., 2012). Chen et al. (2007)  
292 documented a rapid onset of recovery at Meishan section, however, only nine species  
293 of macrofossils were found during late Griesbachian. Our data show that moderately  
294 diverse marine communities flourished soon after the PTME, during the late  
295 Griesbachian, at least in well-oxygenated environments. Rarefaction analyses indicate  
296 that the highest diversity is in the *Jieshaniceras guizhouense* beds both at Gujiao and  
297 Jianzishan (Fig. 14). Furthermore, the microbialite community (early Griesbachian) is  
298 rather diverse at Jianzishan that likely results from the non-lethal oxygen poor  
299 conditions and supports the microbialites refuge model (Forel et al., 2013). At Gujiao,  
300 the early Griesbachian witnessed depressed marine community under anoxic-euxinic  
301 conditions in the aftermath of PTME. The diversity in the *Jieshaniceras guizhouense*  
302 beds at Gujiao is lower than that at Jianzishan, 26 and 37 species at Gujiao and  
303 Jianzishan respectively, supporting that redox condition plays a significant role in  
304 diversity rebound. There is no marked shift in facies observed near the  
305 Griesbachian/Dienerian boundary at the both sections indicating that the dramatic  
306 diversity reduction after the late Griesbachian is not a bias resulting from major facies  
307 change. However, the causation of this small crisis still needs more work to decipher.

308 At Jianzishan, a marked Paleozoic holdovers diversity drop is observed between

309 the *Hindeodus parvus* Zone and *Sinolingularia* beds (Figs 4, 15), possibly reflecting  
310 the second pulse of the PTME in the earliest Triassic mass extinction (Song et al., 2013).  
311 Alternatively (or additionally), this diversity loss might be related to the facies shift  
312 between the microbialites and mudstones (Fig. 4). High diversity levels of Paleozoic  
313 holdovers and new originated taxa occurred during the late Griesbachian in both  
314 sections. Additionally, during the late Griesbachian, nektonic taxa show a more marked  
315 rebound than benthic organisms, e.g. ammonoid (proptychitids). This difference  
316 between nektonic and benthic taxa may result from the relatively stronger mobility of  
317 nektonic organisms such as cephalopods, which makes them better able to avoid  
318 spatially variable hostile environments (Bambach et al., 2002). However, it could also  
319 reflect the intrinsically evolutionary rate of ammonoids that tends to be high, compared  
320 to other mollusk groups, at all times during the clades' history (Stanley, 2009).

321 Well-oxygenated conditions during the late Griesbachian have rarely been  
322 documented in South China, and the interval is widely reported to be poorly ventilated  
323 in marine settings (Song et al., 2012; Tian et al., 2014; Huang et al., 2017; Li et al.,  
324 2016). Our data reveal that there were heterogeneous redox conditions at this time.  
325 Gujiao and Jianzishan sections were located on the platform margin, a predicted  
326 "Refuge Zone" sandwiched beneath the potentially warm surface waters and anoxic  
327 deeper waters (Song et al. 2014; Godbold et al., 2017). Our results suggest that diversity  
328 patterns in the Early Triassic were strongly controlled by the presence of anoxia, with  
329 recovery best seen in the limited areas of more ventilated conditions (Fig. 15).

330 Similar high diversity levels rapidly after the PTME are also known from a limited

331 number of other locations. For example, a late Griesbachian fauna with ammonoids,  
332 bivalves and gastropods in oxic strata is seen in Oman (Krystyn et al., 2003; Twitchett  
333 et al., 2004). The overall diversity during Griesbachian and Dienerian at Meishan  
334 section is relative high, probably owing to its overmuch studies, especially near the  
335 Permian-Triassic boundary. In addition, the diverse conodont fauna contributes the  
336 most of overall diversity (Zhang et al., 2007). However, the low diversity of mollusk  
337 taxa and oxygen poor conditions during the late Griesbachian at Meishan also support  
338 that redox condition is a key factor in controlling biotic recovery (Chen et al., 2007,  
339 2014; Li et al., 2016). The Siusi Member (late Griesbachian-Dienerian) of the Werfen  
340 Formation (Dolomites, Italy) also yields relatively diverse mollusk communities in well  
341 oxygenated environment (Wignall and Twitchett, 1996; Hofmann et al., 2015; Foster et  
342 al., 2017). A marked ammonoid diversity richness has been documented in the Northern  
343 Indian Margin (Salt Range and Spiti) during the early Dienerian (Ware et al., 2015),  
344 and this interval was proved to be well oxygenated (Hermann et al., 2011). Combination  
345 of the works above and present work support the heterogeneous recovery pattern and  
346 rapid diversification only present under oxic conditions.

347         Several studies described diverse mollusk faunas from the earliest Triassic  
348 microbialites of South China (Kaim et al., 2010; Hautmann et al., 2011, 2015). The  
349 microbialites may have serve as a refuge for benthic organisms (e.g., Forel et al., 2013),  
350 but reconstructions of paleoredox conditions within these deposits suggests variety of  
351 redox conditions (e.g. Forel et al., 2009, 2013; Liao et al., 2017). However, these  
352 microbialites occur between the two phases of the PTME (Jiang et al., 2014; Brosse et

353 al., 2015) that straddle the Permian-Triassic boundary (Song et al., 2013), and thus  
354 record intra-extinction conditions not those during the recovery. It is noteworthy that  
355 the most microbialite taxa are Paleozoic holdovers (Hautmann et al., 2015), and so just  
356 record assemblages that survived the first (but not the second) stage of the PTME.

357 In South China, biotic recovery was impeded by anoxic events later in the  
358 Dienerian (Figs. 10, 15), as also seen in other basins, e.g. Pakistan (Hermann et al.,  
359 2011) and northern Italy (Foster et al., 2017). Overall, previous works and our results  
360 suggest that recovery after the PTME was spatiotemporally heterogeneous and highly  
361 dependent on local environmental conditions as seen during the late Griesbachian in  
362 our two studied sections with same sample effort and similar preservation conditions in  
363 South China. The most abundant and diversified Griesbachian communities were to be  
364 in shallower (but not the shallowest) and more oxic settings.

### 365 **Fitful recovery pattern**

366 Based on our new data and previous studies, a fitful recovery pattern from the  
367 PTME can be simplified into three distinct phases of relatively high diversity that vary  
368 in time and space according to regional and local environmental conditions: I. late  
369 Griesbachian-early Dienerian; II. early-middle Smithian and III. Spathian. During  
370 phase I, new originated genus became the dominant group in South China, especially  
371 amongst the ammonoid-dominated nektonic communities. This recovery phase was  
372 short-lived and probably curtailed by the return of anoxic conditions during the  
373 Dienerian. Phase II during the early-middle Smithian is well documented by further  
374 recovery seen among ammonoids (Brayard et al., 2009; Brühwiler et al., 2010),

375 conodonts (Orchard, 2007) and foraminifers (Song et al., 2011). This time interval also  
376 witnessed the local development of rather diverse community (Haig et al., 2015) and  
377 sponge bioconstructions (Brayard et al., 2011) and the presence of large-sized  
378 organisms (Brayard et al., 2010, 2015). It was associated with a relatively cooling  
379 interval (Sun et al., 2012; Romano et al., 2013) and well-oxygenated conditions  
380 (Galfetti et al., 2008; Grasby et al., 2013). This phase of diversification was followed  
381 by a severe extinction in the late Smithian (Orchard, 2007; Brayard et al., 2009; Stanley,  
382 2009), seemingly related to high temperatures and anoxia (Sun et al., 2012; Song et al.,  
383 2012; Grasby et al., 2013). At that time, foraminifers only show a diversity low point,  
384 but no real extinction (Song et al., 2011). The Phase III diversification occurred rapidly  
385 after the late Smithian event and is seen among both nektonic and benthic forms  
386 (Brayard et al., 2009, 2017; Stanley, 2009; Song et al., 2011; Chen et al., 2015; Hautmann  
387 et al., 2013). It also witnessed the appearance of top-level predators in the fossil record  
388 (e.g. Scheyer et al. 2014; Motani et al., 2015). The Phase III diversification is associated  
389 with a cooling trend (Sun et al., 2012; Romano et al., 2013) and with the end of anoxic  
390 events characterized by the widespread occurrence of marine red beds (Song et al.,  
391 2017).

