

1 **Two new species and the molecular phylogeography of the**
2 **freshwater crab genus *Bottapotamon* (Crustacea: Decapoda:**
3 **Brachyura: Potamidae)**

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33 **Abstract**

34 *Bottapotamon chenzhouense* sp. n. and *B. luxiense* sp. n. are described from Hunan
35 Province and Jiangxi Province, respectively. These species both have diagnostic features of the
36 genus *Bottapotamon* and discernible characteristics as new species. *B. chenzhouense* sp. n. can
37 be distinguished from congeners by features such as the G1, which has a fold covering the
38 surface of the entire subterminal article with a transparent distal region. *B. luxiense* sp. n. has an
39 elliptical carapace, and a sturdy and blunt terminal article of G1. The molecular phylogeny and
40 biogeography of the genus *Bottapotamon* (Decapoda: Brachyura: Potamidae) were studied, using
41 mitochondrial cytochrome oxidase I, 16S rRNA and nuclear histone H3 gene fragments. The
42 results support the assignment of the two new species to the genus *Bottapotamon*. In addition,
43 the divergence time of the genus *Bottapotamon* was estimated to be 3.49-1.08 mya, which
44 coincided with various vicariant and dispersal events that occurred in the geological area where
45 the genus *Bottapotamon* is commonly distributed. Mountains appear to have played an important
46 role in the distribution of the genus. The Wuyi Mountains gradually formed offshore and inland
47 of southeastern China by the compression of the Pacific plate and the Indian plate in the
48 Neogene-Quaternary, and the Luoxiao Mountains formed continuously in the continued forming
49 in the north-south direction because of neotectonic movement. Thus, the geographical
50 distribution pattern of the genus *Bottapotamon* is also established gradually.

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52 Please be consistent:

53 1) All scientific names must be in italics or underlined for example:

54 *Bottapotamon nanan* or Bottapotamon nanan,

56 2) The author(s) and year are not in italics,

58 3) If there are 2 authors, decide if you want to use 'and' or '&'.

60 4) If there are more than 3 authors, use 'et al.' after the first author. There is n need
61 to type out all the names of all the authors in the text.

63 5) Please be consistent about et al or *et al* in the text.

65 **Introduction**

66 The genus *Bottapotamon* is a unique genus of freshwater crabs from the China mainland. In
67 1997, three species of the genus *Malayopotamonon* (*Bott*, 1967; *Cheng*, *YZ*, *JX* Lin & *XQ* Luo,
68 *et al.*, 1993; *Dai et al.*, 1979) and one new species were identified as *Bottapotamon* on the basis
69 of its morphological characteristics, such as the form of carapace and first gonopod (G1) (*Türkey*
70 & *Dai*, 1997). Until the current study, the genus *Bottapotamon* contained *B. fukiense*, *B.*

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74 *engelhardti*, *B. yonganense*, *B. lingchuanense* (Türkay & Dai, 1997), *B. youxiense* (Cheng *et al.*,
75 2010) and *B. nanan* (Zhou *et al.*, 2008).

76 The relatively low fecundity and poor dispersal abilities of freshwater crabs (Daniels *et al.*,
77 2003; Yeo *et al.*, 2008) mean that these crabs are easily isolated by barriers such as mountains or
78 seas. Geographically isolated populations then become genetically distinct and result in allopatric
79 speciation (Shih *et al.*, 2006; Yeo *et al.*, 2007). In mainland China, the distribution of the genus
80 *Bottapotamon* is restricted within the area of the Wuyi Mountain Range; *B. engelhardti*, *B.*
81 *yonganense*, *B. youxiense* and *B. nanan* are distributed east of the Wuyi Mountain Range, *B.*
82 *fukiense* occurs on both sides of the Wuyi Mountains (Fujian and Jiangxi Provinces), and only *B.*
83 *lingchuanense* has been isolated in the Nanling Mountain Range (Dai, 1997) (Fig. 1). The
84 geographic barrier separating the Wuyi Mountains from the Nanling Mountains is the Luoxiao
85 Mountain Range, which is the highest range in the area, exceeding 2120 m in height (Gong ~~HL~~,
86 Zhuang WY, Liao WB, *et al.*, 2016). The terrain the genus *Bottapotamon* now inhabits is
87 geologically relatively stable and experienced little orogenic activity during the Cenozoic (Yi,
88 1996; Zhou & Li, 2000). Therefore, we hypothesize that the current distribution of the genus
89 *Bottapotamon* in mainland China was caused by the emergence of these mountains.

