

1 **A nearshore Hirnantian brachiopod fauna from South China and**
2 **its ecological significance**

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16 **Running Header:** A nearshore Hirnantian brachiopod fauna from S China

17
18 **Abstract.**—The brachiopods collected from the Kuanyinchiao Beds (Hirnantian, uppermost
19 Ordovician) in Meitan and Zunyi counties, northern Guizhou include 13 species and one
20 undetermined taxon, dominated by *Hirnantia sagittifera* (M’Coy) accounting for over one-
21 third of the specimens, together with common *Eostropheodonta hirnantensis* (M’Coy). They
22 are assigned to the *Hirnantia-Eostropheodonta* Community which probably inhabited a

23 shallow-water, near-shore BA2 to upper BA3 environment. Population analysis shows that
24 the community was well adapted to this environment after the first phase of the end
25 Ordovician mass extinction. Representative specimens of all the species are illustrated, and a
26 new species, *Minutomena missa* n. sp. is described herein. The variation in *Hirnantia*
27 *sagittifera* (M'Coy) was noted in many of previous studies but was not statistically
28 evidenced. Here we have measured representative specimens of that famous species from the
29 major paleoplates and terranes in the world, along with other species assigned to the genus
30 from South China. Having used Principal Component Analysis (PCA), significant variations
31 in the species are documented statistically and revised, and three nominal species, one
32 subspecies and two morphotypes are now reassigned to *Hirnantia sagittifera* (M'Coy) *sensu*
33 *stricto*.

34

35 UUID: <http://zoobank.org/references/3f83fb1e-a6dd-4585-9f9f-9586dad28244>

36

37 **Introduction**

38

39 The end-Ordovician marked the first major extinction event of the Phanerozoic. There were
40 significant changes in the brachiopod faunas during the extinction and subsequent recovery.

41 The biotic event heralded the arrival of new groups of brachiopod communities assigned to
42 the *Hirnantia* Fauna (Temple, 1965). The *Hirnantia* Fauna is a worldwide, cool-water

43 brachiopod fauna that originated and flourished in shallow-water marine environments at

44 intermediate to low latitudes (e.g., Rong and Harper, 1988, 1999; Rong et al., 2002). The

45 uniqueness of the *Hirnantia* Fauna throughout geological time, its near global distribution,
46 distinctive and limited stratigraphical range (Hirnantian, uppermost Ordovician), have made
47 it a focus for evolutionary paleobiologists and biostratigraphers.

48 After almost half century of intense investigation, the *Hirnantia* Fauna is now well
49 known for its importance in the study of end-Ordovician macroevolutionary processes (e.g.,
50 Temple, 1965; Wright, 1968; Amsden, 1974; Lespérance and Sheehan, 1976; Rong 1979;
51 Nikitin et al., 1980; Cocks, 1982; Benedetto, 1986; Rong and Harper, 1988, 1999; Owen et
52 al., 1991; Harper and Rong, 1995, 2001; Rong et al., 2002; Jin and Zhan, 2008; Zhan et al.,
53 2010; Hints et al., 2012; Benedetto et al., 2013; Harper and Hints, 2016). Detailed
54 investigation of the *Hirnantia* Fauna on a global scale has already made a significant
55 contribution to the recognition of the essence of the end-Ordovician mass extinction. There
56 are more than 100 localities in nearly 30 countries and regions where the *Hirnantia* Fauna has
57 been documented (Rong, 1979; Rong et al., 2002). In South China, *Hirnantia* faunas from 24
58 localities were summarized and discussed, most of which are in Guizhou, Sichuan, and Hubei
59 provinces on the Upper Yangtze Platform (Rong et al., 2002; Zhan et al., 2010). Among them,
60 the collections from Yichang, Hubei Province where the Hirnantian GSSP is situated, were
61 systematically described in detail (Rong, 1984; Zeng et al., 2016). The brachiopod fauna is
62 highly diverse (over 20 genera) and was assigned to Benthic Assemblage (BA) 3, the
63 bathymetric zone of the deeper parts of the shallow-water platform (Rong, 1984).

64 Fieldwork by the author group during past several decades has dramatically increased
65 collections of the *Hirnantia* Fauna in South China. Recently, we discovered two new
66 associations of *Hirnantia* Fauna in Meitan County and on the east border of the Zunyi area

67 (Figure 1). The two associations dominated by *Hirnantia sagittifera* (M' Coy) with moderate
68 diversity (14 species) are assigned to BA 2-3, a typical shallower-water environment. The
69 sufficiency of specimens of *Hirnantia sagittifera* in the study has permitted analysis of its
70 population dynamics following the previous study (Li, 1996) and detection of intraspecific
71 variation. In this paper, the *Hirnantia* Fauna in Meitan County is reported for the first time
72 and its ecology discussed. One new species, *Minutomena missa* n. sp. is systematically
73 described. For *Hirnantia sagittifera* (M' Coy), Temple (1965) noted that significant variation
74 exists in populations of the species, however, this fact and its significance has largely been
75 ignored. Here, based on measurements of global representative specimens of species, together
76 with related specimens from Yichang and this study, we quantitatively analyze this variation.

77

78 **Geological setting**

79

80 The Kuanyinchiao Beds were also named as Kuanyinchaio Formation (e.g., Wang et al.,
81 2018). However, due to its limited thickness which is mostly less than 2 meters, we retain the
82 term Kuanyinchiao Beds in this study. Graptolites from shales underlying and overlying the
83 Kuanyinchiao Beds in many nearshore areas indicate that the unit mostly lies between the
84 *Metabolograptus extraordinarius* Biozone and the *M. persculptus* Biozone. The
85 Kuanyinchiao Beds in South China is commonly composed of dark gray argillaceous
86 limestone, containing abundant brachiopods, trilobites, corals, and a few other fossil groups
87 (Rong, 1979). However, lithological and faunal variations in the Kuanyinchiao Beds have
88 been reported at different localities (e.g., Rong et al., 2002). In many localities in the near-

89 shore facies belt, the beds are split into two parts on lithological differences, such as those in
90 Bijie, Renhuai, Tongzi, Fenggang, and Yinjiang, northern Guizhou, South China. The lower
91 Bed is commonly a gray-black mudstone whereas the upper Bed argillaceous limestone
92 (Wang et al., 2018).

93 The majority of the specimens in this study were collected from the Hetaoba section,
94 Meitan, northern Guizhou Province (Figure 2). The shelly bed consists of calcareous
95 mudstone with a thickness of only 20 cm, whereas the collection from the Baixiangyan
96 section is confined to 15 cm of strata in calcareous mudstone.

97

98 **Paleoecological significance**

99

100 *Community ecology.*—The term “*Hirnantia* Fauna” was replaced by “*Hirnantia* Community”
101 (Lespérance and Sheehan, 1976), and assigned to Benthic Assemblage 4, or even 5 (Sheehan,
102 1979), which belonged to relatively deeper-water environment settings; this was based on
103 principally their understanding of the depth ranges of the *Hirnantia* Fauna from Pércé,
104 Quebec. However, the *Hirnantia* Fauna may comprise more than one ecological community
105 (Rong, 1979; Harper, 1979, 1981). Wang et al. (1987) recognized 6 communities, with four of
106 them occupying BA3, the other two occupied BA2 and upper BA4, respectively. Detailed
107 studies of the paleogeographic framework for the region indicate that *Hirnantia* bearing
108 faunas may occupy a position equivalent to BA3 (Rong, 1984, 1986) or lower BA2 to BA3
109 (Rong and Harper, 1999), although the relative frequencies of the genera vary. Based on the
110 data from South China, it is suggested that, the commonly, highly diverse *Hirnantia* Fauna

111 (>15 genera) may have lived in a deeper-water environment than that of the more strictly
112 defined *Hirnantia* Fauna (commonly less than 10 genera) and may occupy a position in low
113 Benthic Assemblage 3.

114 The brachiopod specimens collected from two sections in this study are all identified
115 (Table 1). Although the relative abundance of common taxa such as *Hirnantia sagittifera*
116 (M'Coy) and *Eostropheodonta hirnantensis* (M'Coy) differ between the two assemblages, the
117 similarities in composition indicate that they may have belonged to a single paleoecologic
118 community. Two communities dominated by *Hirnantia* and *Aphanomena* are assigned to
119 upper BA3 (Rong, 1986; Wang et al., 1987). Rong (1986) recognized *Aphanomena-Hirnantia*
120 Community based on the materials from Donggongsi section, Zunyi City, that is very close
121 to the Baixiangyan section in this study. Since *Aphanomena* has been reassigned to
122 *Eostropheodonta* (Cocks and Rong, 2000), the two assemblages are recognized as the
123 *Hirnantia-Eostropheodonta* community, probably inhabiting a normal shallow-water
124 environment and occupying an upper BA 3 position (Rong, 1986; Wang et al., 1987). With
125 the assumption that a community dominated by *Paracraniops* lived in a lower BA3
126 environment (Wang et al, 1987), lower diversity, the absence of the two craniid genera
127 *Petrocrania* and *Xenocrania*, together with the fact that there is a great deal of biological
128 debris in shell beds in the Baxiangyan section, implying deposition above normal wave base,
129 it may be inferred that the environmental setting for the *Hirnantia* Fauna in the Hetaoba
130 section is relatively deeper than that of Baixiangyan.

131

132 *Population ecology and its significance.* —Since *Hirnantia* faunas immediately followed the

133 first phase of the end Ordovician mass extinction, the ecological effect of the event was a
134 major driver of the adaption of the *Hirnantia* Fauna to their environment. The environmental
135 effects of the extinction can be monitored by population analysis of the main components of
136 the *Hirnantia* Fauna.