392 As a whole, regional environmental fluctuations controlled the recovery in space  
393 and time. The fitful recovery model gets more and more strong evidences than delayed  
394 claims, at the same time, the deteriorative environment conditions are clarified to be  
395 recurrent (e.g. Sun et al., 2012; Song et al., 2012; Grasby et al., 2013; Huang et al.,  
396 2017). The differences between benthic and nektonic/pelagic organisms recovery

397 patterns and underlying processes are still unclear, but are probably related to  
398 physiology variations (Bambach et al., 2002), intrinsic evolutionary rates (Stanley,  
399 2009), and biological adaptation and competition rates (Hautmann et al., 2015). More  
400 exhaustive integrated works (paleontological and environmental) within a high-time  
401 resolution are necessary to decipher spatiotemporally heterogeneous recovery patterns  
402 among different lineages.

### 403 **CONCLUSION**

404 New evidence from two sections in South China has revealed relatively high  
405 diversity levels of ammonoids, bivalves, brachiopods, conodonts, foraminifers and  
406 gastropods during the late Griesbachian. These high diversity levels are associated with  
407 unusually well-oxygenated conditions for this period. The nektonic taxa show more  
408 stronger diversity rebound than benthic forms, possibly due to their ecology (tiering,  
409 stronger mobility) or evolutionary factors (ammonoids show much higher origination  
410 rates than benthic mollusks (Brayard et al., 2009)). Our results show a spatiotemporally  
411 heterogeneous recovery pattern linked to redox conditions, in both benthic and nektonic  
412 communities in the investigated sections in South China. In contrast to the notion of  
413 persistently harsh environmental conditions during the Early Triassic, our results  
414 suggest that there was a regionally variable recovery pattern after the PTME, linked to  
415 amelioration of environmental stresses, with three main high diversity phases during  
416 the late Griesbachian-early Dienerian, early-middle Smithian and Spathian.

### 417 **ACKNOWLEDGMENTS**

418 We thank Arnaud Brayard for participating in the determination of ammonoids,

419 and discussion of biostratigraphy and interpretations, Daoliang Chu for helpful  
420 discussions and Yunfei Huang for suggestions on the identification of bivalves. We  
421 appreciate comments from Ganqing Jiang, Guang R. Shi, and two anonymous  
422 reviewers that have greatly improved the quality of this paper. This study is supported  
423 by the National Natural Science Foundation of China (41622207, 41672016, 41530104,  
424 41661134047) and the 111 Project (B08030). This is a contribution to the IGCP-630.

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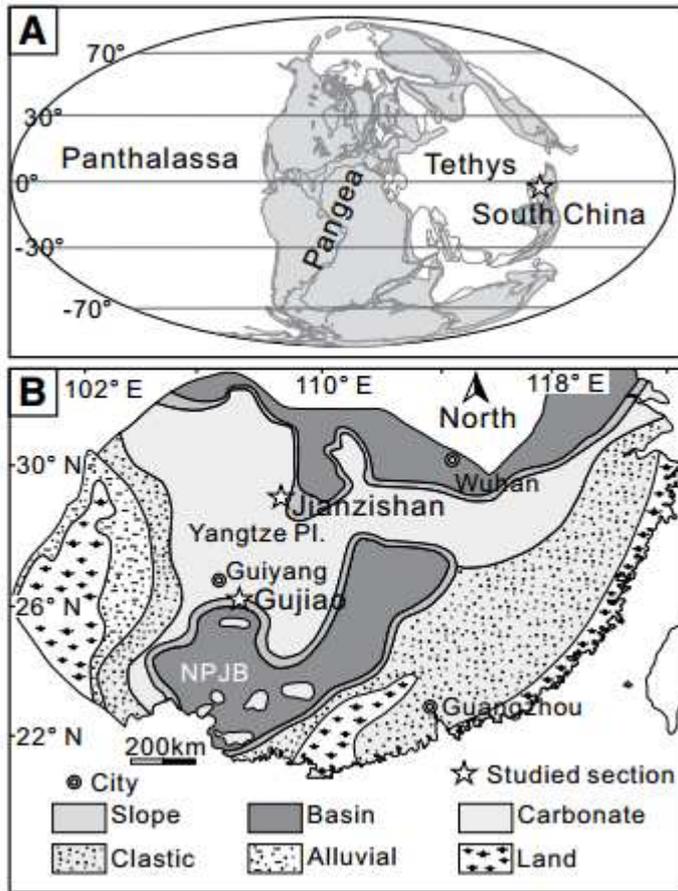
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822

823 **FIGURE CAPTIONS**

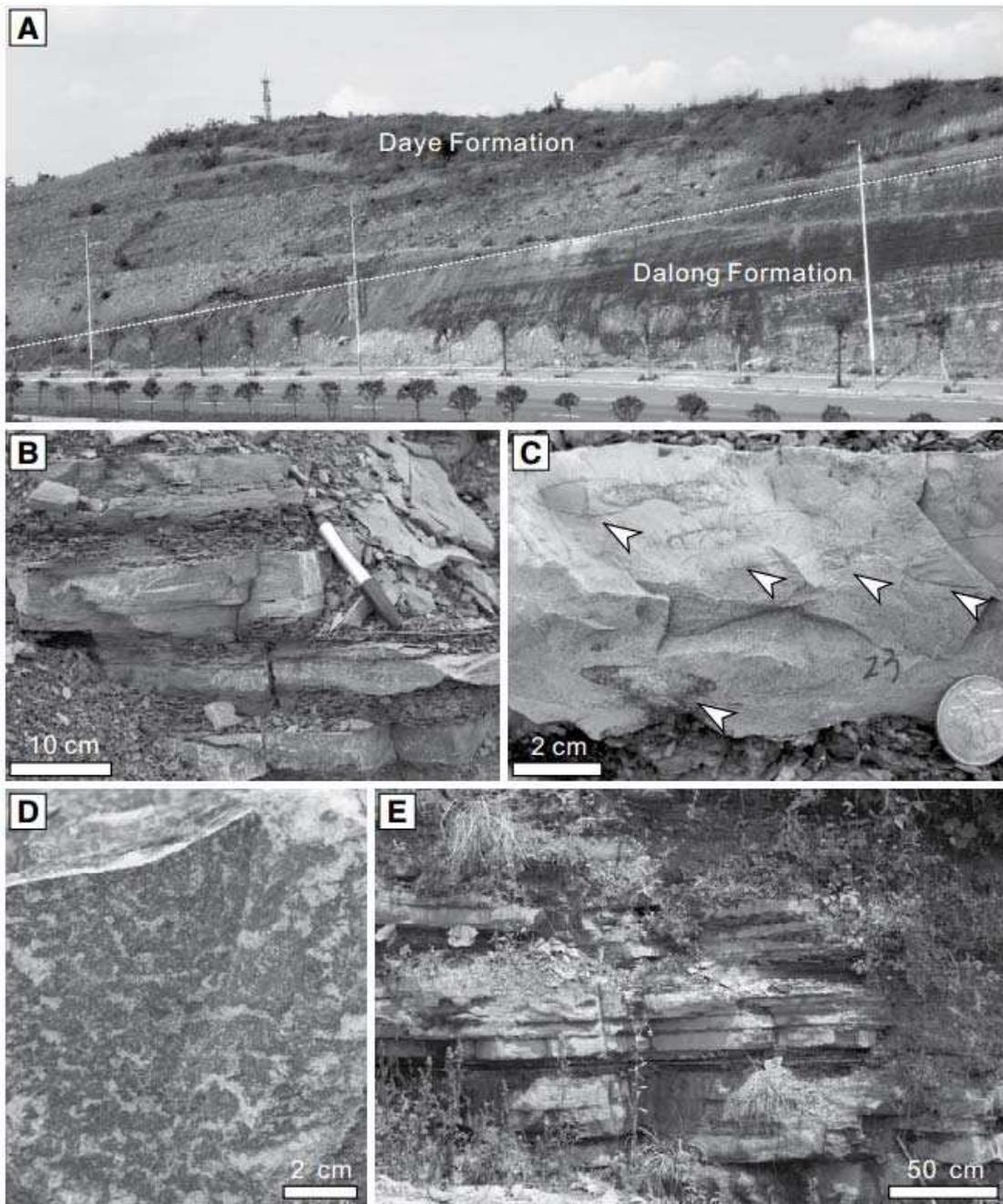


824

825 Figure 1. A. Early Triassic paleogeography, modified from Scotese (2001). B. Early

826 Triassic paleogeographic map of South China, modified from Feng et al. (1997). PL.

827 platform; NPJB. Nanpanjiang Basin.