90 By organizing the existing specimens deposited at the Department of Parasitology of the
91 Medical College of Nanchang University (NCU MCP) and the newly collected specimens, the
92 authors (or who exactly? the first author) discovered two new species in Chenzhou City, Hunan
93 Province, and Luxi County, Jiangxi Province. This paper compares the morphological features of
94 eight species including two new species of the genus *Bottapotamon*, as well as 16S rRNA
95 (Crandall *et al.* 1996), mtDNA COI (Folmer *et al.*, 1994) and nuclear histone H3 (Colgan *et al.*,
96 1998) gene fragments were used to support the establishment of new species in the genus
97 *Bottapotamon*. The phylogenetic relationship, distribution pattern and possible association with
98 major geological and historical events are also discussed.

99

100

101 **Materials & Methods**

102 **Specimen collection**

103 Specimens from Jiangxi, Zhejiang, Fujian and Guangxi, were recently collected and
104 preserved in 95% ethanol. The remaining specimens used in this study were from and deposited
105 at the Department of Parasitology of the Medical College of Nanchang University (NCU MCP),
106 Jiangxi Province, China. The author compared specimens with holotypes of the Institute of
107 Zoology, Chinese Academy of Sciences. All 26 specimens were used for mtDNA COI, 16S
108 rRNA and histone H3 gene fragment amplification (Table 1).

109

110 **Phylogenetic analyses and Divergence time estimation**

111 Genomic DNA was extracted from leg muscle tissue with an OMEGA EZNA™ Mollusc
112 DNA Kit. The 16S rRNA, mtDNA COI, and histone H3 regions were selected for amplification
113 by polymerase chain reaction (PCR) (Table 2). The amplification products were sent to the
114 Beijing Genomics Institute for bidirectional sequencing, and the sequencing results were spliced
115 manually to obtain the sequence data. DNA sequences of *B. yonganense* specimens collected
116 from the suburb of Sanming City, Fujian Province, China, could not be amplified due to poor
117 preservation.

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124 The sequences of four individuals with the same primer sequences were selected from
125 National Center for Biotechnology Information (NCBI) database, as the outgroups
126 (*Candidiopotamon rathbunae* (GenBank accession numbers: mtDNA COI-AB290649, 16S
127 rRNA-AB208609, histone H3-AB290668), *Geothelphusa dehaani* (GenBank accession
128 numbers: mtDNA COI-AB290648, 16S rRNA-AB290630, histone H3-AB290667),
129 *Himalayapotamon atkinsonianum* (GenBank accession numbers: mtDNA COI-AB290651, 16S
130 rRNA-AB290632, histone H3-AB290670), and *Ryukyum yaeyamense* (GenBank accession
131 numbers: mtDNA COI-AB290650, 16S rRNA-AB290631, histone H3-AB290669)). After
132 comparing and selecting the conservative regions, each sequence was 1323 bp in length.
133 According to the Akaike information criterion (AIC), MrMTGui: ModelTest and MrModelTest
134 (phylogenetic analysis using parsimony (PAUP)) determined the best models was GTR+I+G;
135 MEGA 6.06 (Tamura et al., 2013) was used to establish a phylogenetic tree based on the
136 maximum likelihood (ML) (Trifinopoulos et al., 2016). The Bayesian inference (BI) tree was
137 established using MrBayes (Ronquist & Huelsenbeck, 2003).