137 Although fossil populations cannot be defined in the terms as those in modern biology
138 because of the nature of the fossil record, and some attributes of fossil populations are
139 difficult to study, paleoecologists have successfully applied their own approaches to
140 population ecology, particularly the study of population structure or dynamics in analyzing
141 fossil populations through time (see Dodd and Stanton, 1991; Brenchley and Harper, 1998).
142 Many studies revealed that survivorship patterns of time-averaged normal populations
143 without selective destruction can also represent an accurate census of their former living
144 populations (e.g., Cate and Evans, 1992; Li, 1996). During the identification of the specimens
145 from the Hetaoba section (GHH-02), we found that many delicate structures and even the
146 juveniles are well preserved, which suggest that population dynamics in these fossil
147 assemblages can be described and evaluated.

148 The brachiopods from Hetaoba are relatively abundant and assigned to a *Hirnantia*-
149 *Eostropheodonta* community since specimens of two genera account for nearly 60% of the
150 total. We measured complete specimens of the two taxa, and followed the method of
151 Levinton and Bambach (1970), recommended by previous studies (e.g., Li, 1996), and
152 calculated and plotted size frequency and survivorship curves for the two dominant
153 components (Fig. 3). The size-frequency distributions of the two populations with abundant
154 juveniles are all left-skewed, especially the population of *Hirnantia sagittifera*, indicating

155 that there were many young individuals and fewer older ones. These two populations,
156 featured by a high juvenile mortality and a lower mortality later in life, are characterized by
157 survivorship curves that are concave upwards. Most of the brachiopods died young with few
158 individuals living to maturity. The clear high juvenile mortality of *Hirnantia sagittifera*
159 indicates that an ecological threshold may exist during the early life of the population.
160 Compared with the curve of *Hirnantia sagittifera*, there is a lag in the peak for the population
161 of *Eostropheodonta hirnantensis* (M'Coy), suggesting better adaptation to the same
162 environment. The adaptation of both populations indicates that they are eurytopic, or their
163 habitat was not as severe to affect the major components of the fauna even in the shallow-
164 water environment after the end Ordovician mass extinction.

165

166 **Materials and locations**

167

168 The shelly beds of the Hirnantian Stage are widely distributed on the Yangtze Platform of
169 South China. The brachiopod fauna in this study was collected from the lower part of the
170 Kuanyinchiao Beds (for the division of the Kuanyinchiao Beds, see Wang et al., 2018 for
171 details) at two sections in Meitan County, and on the east border of Zunyi (Figure 1) by the
172 authors and their colleagues in 2011, 2017 and 2018. All specimens are preserved as internal
173 and external molds.

174 The section previously investigated is located at Hetaoba, 6 km northwest of Tiancheng
175 village, east of Meitan County Town (GPS: 27°48'07.5"N, 107°34'22.9"E). Common fossils
176 include abundant brachiopods, some trilobites, a few bryozoans and rare gastropods which

177 are preserved in gray-yellow calcareous mudstone. There are 603 specimens and 13 species
178 of brachiopods identified in our collections from the Kuanyinchiao Beds (GHH-02) in the
179 section. More than 90 specimens (33.8%) of *Hirnantia sagittifera* (M'Coy) were collected
180 together with many *Eostropheodonta hirnantensis* (M'Coy) (15.4%), *Paromalomena*
181 *macmahoni* (Reed) (10.3%), *Dalmanella testudinaria* (Dalman) (8.0%), *Palaeoleptostrophia*
182 sp.(6.5%), *Hindella crassa* (Williams) (6.5%), *Leptaena rugosa* (Dalman) (5.8%),
183 *Minutomena missa* n. sp. (4.1%), *Pseudopholidops partibilis* (Rong) (4.8%), and *Cliftonia* cf.
184 *psittacina* (Wahlenberg) (2.5%); and rare *Fardenia* (*Coolinia*) sp. (0.8%), *Xenocrania haimei*
185 (Reed) (0.5%) *Petrocrania* sp. (0.2%) and Discinidae gen. et sp. indet. (0.2%).

186 During recent fieldwork, a relatively low abundance shallow-water shelly brachiopod
187 association was collected from the Baixiangyan section (GMB-01), 7 km east of Yongle
188 village, the east border of Zunyi area (GPS: 27°51'51.8"N, 107°24'36.15" E). The fauna is
189 dominated by brachiopods together with a few trilobites and much crinoidal debris. In Total
190 79 specimens were gathered, including *Hirnantia sagittifera* (M'Coy)(35.4%),
191 *Eostropheodonta hirnantensis* (M'Coy)(31.6%), *Leptaena rugosa* (Dalman)(15.2%),
192 *Hindella crassa* (Williams)(7.1%), *Cliftonia* cf. *psittacina* (Wahlenberg)(5.1%), *Dalmanella*
193 *testudinaria* (Dalman)(4.3%) and *Paromalomena macmahoni* (Reed)(2.9%) are identified in
194 the sample.

195

196 *Repository and institutional abbreviation.*—Types, figures, and other specimens examined in
197 this study are deposited in the following institution: Nanjing Institute of Geology and
198 Paleontology, Chinese Academy of Sciences (the catalog numbers prefixed by NIGP).

199

200 **Systematic paleontology**

201

202 There are 13 species belonging to 13 genera together with an undeterminate species in the
203 two collections under study. The list of species reported is as follows, with some
204 representative specimens being illustrated of each: *Discinidae* gen. et sp. indet. (Fig. 4.7),
205 *Pseudopholidops partibilis* (Rong) (Fig. 4.1–4.3), *Xenocrania haimei* (Reed) (Fig. 4.4–4.5),
206 *Petrocrania* sp. (Fig. 4.6), *Leptaena rugosa* (Dalman) (Fig. 4.8–4.19), *Minutomena missa* n.
207 sp. (Fig. 5), *Paromalomena macmahoni* (Reed) (Fig. 6), *Eostropheodonta hirnantensis*
208 (M’Coy) (Fig. 7.10–7.15), *Palaeoleptostrophia* sp. (Fig. 7.1–7.9), *Cliftonia* cf. *psittacina*
209 (*Wahlenberg*) (Fig. 10.9–10.12), *Fardenia (Coolinia)* sp. (Fig. 7.16–7.18), *Dalmanella*
210 *testudinaria* (Dalman) (Fig. 10.1–10.4), *Hirnantia sagittifera* (M’Coy) (Fig. 8, Fig. 9), and
211 *Hindella crassa* (Williams) (Fig. 10.5–10.8). The one new species and two most dominant
212 species are discussed in detail. Data and some statistical information are also provided.

213

214 Class Strophomenata Williams, Carlson, Brunton, Holmer and Popov, 1996

215 Order Strophomenida Öpik, 1934

216 Superfamily Strophomenoidea King, 1846

217 Family Glyptomenidae Williams, 1965

218 Genus *Minutomena* Zeng et al., 2016

219

220 2016 *Minutomena* Zeng et al., p. 21.

221

222 *Type species.*—*Minutomena yichangensis* Zeng et al., 2016, p. 21, pl. 2, figs. 1–5;

223 Kuanyinchaio Beds (Hirnantian), Yichang, western Hubei, Central China.

224

225 *Minutomena missa* new species

226 Figure 5

227

228 *Type specimens.*—Holotype: NIGP 170742 (Fig. 10.5). Paratypes: NIGP 170738-170741,

229 NIGP 170743-170746. All from Kuanyinchiao Beds (Hirnantian, uppermost Ordovician) of

230 Hetaoba section, Meitan County, northern Guizhou, South China.

231

232 *Diagnosis.*—Larger *Minutomena* with coarser ornamentation and relatively small cardinalia.

233

234 *Occurrence.*—The new species is only known from Kuanyinchiao Beds (Hirnantian,

235 uppermost Ordovician) of Meitan County, Guizhou Province, South China.

236

237 *Description.*—Shells small to medium sized, 3.5 to 11.8 mm long, 4.3 to 13.6 mm wide,

238 transversely semi-circular; ventral valve slightly convex with maximum convexity near

239 umbo, dorsal valve nearly flat to weakly concave, lateral profile rarely resupinate.

240 Ornamentation of multicostellae, unequally round ribs increasing mostly by branching, about

241 3-4 per 2 mm at 10 mm from the umbo (Fig. 5.13); fine and prominent concentric growth

242 lines well developed. Substance of shell coarsely pseudopunctate.

243 Ventral interior with thin and short dental plates, divergent at an angle of about 70°–110°;
244 muscle scars small, broad, faintly visible, rarely extending beyond ends of dental plates
245 anteriorly; diductor scars subtriangular. Pedicle callist broad and well developed (Fig. 5.2,
246 5.4).

247 Dorsal interior with small bilobed cardinal process situated in the depressed posterior end
248 of notothyrial platform; socket ridges short, straight or slightly concave posteriorly, widely
249 divergent about 110°–135°; paired low and conspicuous ridges well developed in front of
250 socket plates (Fig. 5.5–5.10), divergent about 60°–100°; muscle field not seen.

251

252 *Etymology*.—Latin, *missus*, with regards to having overlooked this species during the
253 previous study.

254

255 *Materials*.—Eight ventral internal molds, 14 dorsal internal molds and three external molds.

256

257 *Remarks*.—The genus was published for the first time in a Chinese book, based on the
258 specimens collected from the GSSP area of the Hirnantian Stage (Zeng et al., 2016). Its close
259 relationship with *Paromalomena* was not noticed in the original study. However, the
260 specimens in this study were mistakenly assigned to species of *Paromalomena* by the first
261 author during the earlier study because of a lack of sufficiently informative materials. On the
262 basis of detailed systematic study, the differences between the two genera can be summarized
263 as: 1). Ornamentation of *Paromalomena* is of fine costellae, and increasing by intercalation;
264 concentric wrinkles or rugae commonly irregularly developed; weak and wavy (Fig. 6.2, 6.9).

265 Whereas costae of *Minutomena* are coarser and increasing mostly by branching, the wrinkles
266 or rugae rarely found on the genus (Fig. 5.13). 2). More importantly, side septa or paired low
267 ridges are absent or are inconspicuous in *Paromalomena* (Fig. 6.10), but are essentially
268 developed in front of socket plates in *Minutomena* (Fig. 5.11).