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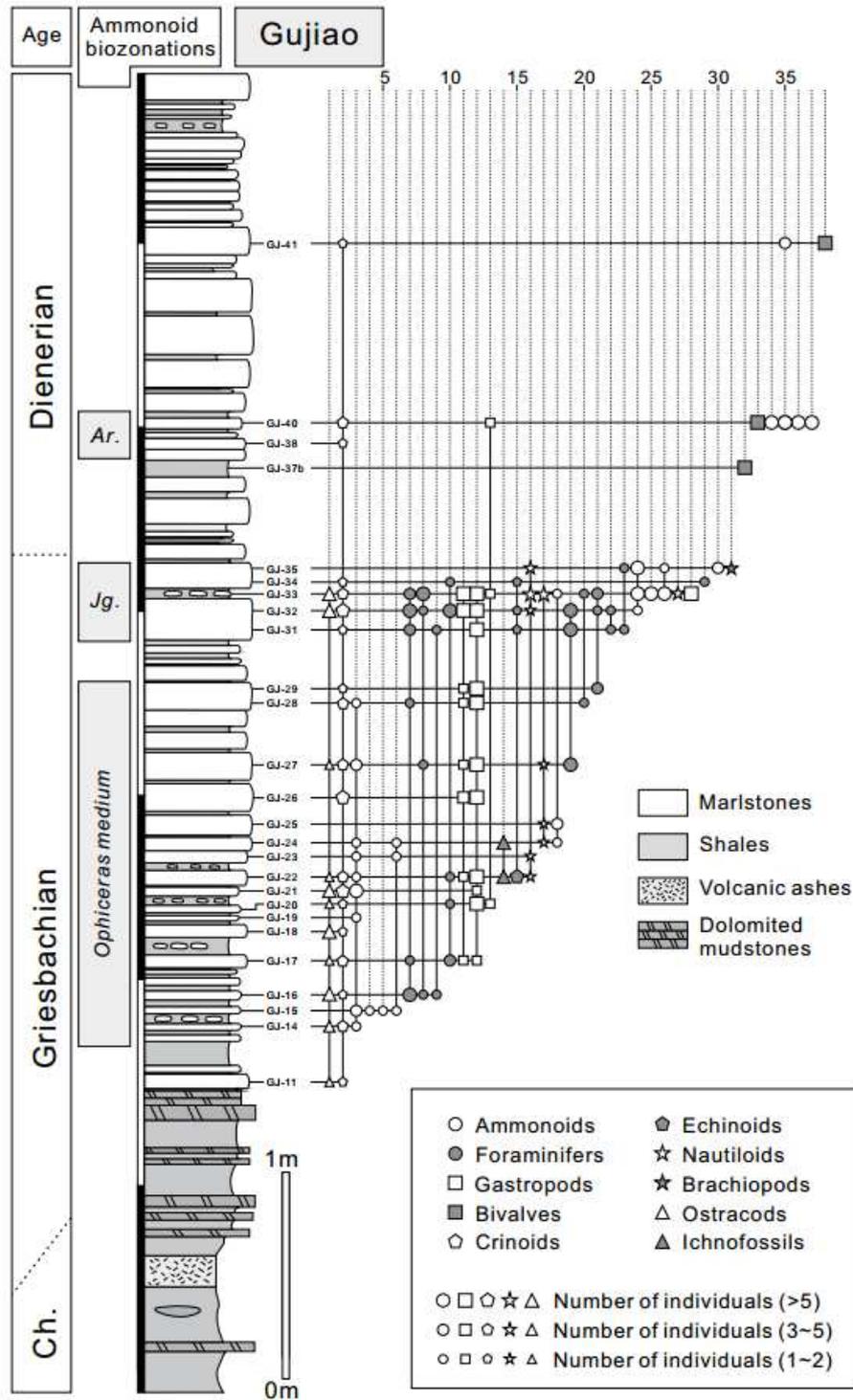
829 Figure 2. A. Landscape view of the Gujiao section. B. Laminated marlstones alternated

830 with black shales of the Daye Formation at Gujiao. C. Light gray marlstones with

831 abundant ammonoids (indicated by white arrows). D. Microbialites from the lower

832 most of the Daye Formation at Jianzishan. E. Mudstones alternating with shales

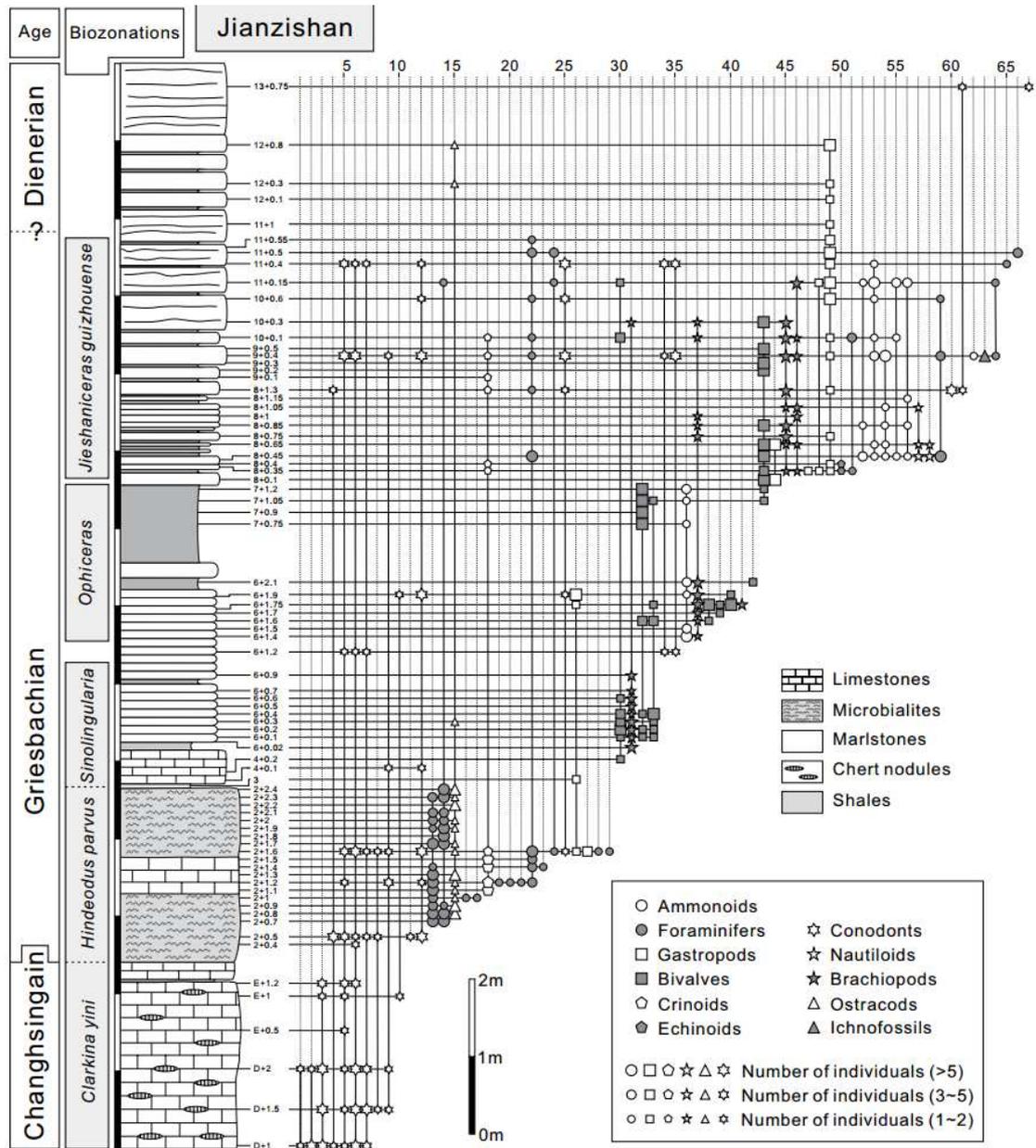
833 overlying the microbialites at Jianzishan.