138 The divergence times of genus *Bottapotamon* were estimated from the combined 16S rRNA
139 and mtDNA COI sequences, based on the Bayesian evolutionary analysis sampling trees
140 (BEAST) program, and four calibration points were used. The Potamidae family has divided into
141 two major subfamilies, Potamiscinae and Potaminae, estimated to have a divergence time of
142 20.9-24.7 mya, which was set as calibration point 1 in our study (Shih et al., 2010). The
143 Parathelphusidae subfamily, *Somanniathelphusa taiwanensis*, which is distributed in Taiwan
144 Island and separated from *Somanniathelphusa amoyensis*, which is distributed in Fujian
145 Province, approximately 0.27-1.53 mya (Jia et al., 2018). The results are consistent with the
146 quaternary glacial period and interglacial period and agree with the separation of Taiwan Island
147 and Fujian Province; this time point was set as calibration point 2. In the geological area where
148 genus *Bottapotamon* is distributed, the Wuyi Mountains gradually formed by the compression of
149 the Pacific plate and the Indian plate in the Neogene-Quaternary (1.64-23.3 mya) (Li, 1984);
150 this time point was set as calibration point 3. A Yule speciation model was constructed for
151 speciation within the genus *Bottapotamon*. We used a GTR+G model with parameters obtained
152 from MrMTGui: ModelTest and MrModelTest (PAUP) for each gene. Seventeen independent
153 MCMC chains were run for 200,000,000 generations, and every 20,000 generations were
154 sampled. The convergence of the 17 combined chains was determined by the evolutionary stable
155 strategy (ESS) (>200 as recommended) for each parameter in Tracer after the appropriate burn-in
156 and cutoff (default of 10% of sampled trees). Trees in the 17 chains were combined using
157 LogCombiner (v. 1.6.1, distributed as part of the BEAST package) and were assessed using
158 TreeAnnotator (v. 1.6.1, distributed as part of the BEAST package). A chronogram was
159 constructed by FigTree.

161 Nomenclatural note

162 The electronic version of this article in Portable Document Format (PDF) will represent a
163 published work according to the International Commission on Zoological Nomenclature (ICZN),

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167 and hence the new names contained in the electronic version are effectively published under that
168 Code from the electronic edition alone. This published work and the nomenclatural acts it
169 contains have been registered in ZooBank, the online registration system for the ICZN. The
170 ZooBank LSIDs (Life Science Identifiers) can be resolved and the associated information viewed
171 through any standard web browser by appending the LSID to the prefix <http://zoobank.org/>. The
172 LSID for this publication is: [urn: lsid: zoobank.org: pub:211926FF-6950-4DFE-95C4-
173 F5247CA9E0BA]. The online version of this work is archived and available from the following
174 digital repositories: PeerJ, PubMed Central and CLOCKSS.

175

176 Results

177 Systematics

178

179 Potamidae Ortmann, 1896

180 *Bottapotamon* Türkay et Dai, 1997

181 ~~*Bottapotamon chenzhouense* sp.n., *Bottapotamon luxiense* sp.n.~~

182

183 *Bottapotamon chenzhouense* sp. n. (Fig. 2-6)

184 urn: lsid zoobank. org: art: E43C4BBB-E429-4C17-8ACD-E4295F426BCB

185

186 Materials examined

187 Holotype: 1 ♂ (25.72 × 15.69 mm) (NCU MCP 643), Huangcao Village, Chenzhou City,
188 Hunan Province, China, 25°39'24.60"N, 113°30'4.07"E, 141 m asl. Coll. Dingmei Luo, July 26th
189 2006. Paratypes: 1 ♀ (18.7 × 13.7 mm) (NCU MCP 643), the same data as the holotype.