269 The new species is different from the type species mainly in two aspects: 1). The size of
270 *Minutomena yichangensis* Zeng et al., 2016 is commonly less than 8 mm in width; whereas
271 the specimens of *M. missa* n. sp. are mostly larger than 10 mm in width. 2). For the type
272 species, a pair of lateral ridges are often very prominent, with its anterior end even reaching
273 half of the valve length (Zeng et al., 2016, pl. 2 fig. 4a); however, the lateral ridges of *M.*
274 *missa* n. sp. are always weaker, with their anterior ends at 1/4 to 1/5 of valve length.

275

276 Order Strophomenida Öpik, 1934

277 Superfamily Strophomenoidea King, 1846

278 Family Leptostrophiidae Caster, 1939

279 Genus *Eostropheodonta* Bancroft, 1949

280

281 *Type species.*—*Orthis hirnantensis* M'Coy, 1851; Hirnant Formation (Hirnantian), Upper
282 Ordovician, Aber Hirnant, near Bala, Wales, UK.

283

284 *Eostropheodonta hirnantensis* (M'Coy, 1851)

285

Figure 7.10–7.15

286

287 *Type specimens.*—Lectotype: Internal mould of pedicle valve, A 28831, from Aber Hirnant,
288 North Wales (M'Coy, 1852, pl. 1H, fig. 11c; Temple, 1965, pl. 19, fig. 1).
289
290 *Remarks.*—The ornamentation of species assigned to *Eostropheodonta* has been considered
291 to be of taxonomic significance. However, it is highly variable even in specimens from the
292 same population (see discussion in Rong et al., 2019). Furthermore, the development of
293 dental plates is also variable. During the early study of the fauna, all related specimens were
294 assigned to *Eostropheodonta*. After detailed analysis of the variation, we consider that
295 specimens with vestigial or absent dental plates, together with the triangular and weakly
296 impressed muscle field, are better assigned to *Palaeoleptostrophia*, rather than
297 *Eostropheodonta*. Based on the dorsal internal molds of the species of the two genera, it is
298 difficult to differentiate the two (Rong and Cocks, 1994). Since species of
299 *Palaeoleptostrophia* have numerous coarse pseudopunctae in the posterolateral parts of its
300 both ventral and dorsal internal molds, we assign the dorsal internal molds with the
301 developed coarse pseudopunctae to *Palaeoleptostrophia*. The genus *Palaeoleptostrophia* was
302 thought to originate in the Rhuddanian, and the oldest Leptostrophiidae lacked dental plates
303 (Rong and Cocks 1994). There are some specimens of that family recorded as leptostrohiids
304 in the Hirnantian rocks of Latvia and Estonia (Hints and Harper, 2015). Thus, it could be
305 regarded as a progenitor taxon evolved during the crisis interval (Rong et al., 2019). The
306 evolutionary relationship between *Eostropheodonta* and *Palaeoleptostrophia* will be
307 discussed together with other genera of Leptostrophiidae in detail elsewhere.
308

309 Order Orthida Schuchert & Cooper, 1932

310 Suborder Dalmanellidina Moore, 1952

311 Superfamily Enteletoidea Waagen, 1884

312 Family Draboviidae Havlíček, 1950

313 Subfamily Draboviinae Havlíček, 1950

314 Genus *Hirnantia* Lamont, 1935

315

316 *Type species.*—*Orthis sagittifera* M' Coy, 1851; Hirnant Formation (Hirnantian), Upper

317 Ordovician, Aber Hirnant, near Bala, Wales, UK.

318

319 *Hirnantia sagittifera* (M' Coy, 1851)

320 Figure 8–9

321

322 Here only the data used in the analysis are listed, see Appendix of Rong et al., 2019 for detail

323 list.

324 1851 *Orthis sagittifera* M' Coy, p. 398.

325 1935 *Hirnantia sagittifera* (M' Coy); Lamont, p. 313, pl.7, figs. 20–22.

326 1965 *Hirnantia sagittifera*; Temple, p. 349–401, pl.11, fig. 8; pl. 12, figs. 1–10; pl.13, figs.

327 1–10; pl. 14, figs.1–8.

328 1968 *Hirnantia sagittifera*; Bergström, p. 10, pl. 3, fig. 13; pl. 4, figs. 1, 2.

329 1974 *Hirnantia magna* Rong, Xu and Yang, p. 196, pl. 92, fig. 25, 26.

330 1976 *Hirnantia sagittifera*; Lespérance and Sheehan, p. 723, pl. 109, figs. 3–11.

- 331 1977 *Hirnantia sagittifera*; Havlíček, p. 266, pl. 29, figs. 15–23.
- 332 1979 *Hirnantia sagittifera fecunda* Rong, pl. 1, figs. 10, 14, 15.
- 333 1980 *Hirnantia sagittifera*; Nikitin, p. 43, pl. 11, figs. 1–4.
- 334 1982 *Hirnantia sagittifera*; Cocks, p. 763–764, pl. 78, figs. 6–10; pl. 79, figs. 1–3; pl. 80, fig.
- 335 8.
- 336 1983 *Hirnantia sagittifera*; Mergl, p. 343–344, pl. 4, figs. 14–16.
- 337 1984 *Hirnantia sagittifera*; Rong, 139. 141, pl. 6, figs 1–17; pl. 7, figs 1–8; pl. 12, fig. 15;
- 338 text-figs 10, 11.
- 339 1986 *Hirnantia sagittifera*; Benedetto, p. 444, pl. 1, figs 11–15; pl. 2, figs 1–3.
- 340 1987 *Hirnantia sagittifera*; Rong and Xu, p. 5–6, pl. 1, figs 9, 12, 13, 16, 17, 20–22, 24, 26.
- 341 1999 *Hirnantia sagittifera*; Villas, Lorenzo and Gutierrez-Marco, p. 193, fig. 4 (a–i).
- 342 2012 *Hirnantia sagittifera*; Hints, Parnaste and Gailite, p. 67–70, fig. 2. A–J.
- 343 2016 *Hirnantia sagittifera* morph. Bohemia Havlíček (1977), Zeng et al., p. 69–70, pl. 35,
- 344 figs. 1–12; pl. 36, figs. 1–2.
- 345 2016 *Hirnantia sagittifera* morph. Poland Temple (1965), Zeng et al., p. 70–71, pl. 37,
- 346 figs.1–11.
- 347 2016 *Hirnantia magna*; Zeng et al., p. 71–72, pl. 36, figs. 3–12.
- 348 2016 *Hirnantia septumis* Zeng et al, p. 72–73, pl. 38, figs. 1–12.
- 349 2016 *Hirnantia fecunda* (Rong), Zeng et al., p. 74, pl. 34, figs. 6–12.
- 350
- 351 *Type specimens*.—Lectotype: Internal mold of brachial valve, A 41217, from Aber Hirnant,
- 352 North Wales (M’Coy, 1852, pl. 1H, fig. 17; Temple, 1965, pl. 11, fig. 8).

353

354 *Remarks.*—*H. sagittifera* has been described in detail by many different authors (e.g.,
355 Temple, 1965; Havlíček, 1977; Cocks, 1982; Villas et al., 1999; Hints et al., 2012 and
356 therein). The population variation in the species was noted mainly in size and outline
357 (Temple 1965; Cocks 1982; Cocks and Torsvik, 2002). Its adult shell width can reach 50 mm
358 (Bergström, 1968; Hints et al., 2012) or less than 15 mm in different populations. The
359 variation of its internal characters is also significant. Rong (1984) initially discussed the
360 morphology variation of the cardinal process and considered it can be both simple and
361 bilobate with peculiar crenulations on its top and sides. The variation in *H. sagittifera* is so
362 considerable that some specimens in the same population could be easily identified as
363 separate species.

364 Recently Zeng et al. (2016) monographed a *Hirnantia* Fauna from Yichang, where the
365 GSSP of Hirnantian Stage is situated. With abundant and well-preserved specimens of
366 *Hirnantia* from the same beds, they identified 5 species (or morphotypes), *H. sagittifera*
367 (M'Coy) morph. Bohemia, *H. sagittifera* (M'Coy) morph. Poland, *H. magna* (Rong et al.), *H.*
368 *septumis* (Zeng) and *H. fecunda* (Rong). The main criteria for discrimination were shell size
369 and the development of muscle fields. However, these key differences are not clear with
370 reference to their specimens.

371 In order to assess the value of the differences between the “5 species”, we measured most
372 of the taxa from published figures. We also compared several specimens from other
373 collections from South China: 1) Donggongsi section (AAE-080) of Zunyi city, Guizhou
374 province; 2) Tianbatou section of Gongxian, Sichuan province; 3) Wangjiawan section of

375 Yichang area, Hubei Province (the GSSP of the Hirnantian). Seven parameters are selected
376 for dorsal internal molds, such as length, width, length of cardinalia (from anterior margin of
377 brachiophore bases to the apex of shell), width of cardinalia (the maximum distance between
378 two brachiophore bases), muscle field position (distance from anterior margin of muscle
379 fields to the apex of shell), width of muscle fields, and angle of brachiophore supporting
380 plates; six parameters are available for the ventral internal molds, as length, width, length of
381 muscle field, width of muscle field, width of adductor muscle scar, and angle of dental plates
382 (Appendix 1 and 2). Using PCA (principal components analysis) to interrogate the
383 measurements above, most of the variation in these specimens can be analyzed and evaluated.
384 Since related specimens from a previous study (Rong, 1979, 1984) were collected from the
385 same locality, we also included them in the analysis. The results (Fig. 10.1) displayed a clear
386 pattern of variation. No single morphological group can be clearly recognized from the
387 scatters, indicating that the variation of each “species” is significant but continuous for both
388 outline and internal structures.