834

835 Figure 3. Fossil occurrences in the Gujiao section. See Table 1 for fossil list. Ch.

836 Changhsingian; *Ar. Ambites radiatus* bed; *Jg. Jieshaniceras guizhouense* beds.

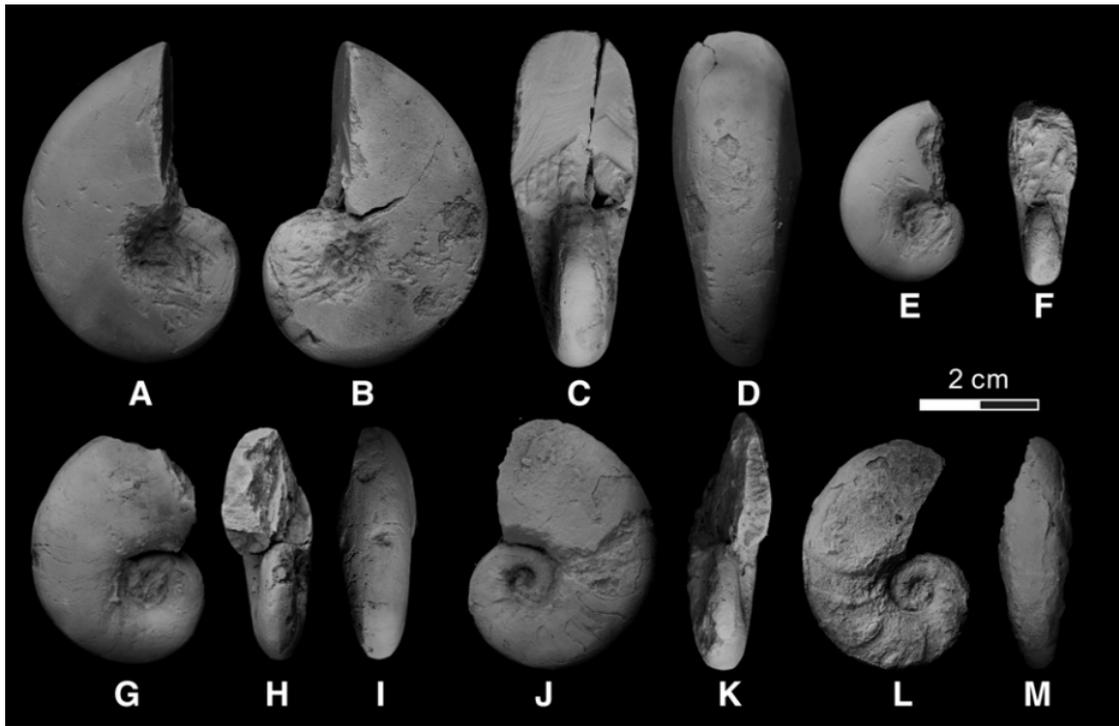


837

838 Figure 4. Fossil occurrences in the Jianzishan section. Conodont data and part of

839 ammonoid data are from Bai et al. (2017). Brachiopod data are from Wang et al. (2017).

840 See Table 2 for fossil list.

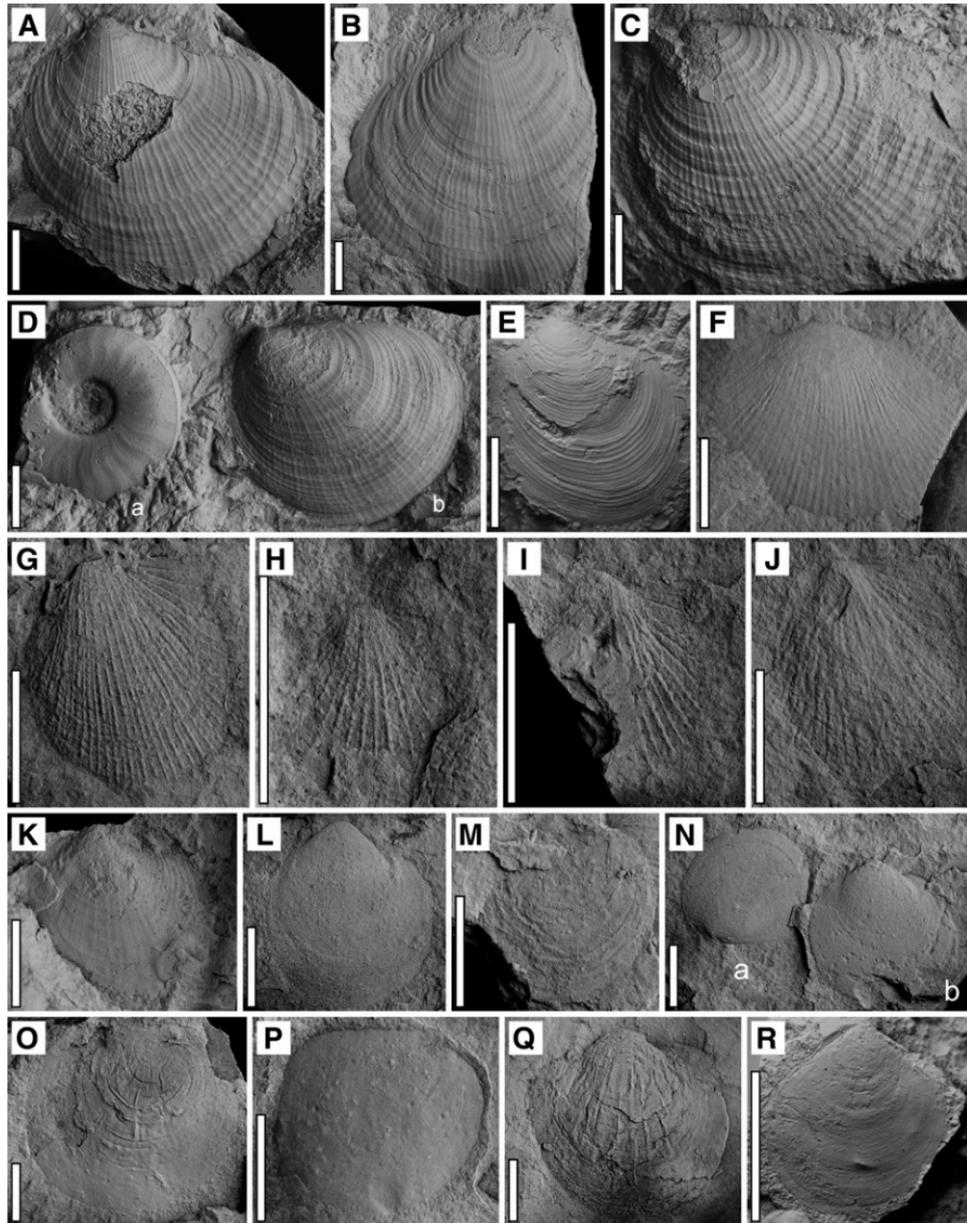


841

842 Figure 5. Nautiloids from bed GJ-33, Gujiao section. A-D. *Xiaohenautilus sinensis* Xu,

843 1988; E-F. *Xiaohenautilus sinensis* Xu, 1988; G-I. *Xiaohenautilus huananensis* Xu,

844 1988; J-K. *Xiaohenautilus huananensis* Xu, 1988; L M. undetermined nautiloid.



845

846 Figure 6. Mollusks from the Gujiao and Jianzishan sections. Scale bar is 1 cm. A-C.

847 *Claraia radialis* (Leonardi, 1935), A, left valve, from GJ-40, B, right valve, from GJ-

848 40, C, left valve, from GJ-40; D. a. *Ambites radiatus* (Brühwiler, Brayard, Bucher and