190

191 Comparative materials

192 *B. fukiense* (Türkay & Dai, 1997): 4 ♂ (25.21 × 15.02 mm, 25.03 × 14.97 mm) (NCU MCP
193 4089), Xiapu Village, Ningde County, Fujian; (26.08 × 15.45 mm) (NCUMCP4156), Shangshan
194 Village, Zhenghe County, Fujian; (26.08 × 15.45 mm) (NCUMCP4090), Siqian Village,
195 Shouning County, Fujian; 1 ♀ (26.01 × 15.57 mm) (NCU MCP 4156), Shangshan Village,
196 Zhenghe County, Fujian. *B. engelhardti* (Türkay & Dai, 1997): 5 ♂ (23.01 × 14.03 mm, 24.68 ×
197 15.69 mm, 24.81 × 15.87 mm, 25.02 × 15.47 mm) (NCU MCP 4157), Tansan Village, Youxi
198 County, Fujian; (25.21 × 15.16 mm) (NCU MCP 4091), Chimu Village, Youxi County, Fujian; 1
199 ♀ (26.01 × 16.35 mm) (NCU MCP 4091), Chimu Village, Youxi County, Fujian. *B. yonganense*
200 (Türkay & Dai, 1997): 1 ♂ (25.87 × 15.95 mm) (NCU MCP 4096), Sanming City, Fujian; *B.*
201 *lingchuanense* (Türkay & Dai, 1997), 3 ♂ (24.78 × 14.89 mm, 25.04 × 15.06 mm) (NCU
202 MCP4076), Yuanpu Village, Gongcheng County, Guangxi Zhuang Autonomous Region; (25.25
203 × 15.11 mm) (NCU MCP 3281), Bindong Village, Lingchuan County, Guangxi Zhuang
204 Autonomous Region; 3 ♀ (25.48 × 14.92 mm, 25.14 × 15.09 mm, 25.78 × 14.79 mm), (NCU
205 MCP 3281), Bindong Village, Lingchuan County, Guangxi Zhuang Autonomous Region. *B.*
206 *youxiense* (Cheng et al., 2010): (24.91 × 15.72 mm) (NCU MCP 4092), (25.11 × 15.16 mm)
207 (NCU MCP 4158), (25.34 × 15.52 mm) (NCU MCP 4059), Xiwei Village, Youxi County,
208 Fujian; 1 ♀ (26.04 × 14.92 mm) (NCU MCP 4059), Xiwei Village, Youxi County. *B. nanan*

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209 (*Zhou et al., 2008*):6♂ (26.26 × 15.72 mm, 26.03 × 15.55 mm) (NCU MCP4090), Siqian
210 Village, Shouning County, Fujian; (26.15 × 15.43 mm, 26.16 × 15.32 mm) (NCU MCP4038),
211 Yongjia County, Zhejiang; (26.25 × 15.36 mm, 26.06 × 15.52 mm) (NCU MCP4039), Yongjia
212 County, Zhejiang; 1♀(26.11 × 15.12 mm) (NCU MCP4039), Yongjia County, Zhejiang.

213

214 **Diagnosis**

215 Carapace approximately about 1.6 times broader than long, dorsal surface gently convex
216 longitudinally and transversely; cervical groove indistinct, H-shaped groove between gastric and
217 cardiac regions distinct. Male pleon triangular, sixth somite width 2.5 times length; telson
218 triangular, tip rounded, with proximal width 1.7 times length. G1 long, tip of terminal segment
219 reaching suture between thoracic sternites 4, 5 in situ; subterminal segment 1.3 times as long as
220 terminal segment; terminal segment slightly elongated inward, distal part of terminal segment
221 elongated with ventrally directed semicircular lobe. Female vulvae partially exposed anteriorly to
222 the thoracic sternites 5, 6 in situ, ovate, deep, posteromesial margin with a low raised rim,
223 opened inward.

224

225 **Description**

226 Carapace approximately about 1.6 times broader than long, dorsal surface gently convex,
227 surface slightly pitted. Cervical groove shallow, indistinct. H-shaped groove between the gastric
228 region and cardiac region shallow but distinct. Postfrontal lobe blunt, separated medially by a Y-
229 shaped groove extending to frontal region; postorbital crest indistinct, postorbital region slight
230 concave. Frontal region deflexed downwards. Dorsal orbital margin ridged, external orbital angle
231 triangular outer margin smooth; Anterolateral margin cristate, epibranchial tooth pointed,
232 indistinct, clearly demarcated from external orbital tooth (*Fig. 2*).