389 As Zeng (2016) considered that ‘*H. sagittifera* (M’Coy) morph.’ Bohemia is different
390 from ‘*H. sagittifera* (M’Coy) morph.’ Poland mainly on its larger size. However, as the
391 specimens were collected from the same beds, we consider that many of ‘*H. sagittifera*
392 (M’Coy) morph. Poland’ are immature specimens lacking the development of the dorsal
393 muscle field. Since the material from Bohemia and Poland are both assigned to *H. sagittifera*,
394 this morphotype is not necessarily present. Similarly, *H. magna* (Rong et al.) was erected
395 mainly by virtue of its larger size (It can reach 40 mm) and the development of cardinalia and
396 muscle fields based on specimens from very shallow-water (BA2) populations and is

397 represented only by large adult shells rather than different species. Most of the shells were
398 significantly larger than those collected from other localities at that time (Rong et al., 1974).
399 However, large and even larger specimens *H. sagittifera* have been reported from Sweden
400 (Bergström, 1968), Oslo (Cocks, 1982), Argentina (Benedetto, 1986) and Estonia (Hints et
401 al., 2012), which undermine the validity of the species *H. magna*. Here, we figure the
402 holotype of this species (Fig. 9.11) and note the similarity between the holotype of *H. magna*
403 and the large specimens from this study (Fig. 8.11), suggesting *H. magna* can be treated as a
404 junior synonym of *H. sagittifera* (M'Coy).

405 *H. fecunda* Rong (Zeng et al., 2016) was firstly erected as *H. sagittifera fecunda* (Rong,
406 1979) mainly on the basis of its shorter and narrower cardinalia and ventral muscle fields.
407 Both the measurements and the results of the PCA analysis indicates that the variation of *H.*
408 *sagittifera* encompasses that of *H. fecunda*. In this study, specimens collected from Meitan
409 with short brachiophore bases and muscle fields are also similar to “*H. sagittifera fecunda*”.
410 However, we consider that the subspecies should be suppressed and assigned to *H.*
411 *sagittifera*. The typical character of *H. septumis* Zeng is a short and narrow medium septum
412 developed in the ventral valves. We consider that this character is an indicator of adult or
413 gerontic individuals; these specimens are quite large with a well-developed muscle field, and
414 some of the ventral valves have even muscle bounding ridges. Similar but weaker muscle-
415 bounding ridges are also identified on the specimens collected from Guizhou (Fig. 9.5, 9.6).

416 We suggest that the “five species” of Zeng et al. (2016) together with *H. magna* (Rong et
417 al., 1974) and *H. sagittifera fecunda* (Rong, 1979) should be reassigned to *H. sagittifera*
418 *sensu stricto*. The “unique characters” described for each of the above taxa are part of wide

419 ranging, continuous intra-specific variation rather than criteria for the establishment of
420 different species. Besides the data from Yichang of China presented above, representative
421 specimens of *H. sagittifera* from different regions and paleo-plates reveal further variation in
422 these species. We measured the figured specimens from Wales (Temple, 1965; Walmsley et
423 al., 1969; Brenchley et al., 2006), Sweden (Bergström, 1968), Quebec, Canada (Lespérance
424 and Sheehan, 1976), Bohemia (Havlíček, 1977), Kazakhstan (Nikitin, 1980), Oslo (Cocks,
425 1982), Algeria (Mergl, 1983), Argentina (Benedetto, 1986), Tibet of China (Rong and Xu,
426 1987), Spain (Villas et al., 1999) and Estonia (Hints et al., 2012). Most of the specimens are
427 preserved as internal molds, with the exception of those from Estonia with calcareous shells,
428 mainly from drill cores. Using the specimens from Yichang as a template, based on their
429 measurements (Appendix 1 and 2), the results from PCA clearly indicate that the variation of
430 the global species can be captured in the data from Yichang, China (Fig. 10.2). Some outliers,
431 such as the data from Estonia and Sweden, may be due to different modes of preservation,
432 which introduce a bias into the measurements. The differences between specimens from other
433 regions were emphasized (e.g., Zeng et al., 2016). However, we consider that given the wide
434 variability of the Yichang material, these differences may be accommodated within the intra-
435 specific variability of *H. sagittifera*.

436 Specimens of *Hirnantia sagittifera* collected from Guizhou, South China display a
437 marked range of variation. Dorsal valves can be relatively flat or occasionally very convex
438 (Fig. 8.13). The cardinal process can be simple (Fig. 8.15) or bilobed with crenulations (Fig.
439 9.10) in adult specimens. Dorsal muscle fields also vary from weak to well developed, with
440 different outlines. The anterior pairs of dorsal muscle scars can be equal to (Fig. 8.10) or

441 smaller (Fig. 9.11) than the posterior pairs. The angle between the brachiophore supporting
442 plates or dental plates is clearly varied. The ventral adductor muscle field can be wide to very
443 narrow. The anterior extremity of adductor field on a few specimens can form a short septum
444 as described in the different species (Zeng et al., 2016). With sufficient sampling of
445 populations, more variation will be discovered.

446

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448

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618

619 **Figure and table Captions**

620

621 Figure 1. Map showing the locations of the two sections, from where the near-shore

622 *Hirnantia* brachiopod Fauna was discovered.

623

624 Figure 2. The *Hirnantia* Fauna occurring in the Kuanyinchiao Beds at two sections from

625 northern Guizhou, South China.

626

627 Figure 3. Size frequency histograms and survivorship curves for the two populations of the

628 dominant components (*Hirnantia* and *Eostropheodonta*) in the Hetaoba section (GHH-02).

629

630 Figure 4. *Pseudopholidops partibilis* (Rong) (1–3), (1) a dorsal (?) internal mold (NIGP

631 170724); (2, 3) and two dorsal(?) external molds (NIGP 170724, 170726); *Xenocrania haimei*

632 (Reed) (4, 5), dorsal external mold (4) and its local enlargement (5) showing detail of

633 ornamentation (NIGP 170727); *Petrocrania* sp., (6) dorsal internal mold (NIGP 170728);

634 Discinidae gen. et sp. indet. (7) dorsal valve exterior (NIGP 170729); *Leptaena rugosa*

635 (Dalman) (8–19), (8–11) four ventral internal molds (NIGP 170730–170733); (12) dorsal

636 internal mold (NIGP 170734); (13–15) dorsal internal mold (13), its latex cast (15) with local

637 enlargement showing detail of cardinalia (14) (NIGP 170735); (17, 18) dorsal internal mold

638 (17) and its latex cast (18) (NIGP 170736); (16, 19) dorsal external mold (19) and its local

639 enlargement (**16**) showing detail of ornamentation (NIGP 170737). All specimens from GHH-
640 02 except (**17**) and (**18**) from GMB-01. Scale bar represent 2 mm except where noted.

641
642 Figure 5. *Minutomena missa* n. sp. (**1–4**) four ventral internal molds (NIGP 170738–170741);
643 (**5, 6, 8**) dorsal internal mold (**5**), its latex cast (**8**) with local enlargement showing detail of
644 cardinalia (**6**) (NIGP 170742); (**7, 11**) dorsal internal mold (**11**) and local enlargement of its
645 latex (**7**) showing detail of cardinalia (NIGP 170743); (**9, 10**) two dorsal internal molds
646 (NIGP 170744, 170745); (**12, 15, 16**) dorsal internal mold (**12**), its external mold (**16**) with
647 local enlargement of its latex (**15**) showing detail of ornamentation (NIGP 170746); (**13, 14**)
648 dorsal external mold (**13**) and its local enlargement (**14**) showing detail of ornamentation
649 (NIGP 170747). All specimens from GHH-02. Scale bar represents 2 mm except where
650 noted.

651
652 Figure 6. *Paromalomena macmahoni* (Reed). (**1, 2**) two ventral internal molds (NIGP
653 170748, 170749); (**3, 4, 10, 11**) four dorsal internal molds (NIGP 170750–170753); (**5–8**)
654 dorsal internal mold (**5**) and its latex cast (**8**), with local enlargement of mold (**6**) and cast (**7**)
655 showing detail of cardinalia (NIGP 170754); (**9, 12**) dorsal external mold (**12**) and its local
656 enlargement (**9**) showing detail of ornamentation (NIGP 170755). All specimens from GHH-
657 02. Scale bars represent 2 mm except where noted.

658
659 Figure 7. (**1–9**) *Palaeoleptostrophia* sp. (**1, 4**) two ventral internal molds (NIGP 170756,
660 170757); (**2, 3**) ventral internal mold and its local enlargement showing absence of dental

661 plates (NIGP 170758); **(5, 8, 9)** dorsal internal mold **(5)**, its external mold **(8)** with local
662 enlargement **(9)** showing detail of ornamentation (NIGP 170759); **(6, 7)** two dorsal internal
663 molds (NIGP 170760, 170761); *Eostropheodonta hirnantensis* (M'Coy) **(10–15)**, **(10–12)**
664 three ventral internal molds (NIGP 170762–170764); **(13, 14)** two dorsal internal molds
665 (NIGP 170765, 170766); **(15)** local enlargement of a dorsal internal mold (NIGP 170767);
666 *Fardenia (Coolinia)* sp. **(16–18)**, **(16, 18)** ventral internal mold and its external mold (NIGP
667 170768); **(17)** dorsal internal mold (NIGP 170769). All specimens from GHH-02. Scale bars
668 represent 2 mm.

669
670 Figure 8. *Hirnantia sagittifera* (M'Coy). **(1–3, 6)** four ventral internal molds (from GHH-02,
671 NIGP 170770–170773); **(4, 5, 7–9, 14)** seven dorsal internal molds (all from GHH-02 except
672 7 from GMB-01, NIGP 170774–170779); **(10, 16)** dorsal internal mold and its external mold
673 (from GHH-02, NIGP 170780); **(11–13, 15)** dorsal internal mold **(11)** and its posterior view
674 **(13)**, its latex cast **(15)** with local enlargement **(12)** showing detail of cardinalia (from GMB-
675 01, NIGP 170781). Scale bars represent 2 mm.

676
677 Figure 9. *Hirnantia sagittifera* (M'Coy). **(1–3)** ventral internal mold **(1)**, dorsal internal mold
678 **(2)** with its latex cast **(3)**, from Huangtian section of Gongxian, Sichuan province (NIGP
679 170782, 170783); **(4, 7)** ventral internal mold and a dorsal internal mold, from Wangjiawan
680 section of Yichang area (the GSSP of Hirnantian Stage) (NIGP 170784, 170785); **(5, 6)**
681 ventral internal and its latex cast (from AAE-080, NIGP 170786); **(9, 10)** dorsal internal mold
682 and its latex cast (from AAE-080, NIGP 170787); **(8, 11)** ventral internal mold and dorsal

683 internal mold (AAE-080), the two specimens were designated as paratype and holotype of
684 *Hirnantia magna* Rong et al., 1974 separately (NIGP 22275, 22276). Scale bars represent 2
685 mm.