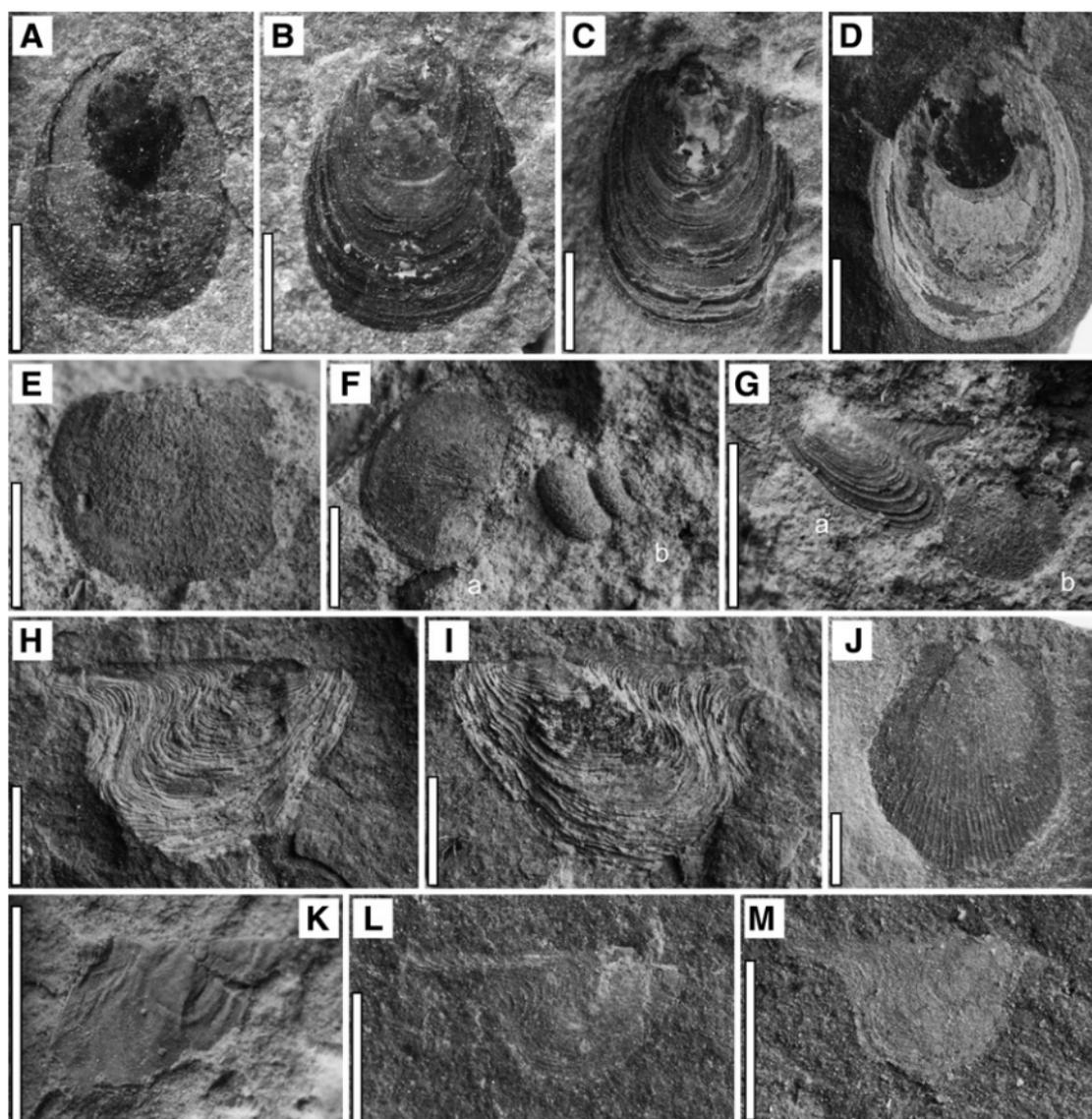
849 Kuang, 2008), b. *Claraia radialis* (Leonardi, 1935), left valve, from GJ-40; E. *Claraia*

850 *aurita* (von Hauer, 1850), left valve, from GJ-41; F. *Claraia stachei* (Patte, 1935), left

851 valve, from GJ-37b; G-J. *Eumorphotis venetiana* (von Hauer, 1850), G, left valve, from

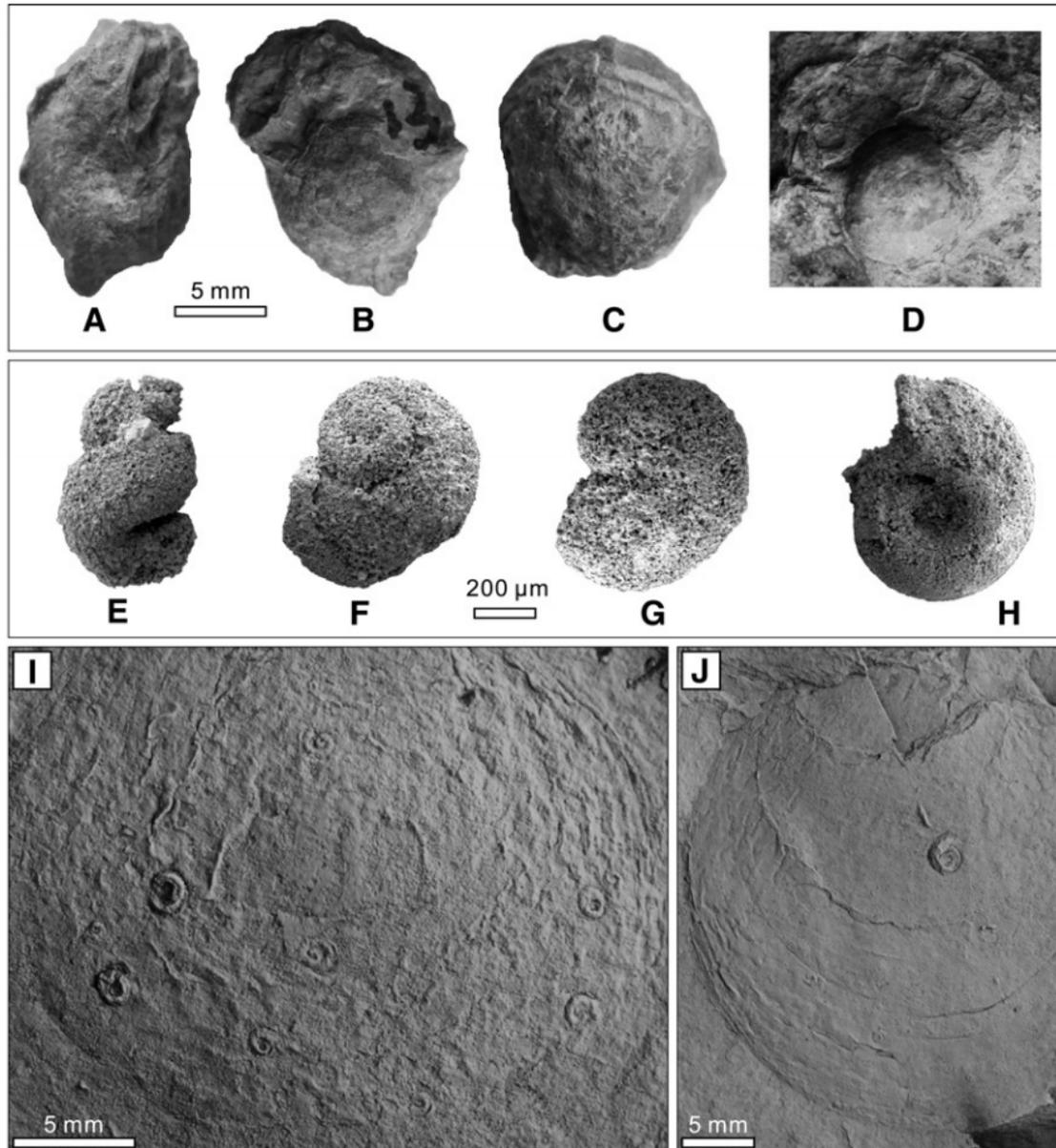
852 JZS-6+0.3, H, left valve, from JZS-6+0.35, I, left valve, from JZS-6+0.3, J, left valve,

853 from JZS-6+0.3; K-N. *Claraia wangi* (Patte, 1935), K, left valve, from JZS-6+0.45, L,  
 854 left valve, from JZS- 7+0.75, M, left valve, from JZS-7+1.05, N, a. left valve, b. right  
 855 value, from JZS-7+1.2; O-Q. *Claraia zhenanica* Chen and Liu in Liu, 1964, O, right  
 856 valve, from JZS-8+0.65, P, right valve, from JZS-8+0.65, Q, left valve, from JZS-9+0.2;  
 857 R. *Scythentolium scutigerulus* Hautmann et al., 2011, left valve, From JZS-6+1.9.



858  
 859 Figure 7. Brachiopods, bivalves and gastropods from the Jianzishan section. The scale  
 860 bar is 2 mm. A-D. *Sinolingularia* sp. indet.; A, from JZS-6+0.02; B, from JZS-6+0.3;  
 861 C, from JZS- 6+0.9; D, from JZS-10+0.3; E. *Crurithyris* sp. indet., from JZS-6+1.9; F.

862 a. *Crurithyris* sp. indet., b. ?*Pseudomurchisonia* sp. indet., from JZS-6+1.9; G. a. *Pteria*  
863 *ussurica variabilis* Chen and Lan in Gu et al, 1976, left valve; b. *Crurithyris* sp. indet.,  
864 from JZS-6+1.75; H. *Pteria ussurica variabilis* Chen and Lan in Gu et al, 1976, left  
865 valve, external mold, from JZS-6+1.75; I. *Pteria ussurica variabilis* Chen and Lan in  
866 Gu et al, 1976, left valve, from JZS-6+1.75; J. *Eumorphotis* sp. indet., left valve, from  
867 JZS-6+1.75; K. *Eumorphotis* sp. indet., right valve, from JZS-6+1.7; L. *Pteria ussurica*  
868 *variabilis* Chen and Lan in Gu et al, 1976, right valve, from JZS-6+1.75; M. *Pteria*  
869 *ussurica variabilis* Chen and Lan in Gu et al, 1976, right valve, from JZS-6+1.75.