233 Third maxilliped merus about 1.3 times as broad as long; Ischium about 1.5 times as long as
234 broad, with distinct median sulcus; exopod reaching proximal third of merus length, without
235 flagellum (*Fig. 3A*).

236 The male sternum is relatively flat and has granular small pits. The first section is triangular
237 and the second to fourth sections are fused. The interruption between sternite sutures is
238 intermediate in depth and wide. The median longitudinal sutures of sternites 7/8 are shorter; the
239 tubercle of abdominal lock is on the medial side of the fifth male ventral nail (*Fig. 4*).

240 Cheliped slightly unequal; margins crenulated; carpus with sharp spine on inner distal
241 angle, with spinule at base; outer surface of manus with convex granules, manus about 1.6 times
242 as long as high, slightly longer than movable finger, gape wide when fingers closed, cutting edge
243 lined with low teeth (*Fig. 3C*).

244 Ambulatory legs slender; margins of propodus smooth; last leg with propodus about 1.8
245 times as long as broad, slightly shorter than dactylus (*Fig. 3B*).

246 G1 slender, a fold covering the surface of the entire subterminal article with a transparent
247 distal region. tip of terminal segment slightly reaching beyond sternal press-button in situ,
248 subterminal segment about 1.3 times as long as terminal segment. G1 slightly curved

249 ventrolaterally; distal part of G1 terminal segment distinctly broader than proximal part. G2
250 subterminal segment about 2.3 times as long as terminal segment (Fig. 5A, 6A).

251

252 **Remarks**

253 The new species fits well within the morphological definition of the hitherto monotypic
254 *Bottapotamon* (Türkay & Dai, 1997; Cheng et al., 2010; Zhou et al., 2008): G1 is slender, tip of
255 terminal segment reaching suture between thoracic sternites 4, 5 in situ; terminal segment
256 slightly elongated inward (Table. 3). Nonetheless, the new species can be distinguished from
257 comparative specimens, by the Carapace surface gently convex, cervical groove indistinct; H-
258 shaped groove shallow but distinct. epibranchial tooth pointed and indistinct, third maxilliped
259 without flagellum; chelipeds carpus with sharp spine on inner distal angle; G1 is sturdy and blunt
260 (Table. 3).

261

262 **Etymology**

263 The species is named after the type locality: Chenzhou city, Hunan Province, China.

264

265 **Distribution**

266 *B. chenzhouense* sp. n. was found under stones in a mountain stream in Huangcao village,
267 Chenzhou city, Hunan Province, China.

268

269 ***Bottapotamon luxiense* sp.n. (Fig. 5-10)**

270 urn: lsid:zoobank.org:art:1C1CC520-193A-405E-9A2D-DC79E7D4AA87.

271

272 **Materials examined**

273 Holotype: 1 ♂ (18.72×15.69 mm) (NCU MCP 4200), Yixiantian Wugongshan Mountain,
274 Luxi county, Pingxiang city, Jiangxi Province, China, 27°28'56.16"N, 114°10'27.51"E, 1331 m
275 asl. Coll. Jiexin Zou, May 6th 2019. Paratypes: 1 ♂ (19.22 × 16.38 mm) (NCU MCP 4200).
276 Others: 12 ♀ (16.7 × 15.7 mm, 15.41×15.36 mm, 14.23×12.98 mm, 15.63×14.52 mm,
277 16.13×15.86 mm, 16.23×14.97 mm, 13.65×12.33 mm, 14.56×13.15 mm, 15.27×14.10 mm,
278 16.02×15.43 mm, 15.89×15.01 mm, 13.13×12.46 mm) (NCU MCP 4200), 12 ♂ (15.66×13.89
279 mm, 14.21×13.11mm, 13.69×12.01 mm, 14.23×13.69 mm, 15.17×14.31 mm, 14.19×13.69 mm,
280 14.69×13.54 mm, 14.73×13.52 mm, 12.87×11.36 mm, 13.00×12.13 mm, 13.58×12.29 mm,
281 15.26×14.36 mm) (NCU MCP 4200), same data as holotype.