686

687 Figure 10. Principal components analysis of data for 6 parameters for seven “species” of
688 *Hirnantia* (1), and globally-distributed *Hirnantia sagittifera* (2), to show the variation.

689

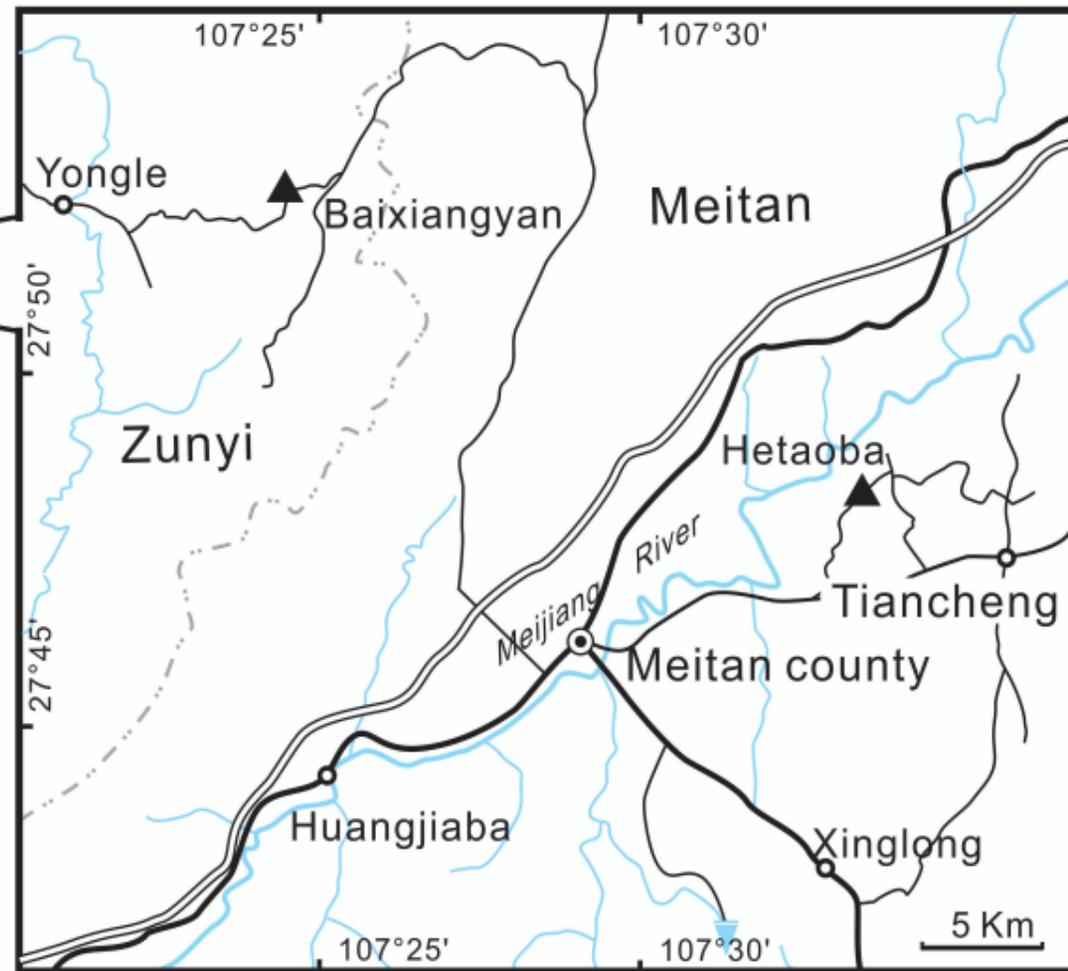
690 Figure 11. *Dalmanella testudinaria* (Dalman) (1–4), (1) ventral internal mold (NIGP
691 170788); (2–4) three dorsal internal molds (NIGP 170789–170791); *Hindella crassa*
692 (Williams) (5–8), (5, 6) two ventral internal molds (NIGP 170792, 170793); (7, 8) two dorsal
693 internal molds (NIGP 170794, 170795); *Cliftonia* cf. *psittacina* (Wahlenberg) (9–12), (9)
694 ventral internal mold (NIGP 170796); (10) ventral external mold (NIGP 170797); (11, 12)
695 two dorsal internal molds (NIGP 170798, 170799). All from GHH-02 except 11 and 12 from
696 GMB-01. Scale bars represent 2 mm.

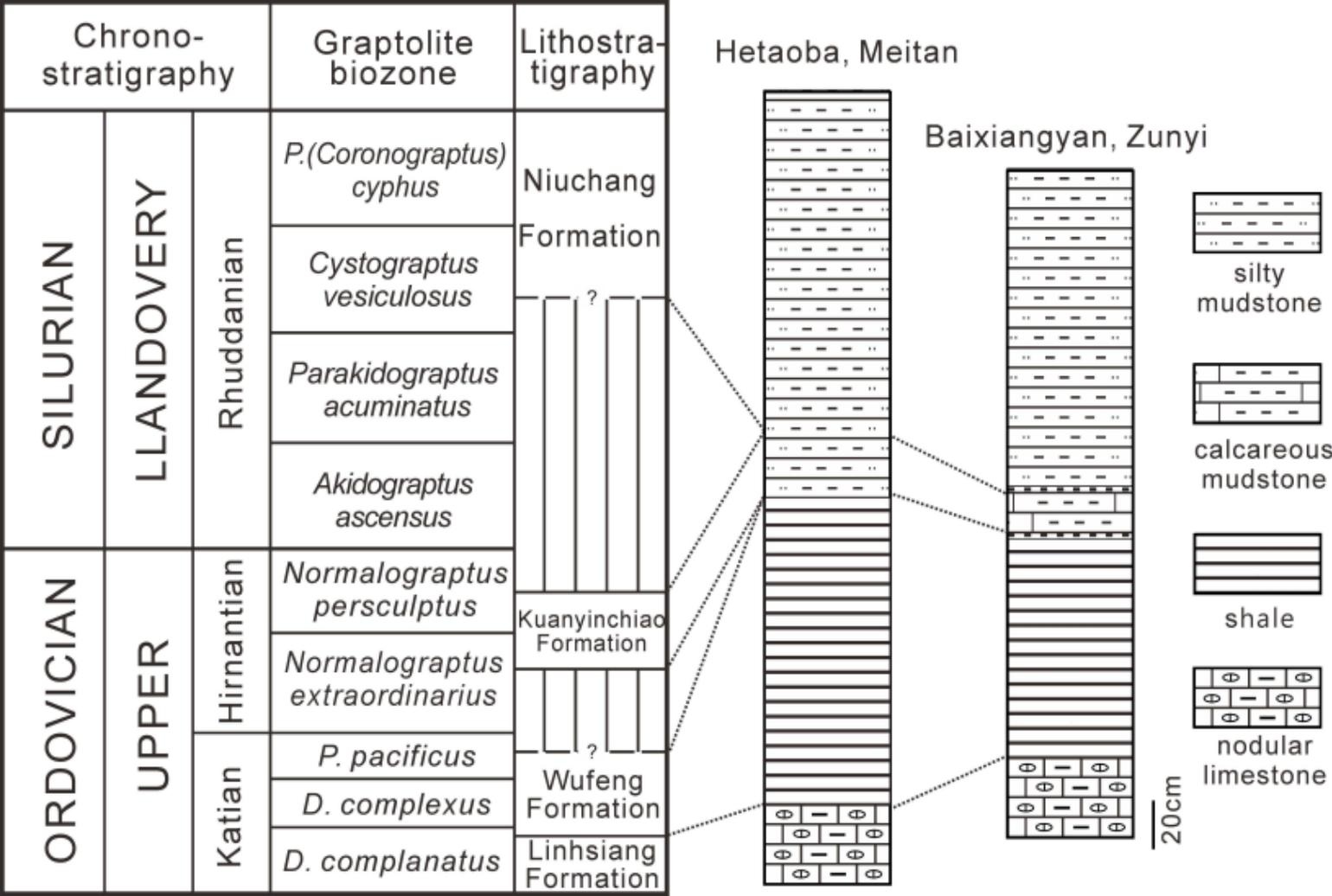
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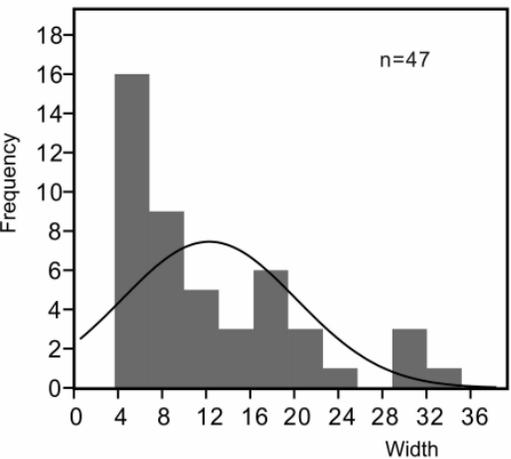
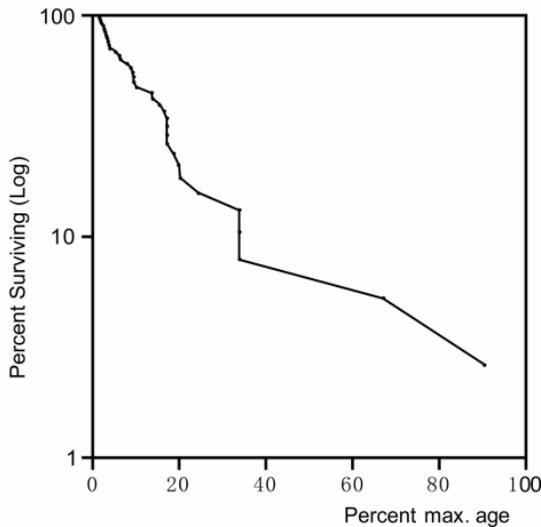
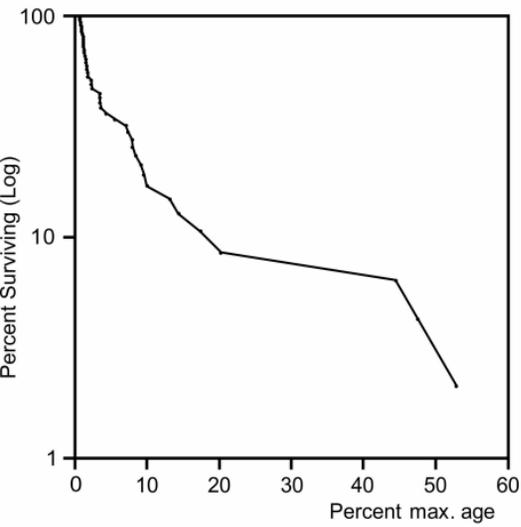
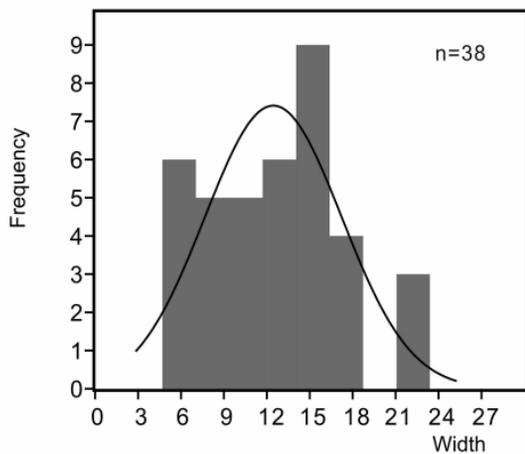
698 Table 1. Taxonomic list at species level for this study with numbers of specimens from the
699 two collections.