870

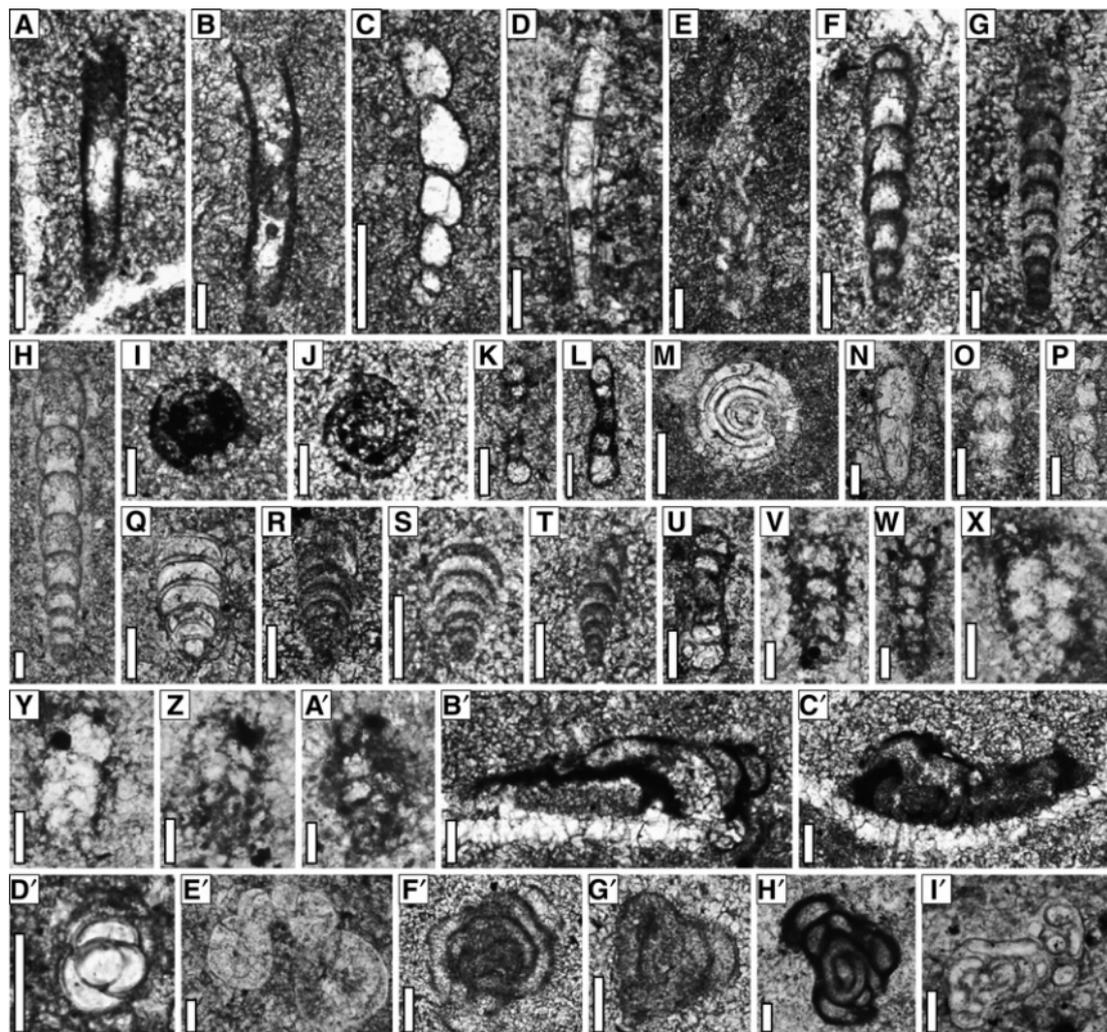
871 Figure 8. Gastropods and microconchids from the Gujiao and Jianzishan sections. A-D.

872 Bellerophontidae gen. indet., from bed GJ-33; E. undetermined Gastropod A., from bed

873 JZS-6+1.9; F-G. ?*Naticopsis* sp. indet., from bed JZS-6+1.9; H. Bellerophontidae gen.

874 indet., from bed JZS-6+1.9. I-J, epizoan microconchids on *Claraia* shells, from bed

875 JZS-8+0.65.



876

877 Figure 9. Foraminifera from the Gujiao and Jianzishan sections. The scale bar is 50  $\mu\text{m}$ .

878 A-B. *Earlandia* sp. indet., from JZS-2+0.7; C. *Dentalina* sp. indet., from JZS-2+1; D.

879 *Tezaquina* sp. indet., from JZS-2+1.2; E. *Vervilleina* sp., indet., from JZS-11+0.4; F-H.

880 *Nodosinelloides sagitta* (Miklukho-Maklay, 1954), F-G, from JZS-11+0.5, H, from GJ-

881 28; I-L, *Postcladella kalhori* (Brönnimann, Zaninetti and Bozorgnia, 1972), I-J, from

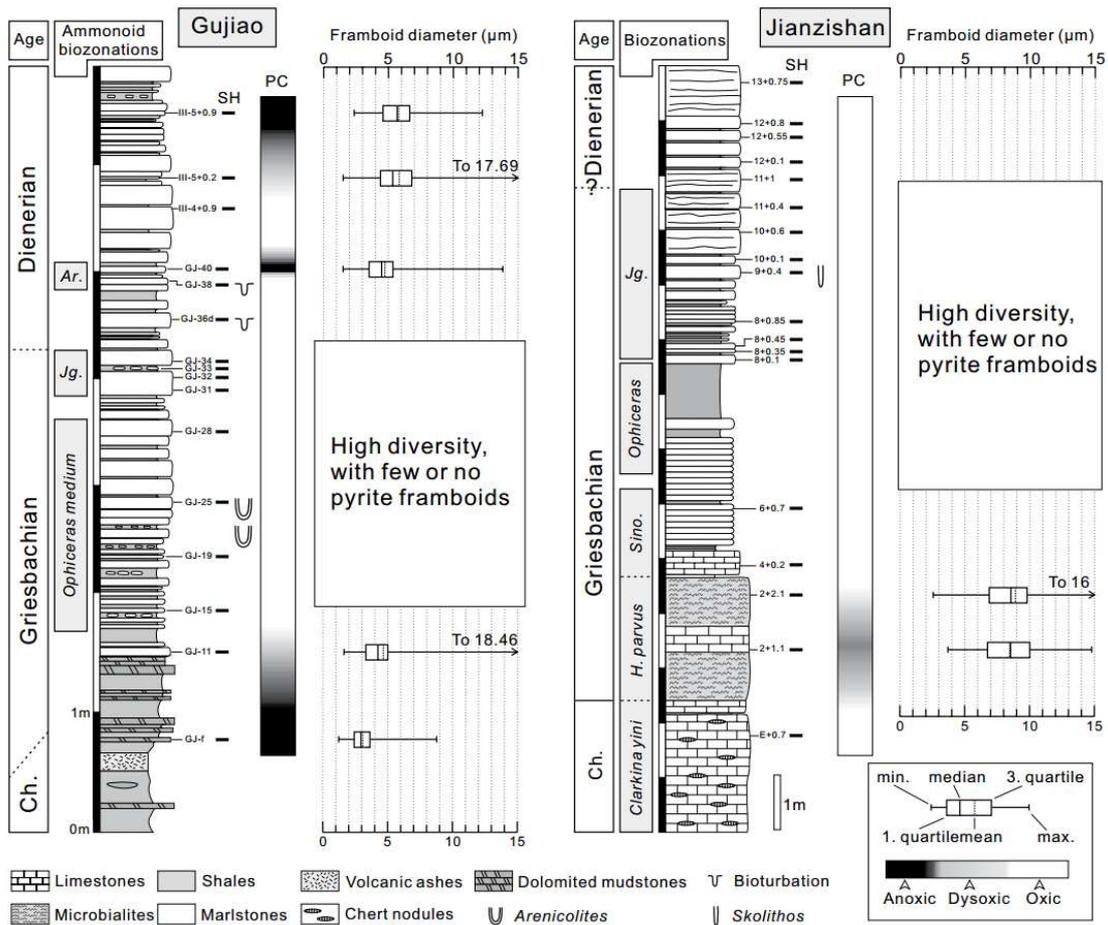
882 JZS-2+0.7, K-L from JZS-2+2.3; M. *Ammodiscus* sp. indet., from JZS-2+1; N.