282

283 **Comparative materials**

284 Same as *Bottapotamon chenzhouense* sp. n.

285

286 **Diagnosis**

287 Carapace about 1.4 times broader than long, dorsal surface gently convex longitudinally and
288 transversely; cervical groove distinct, H-shaped groove between gastric and cardiac regions
289 distinct. Male pleon triangular, sixth somite width 2.3 times length; telson triangular, tip
290 rounded, with proximal width 1.6 times length. G1 long, tip of terminal segment reaching suture

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292 between thoracic sternites 4, 5 in situ; subterminal segment 1.2 times as long as terminal
293 segment; terminal segment slightly elongated inward, distal part of terminal segment elongated
294 with ventrally directed semicircular lobe. Female vulvae partially exposed anteriorly to the
295 thoracic sternites 5, 6 in situ, ovate, deep, posteromesial margin with a low raised rim, opened
296 inward.

297
298 **Description**

299 Carapace nearly ellipse, about 1.4 times broader than long, dorsal surface gently convex,
300 surface slightly pitted. Cervical groove distinct. H-shaped groove between the gastric region and
301 cardiac region shallow but distinct. Postfrontal lobe blunt; postorbital crest indistinct, postorbital
302 region slight concave. Frontal region deflexed downwards. Dorsal orbital margin ridged, external
303 orbital angle triangular, outer margin smooth. Anterolateral margin cristate, epibranchial tooth
304 pointed (*Fig. 7*).

305 Third maxilliped merus about 1.4 times as broad as long, trapezoidal; ischium about 1.5
306 times as long as broad, with distinct median sulcus; exopod reaching proximal third of merus
307 length, with flagellum (*Fig. 8A*).

308 Thoracic sternum pitted; sternites 1, 2 completely fused to form triangular structure; sternites
309 2, 3 separated by continuous suture; boundary between sternites 3, 4 present, indistinct. Sterno-
310 pleonal cavity broad, shallow, with narrow median interruption in sutures 4/5, 5/6, 6/7; median
311 line between sternites 7, 8 moderately long (*Fig. 9*).

312 The male sternum is relatively flat and has granular small pits. The first section is triangular
313 and the second to fourth sections are fused. The interruption between sternite sutures is medium
314 in depth and wide. The median longitudinal sutures of sternites 7/8 are shorter; the tubercle of
315 abdominal lock is on the medial side of the fifth male ventral nail (*Fig. 6B*).

316 Chelipeds slightly unequal; margins crenulated; outer surface of manus with convex
317 granules, manus about 1.5 times as long as high, slightly longer than movable finger, gape wide
318 when fingers closed, cutting edge lined with low teeth (*Fig. 8B*).

319 Ambulatory legs slender; margins of propodus smooth; last leg with propodus about 1.7
320 times as long as broad, slightly shorter than dactylus (*Fig. 8C*).

321 G1 is sturdy and blunt, tip of terminal segment slightly reaching beyond sternal press-button
322 in situ, subterminal segment about 1.4 times as long as terminal segment. G1 slightly curved
323 ventrolaterally; distal part of G1 terminal segment distinctly broader than proximal part. G2
324 subterminal segment about 2.2 times as long as terminal segment (*Fig. 5B, 6B*).