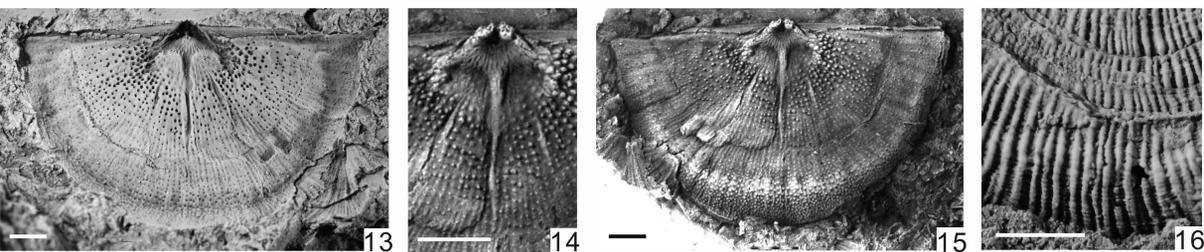
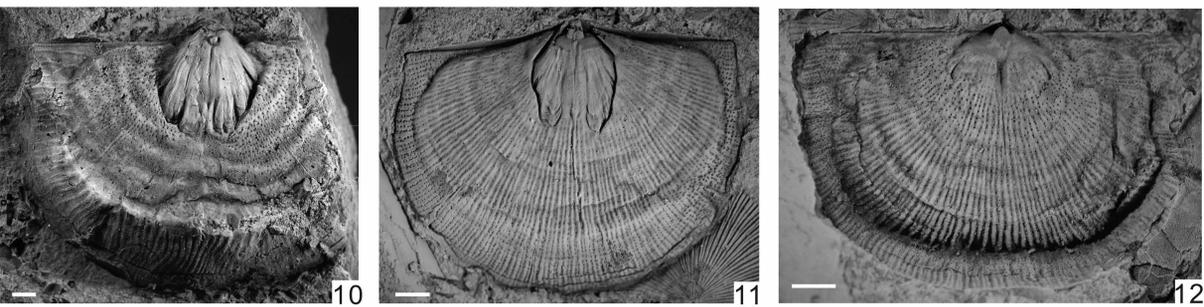
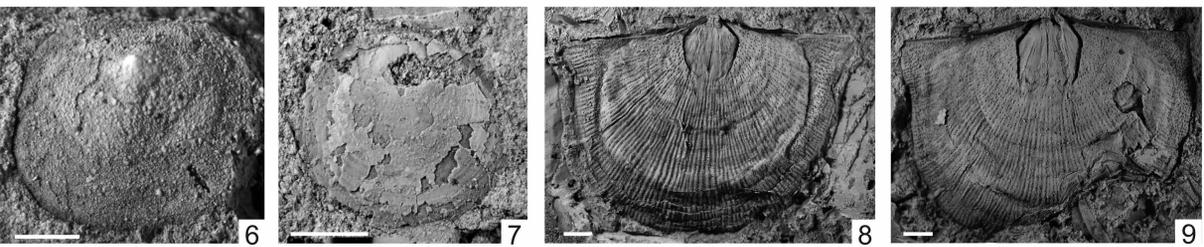
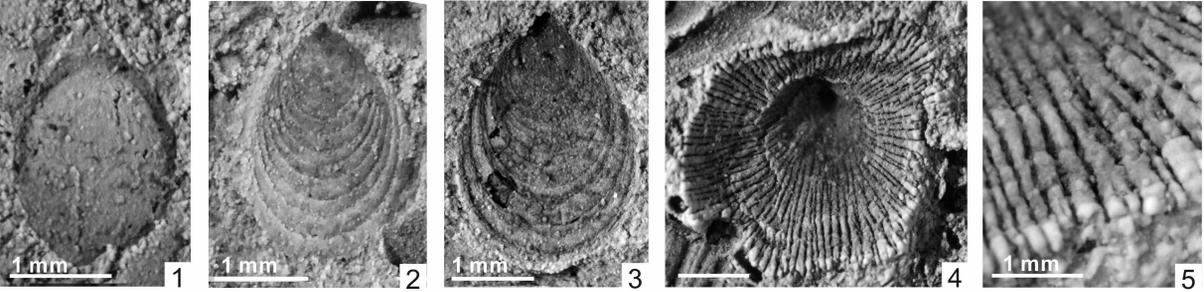
Table 1. Taxonomic list at species level for this study with numbers of specimens from the two collections.

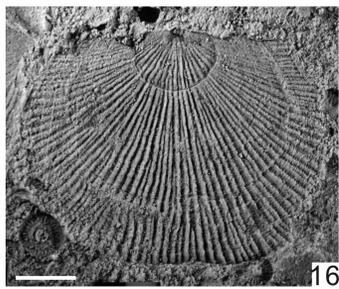
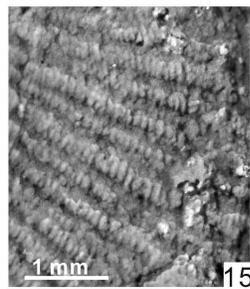
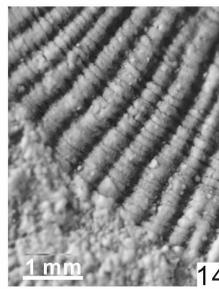
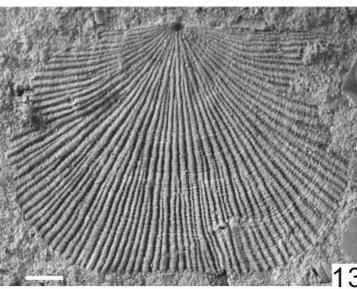
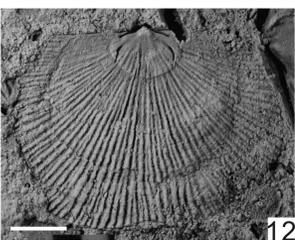
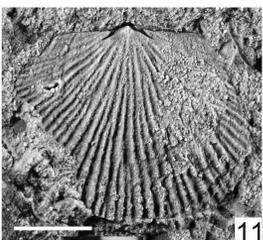
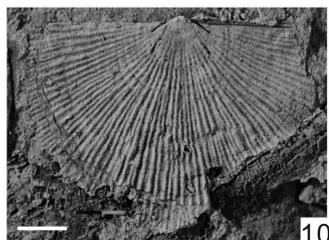
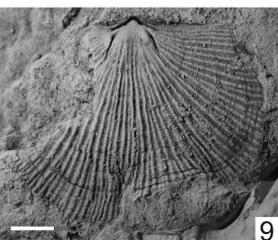
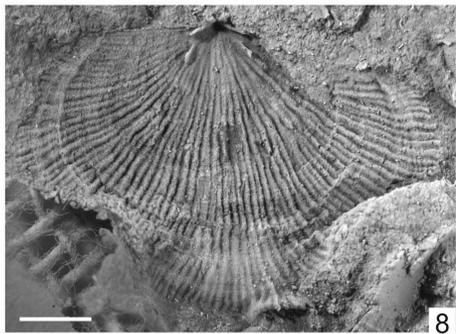
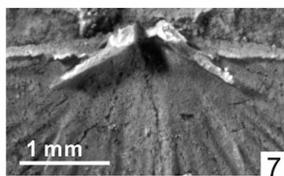
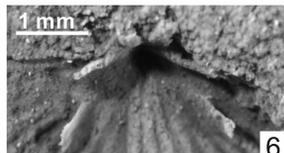
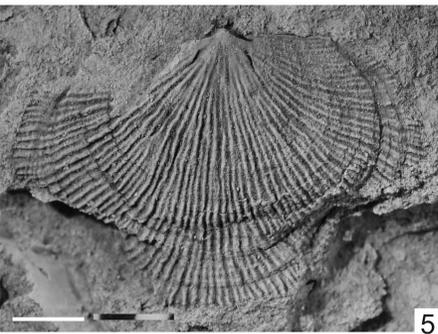
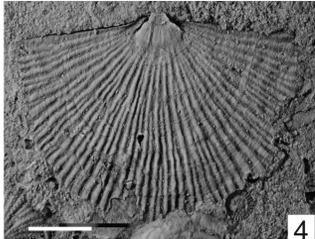
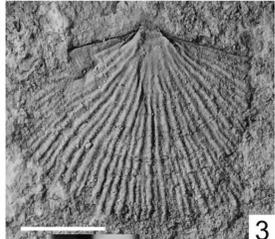
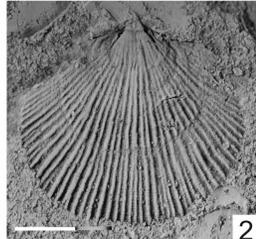
Species	Hetaoba section (GHH-02)			Baixiangyan section (GMB-01)		
	Ventral internal molds	Dorsal internal molds	External molds	Ventral internal molds	Dorsal internal molds	External molds
<i>Pseudopholidops partibilis</i> (Rong)	12	8	9			
<i>Petrocrania</i> sp.		1	1			
<i>Xenocrania haimeii</i> (Reed)		1	2			
<i>Leptaena rugosa</i> (Dalman)	15	11	9	5	4	3
<i>Paromalomena macmahoni</i> (Reed)	19	27	16		2	
<i>Minutomena missa</i> n.sp.	8	14	3			
<i>Eostropheodonta hirsantensis</i> (M'Coy)	43	21	30	12	2	11
<i>Palaeoleptostrophia</i> sp.	26	4	9			
<i>Fardenia</i> (<i>Coolinia</i>) sp.	1	1	3			
<i>Hirnantia sagittifera</i> (M'Coy)	87	78	41	11	14	3
<i>Dalmanella testudinaria</i> (Dalman)	9	31	8	1	2	
<i>Cliftonia</i> cf. <i>psittacina</i> (Wahlenberg)	6	2	7	1	2	1
<i>Hindella crassa</i> (Williams)	16	14	9	1	4	
Discinidae gen. et sp. indet.		1				