883 “*Nodosaria*” sp. indet., from JZS-2+1.2; O. “*Nodosaria*” *elabugae* Cherdyntsev, 1914,

884 from JZS-2+1.6; P. “*Nodosaria*” *skyphica* Efimova, 1974, from JZS-2+1.6; Q-S.

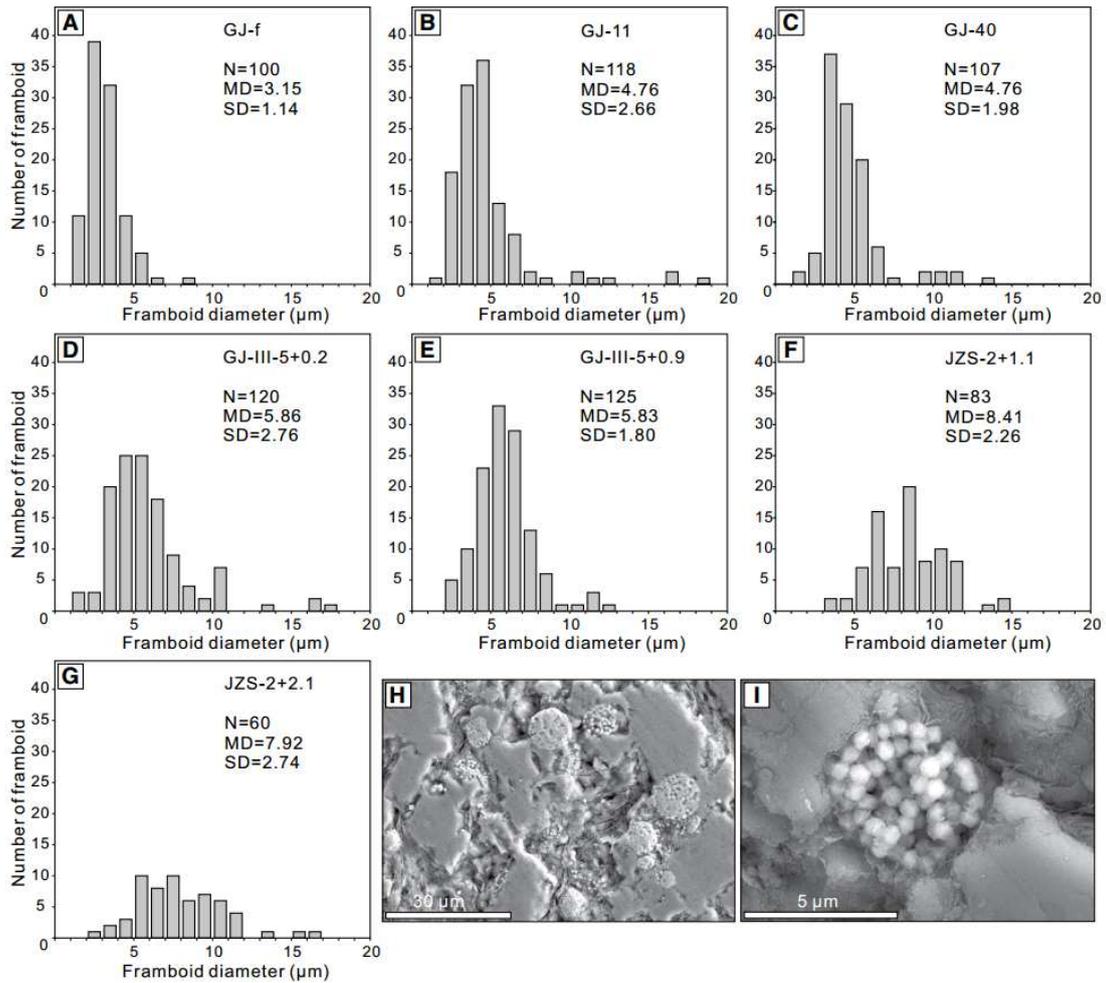
885 *Geinitzina* sp. indet., Q, from JZS-2+1.6, R-S, from JZS-11+0.4; T. *Fronidina* sp. indet.,

886 from JZS-9+0.4; U. *Hemigordius* sp. indet., from JZS-2+1.2; V-W. *Dagmarita* sp.  
 887 indet., V, from GJ-29, W, from GJ-33; X-A'. *Gaudryina* sp. indet., X-Z, from GJ-32, A',  
 888 from GJ-33; B'-C'. *Tolypamina* sp. indet., from JZS-9+0.4; D'. *Duotaxis* sp. indet.,  
 889 from JZS-2+1.4; E'. *Duotaxis* sp. indet., from GJ-28; F'-I'. *Glomospira* sp. indet., F'-  
 890 G', from JZS-10+0.1, H' from JZS-8+0.35, I' from GJ-31.



891

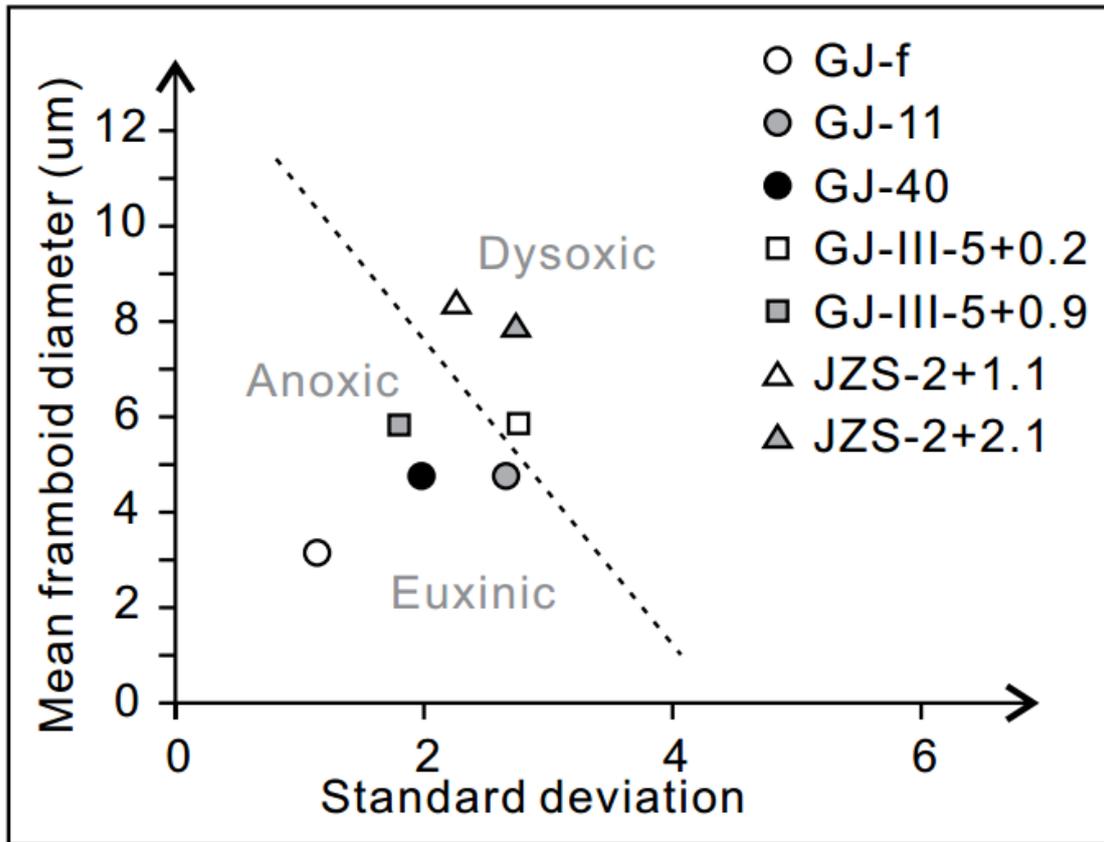
892 Figure 10. Griesbachian-Dienerian paleoredox conditions of the Gujiao and Jianzishan  
 893 sections. Only five samples yield pyrite frambooids at Gujiao, and two at Jianzishan. SH:  
 894 pyrite sample horizon. PC: Paleoredox conditions. Biozonations abbreviation. Jg.  
 895 *Jieshaniceras guizhouense* beds; Ar. *Ambites radiatus* bed; H. *Hindeodus*; Sino.  
 896 *Sinolingularia* beds.



897

898 Figure 11. A-G. Size distributions of pyrite framboids. H-I. SEM photos of typical

899 pyrite framboids, from sample GJ-III- 5+0.9.