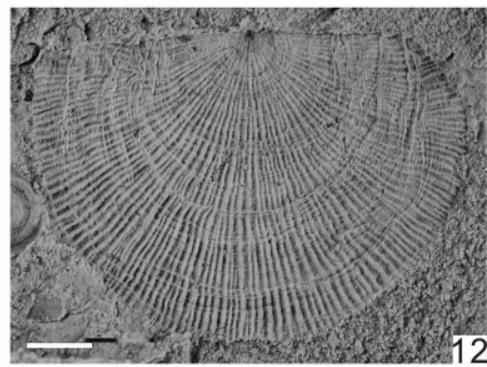
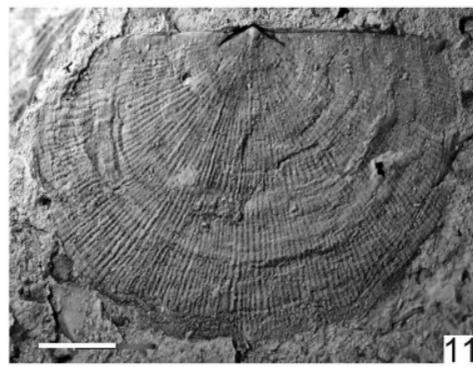
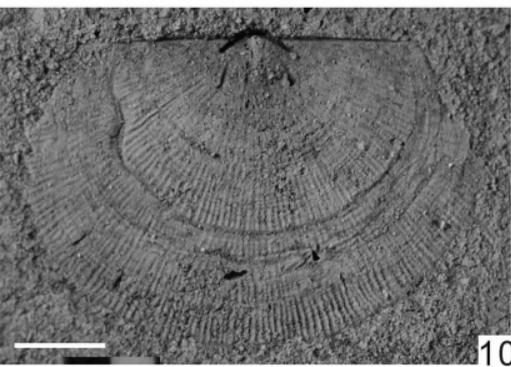
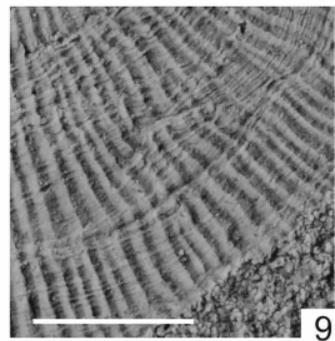
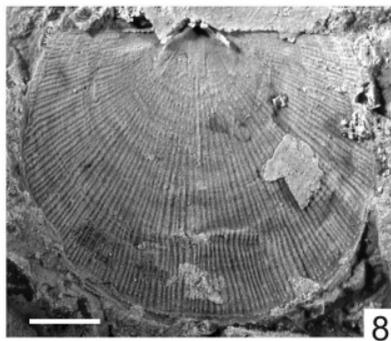
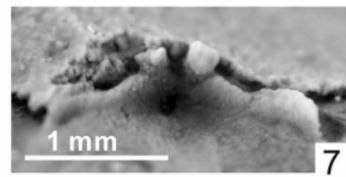
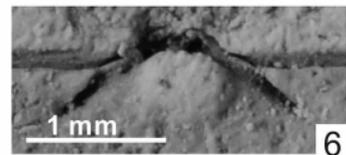
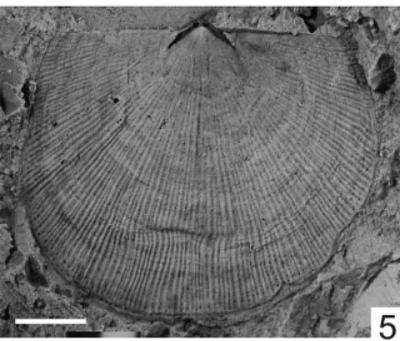
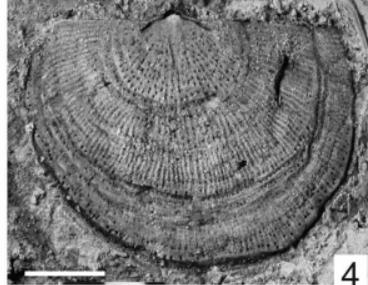
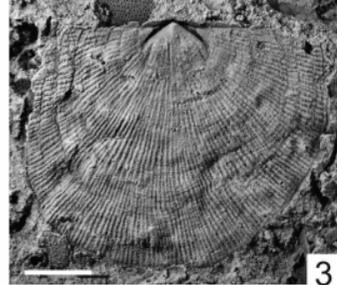
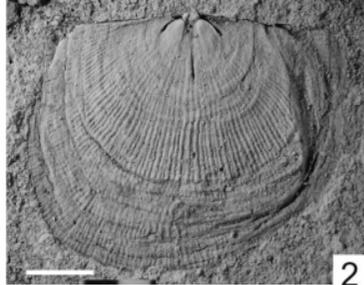
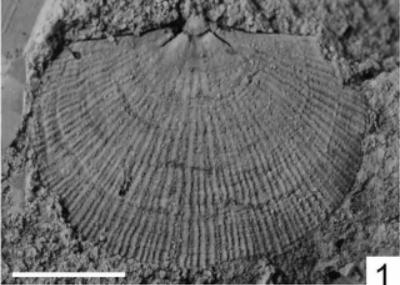