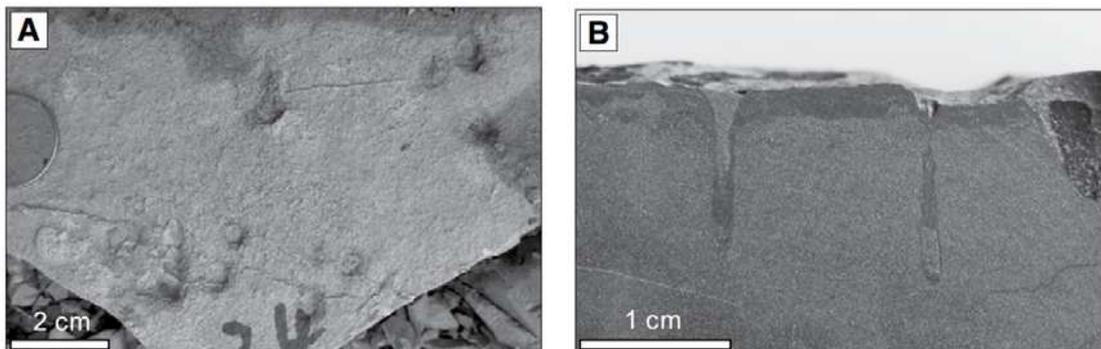


900

901 Figure 12. Mean diameter versus standard deviation of pyrite framboids. The dotted

902 line separating euxinic/anoxic from dysoxic facies is from Bond and Wignall (2010)

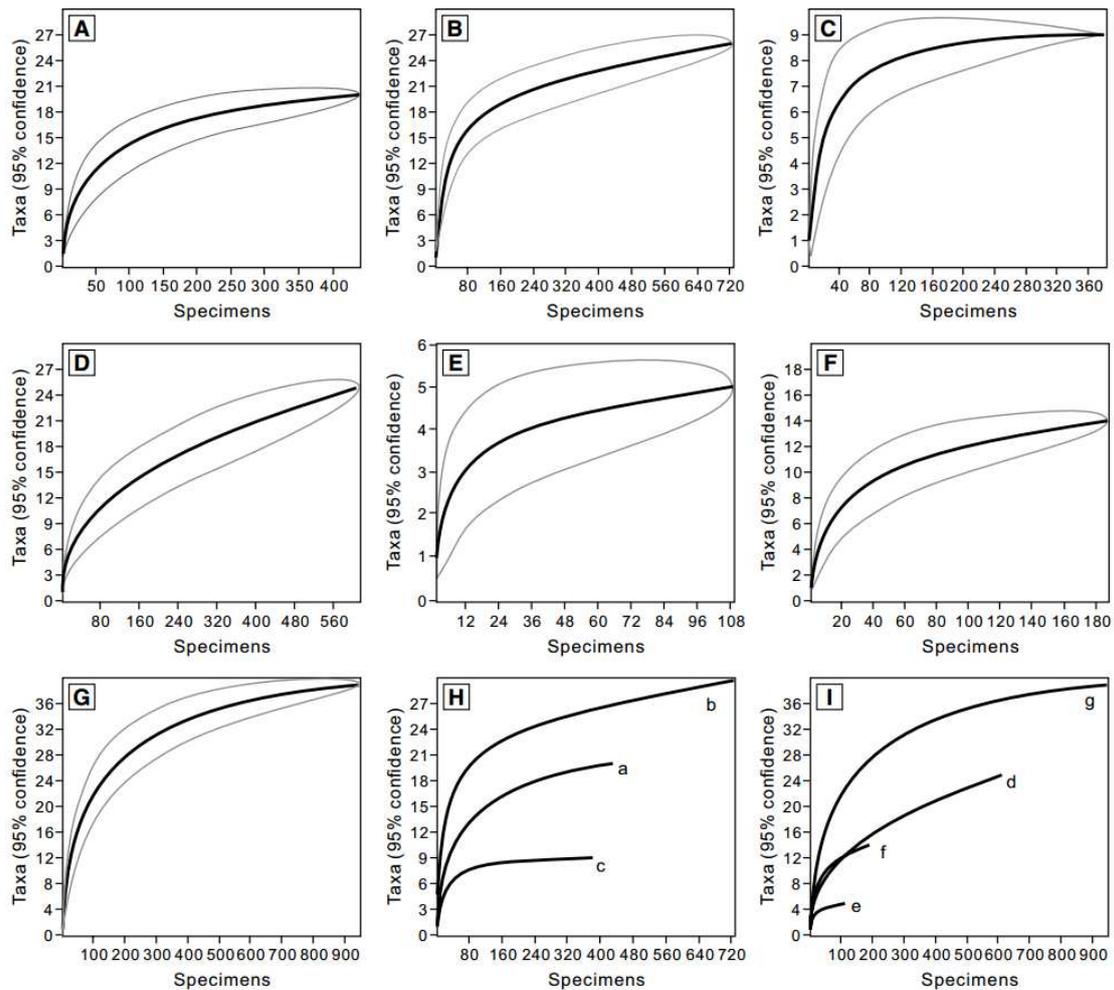
903 and Tian et al., (2014) and is derived from measurements in modern environments.



904

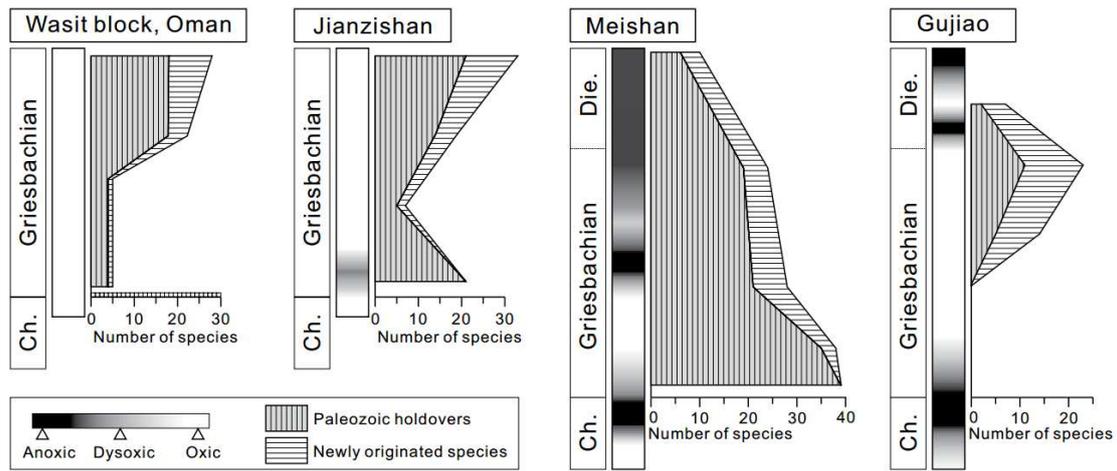
905 Figure 13. Ichnofossils from the Gujiao and Jianzishan sections. A, *Arenicolites* isp.,

906 from bed GJ-24. B, *Skolithos* isp., from bed JZS-9+0.4.



907

908 Figure 14. Rarefaction analyses for each analyzed zone or beds. The grey lines indicate  
 909 95% confidence intervals. The *Jieshaniceras guizhouense* beds in both sections contain  
 910 the highest taxonomic richness. A, *Ophiceras medium* beds, Gujiao section. B,  
 911 *Jieshaniceras guizhouense* beds, Gujiao section. C, *Ambites radiatus* bed, Gujiao  
 912 section. D, *Hindeodus parvus* Zone, Jianzishan section. E, *Sinolingularia* beds,  
 913 Jianzishan section. F, *Ophiceras medium* beds, Jianzishan section. G, *Jieshaniceras*  
 914 *guizhouense* beds, Jianzishan section. H, rarefaction for each bed at Gujiao, a,  
 915 *Ophiceras medium* beds; b, *Jieshaniceras guizhouense* beds; c, *Ambites radiatus* bed.  
 916 I, rarefaction for each biozone/assemblage at Jianzishan, d, *Hindeodus parvus* Zone, e,  
 917 *Sinolingularia* beds, f, *Ophiceras medium* beds, g, *Jieshaniceras guizhouense* beds.



918

919 Figure 15. Correlation of paleoredox conditions and taxonomic richness of Paleozoic

920 holdovers and new originated species. The redox conditions of Wasit block, Oman is

921 from Clarkson et al. (2016) and the paleontological data are from Twitchett et al. (2004).

922 The redox conditions of Meishan is from Chen et al. (2014) and Li et al. (2016) and the

923 paleontological data is from Chen et al. (2007) and Song et al. (2013). Ch.

924 Changhsingian; Die. Dienerian.

925