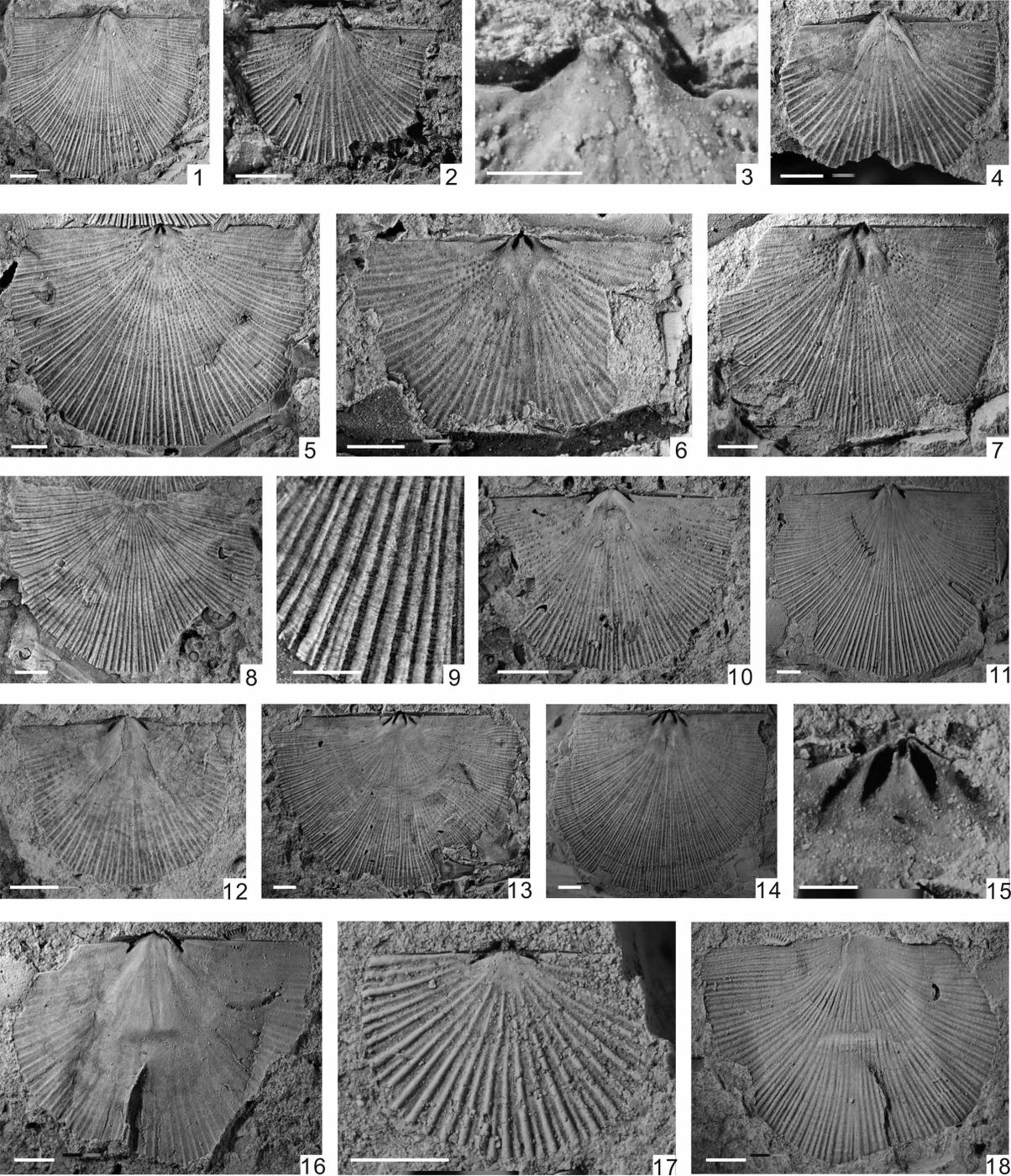


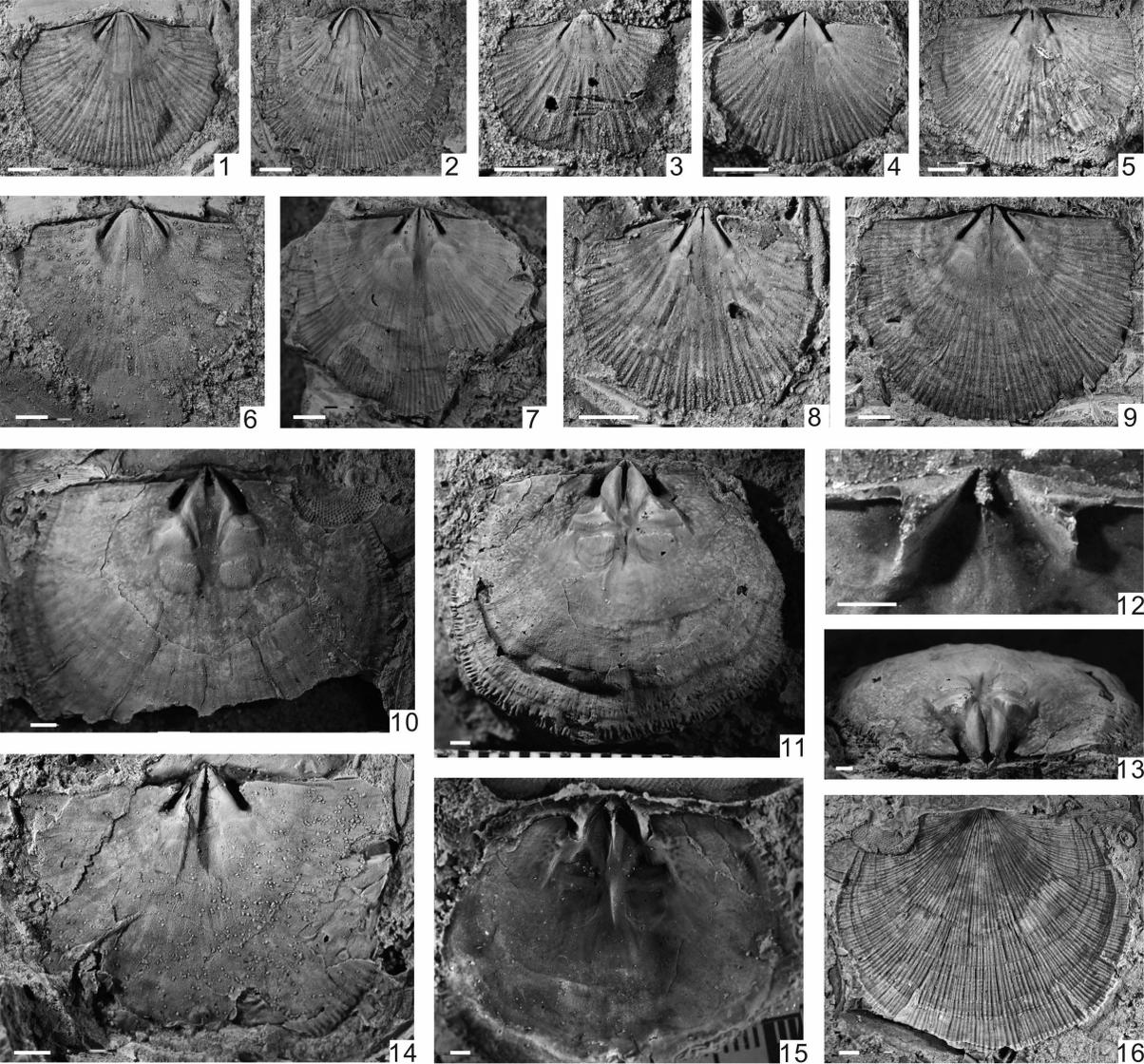
Hirnantia sagittifera (M'Coy)*Eostropheodonta himantensis* (M'Coy)

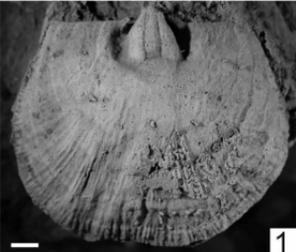












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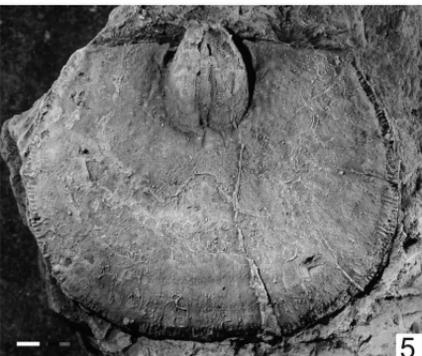
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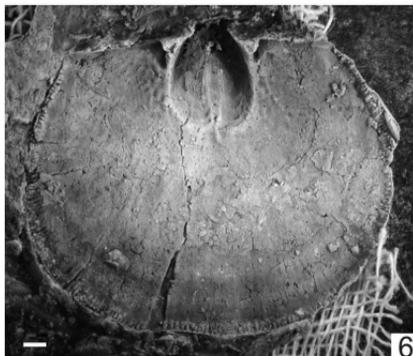
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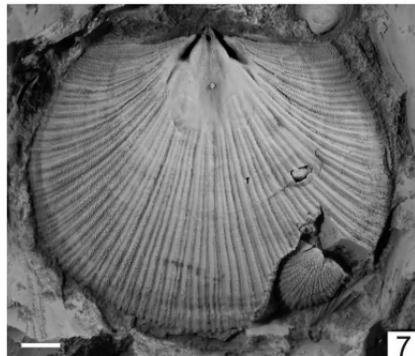
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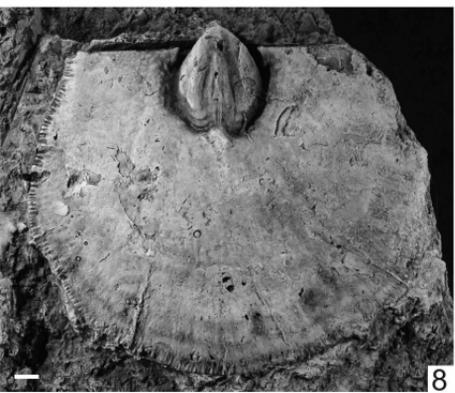
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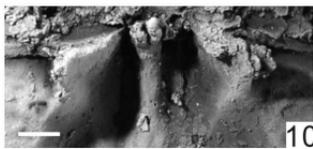
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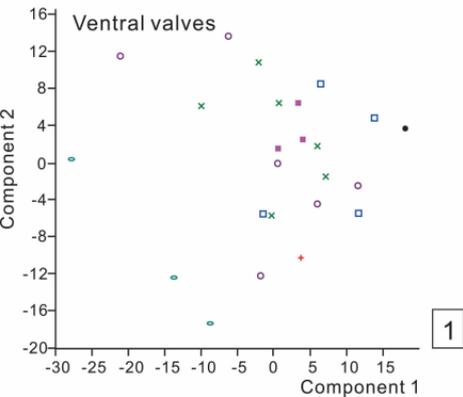
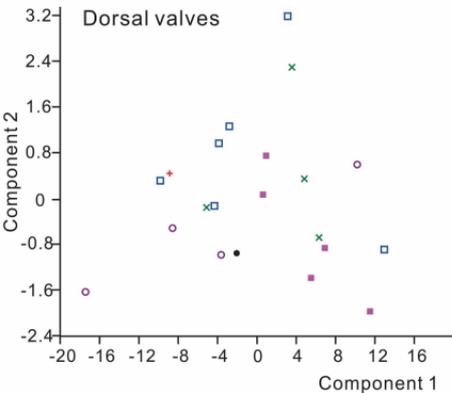


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- *Hirnantia magna* Rong (Zeng et al., 2016)
- × *Hirnantia septumis* Zeng (Zeng et al., 2016)
- + *Hirnantia fecunda* Rong (Zeng et al., 2016)
- *Hirnantia sagittifera fecunda* Rong (Rong, 1979)
- *Hirnantia sagittifera* morph. Bohemia (Zeng et al., 2016)
- *Hirnantia sagittifera* morph. Poland (Zeng et al., 2016)
- *Hirnantia sagittifera* M'Coy (Rong, 1984)



- ▲ Meitan, S. China This study
- Yichang S. China (Rong 1984; Zeng et al., 2016)
- Zunyi, S. China (Rong et al., 1974; Rong, 1979)
- ▢ Tibet, China (Rong and Xu, 1987)
- South Ougarta, Algeria (Mergl, 1983)
- × Kazakhstan (Nikitin 1980)
- San Juan, Argentina (Benedetto, 1986)
- ◇ Estonia (Hints et al., 2012)
- Wales (Temple 1965; Walmsley et al., 1969; Brenchley et al., 2006)
- ▲ Oslo, Norway (Cocks, 1982)
- ◇ Spain (Villas et al., 1999)
- ▽ Bohemia (Havlicek, 1977)
- Västergötland, Sweden (Bergström, 1968)