325
326 **Remarks**

327 The new species fits well within the morphological definition of the hitherto monotypic
328 *Bottapotamon* (*Türkay & Dai, 1997; Cheng et al., 2010; Zhou et al., 2008*): cervical groove
329 indistinct, H-shaped groove between gastric and cardiac regions distinct, G1 long, tip of terminal
330 segment reaching suture between thoracic sternites 4, 5 in situ; terminal segment slightly
331 elongated inward (*Table. 3*). Nonetheless, the new species can be distinguished from
332 comparative specimens, by the carapace surface gently convex, cervical groove shallow and

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338 indistinct; H-shaped groove shallow but distinct. epibranchial tooth pointed and indistinct, third
339 maxilliped without flagellum; chelipeds carpus with sharp spine on inner distal angle, with
340 spinule at base; G1, a fold covering the surface of the entire subterminal article with a
341 transparent distal region (Table. 3).

342

343 **Etymology**

344 The species is named after the type locality: Yixiantian Wugongshan Mountain, Luxi county,
345 Pingxiang city, Jiangxi Province, China.

346

347 ~~Living color~~ **Live colouration**

348 The dorsal surfaces of the carapace and pereopods are dark purple-red, and the joints of the
349 cheliped merus and carpus the ambulatory legs are bright red. The inner surface of the
350 immovable finger and distal part of the movable finger are almost milky.

351

352 **Distribution**

353 *B. luxiense* sp. n. was found under stones in a mountain stream in Yixiantian Wugongshan
354 Mountain, Luxi county, Pingxiang city, Jiangxi Province, China (Fig. 10).

355

356 **Ecology**

357 *B. chenzhouense* sp. n. and *B. luxiense* sp. n. were collected in the Luoxiao mountains. This
358 region has a humid subtropical monsoon climate and is in the Xiangjiang River and Ganjiang
359 River watershed, which has rich biodiversity (Wang, 1998). Similar to the natural habitat of other
360 *Bottapotamon* species, *B. chenzhouense* sp. n. and *B. luxiense* sp. n. can be found under small
361 rocks in sandy creek beds in narrow mountain streams or highway drains with clear, slow
362 flowing and cool water surrounded by dwarf shrubs or grasses (Fig. 10).

363

364 **Phylogenetic analyses and Divergence time estimation**

365 The combined mtDNA COI, 16S rRNA and nuclear histone H3 phylogenetic trees were
366 constructed by ML analysis, and the corresponding support values were calculated by ML and BI
367 analyses, both of which had high support values. The results showed that the genus
368 *Bottapotamon* is monophyletic, and confirmed that *B. chenzhouense* sp. n. and *B. luxiense* sp. n.
369 are new species of genus *Bottapotamon* and supported the relationship of the genus
370 *Bottapotamon* (Fig. 11). *B. engelhardti*, *B. yonganense* and *B. nanan*, which are mostly
371 distributed in the Wuyi Mountain Range, form a clade; *B. luxiense* sp. n. forms a sister clade to
372 the clade of *B. engelhardti*, *B. yonganense* and *B. nanan*. The next sister clade is composed of *B.*
373 *chenzhouense* sp. n., which is distributed in the Luoxiao Mountain Range, and the furthest sister
374 clade is composed of *B. lingchuanense*, which is distributed in the Nanling Mountain Range. *B.*
375 *fukiense* and *B. youxiense* are also distributed in the Wuyi Mountain Range, but they do not
376 assemble with *B. engelhardti*, *B. yonganense* and *B. nanan*.

377 The divergence time estimation results are consistent with the four calibration points. The
378 genus *Bottapotamon* diverged approximately 3.49-1.08 mya, *B. fukiense* and *B. youxiense*
379 diverged 1.96 mya (95% confidence interval =2.65-1.31 mya), *B. luxiense* diverged 1.90 mya

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380 (95% confidence interval =2.05-1.09 mya), *B. lingchuanense* and *B. chenzhouense* sp. n.
381 diverged 1.51 mya (95% confidence interval =1.6-0.7 mya); *B. engelhardti* and *B. nanan*
382 diverged 1.08 mya (95% confidence interval =1.76-0.80 mya).(Fig. 12)

383

384 Discussion

385 In mainland China, the genus *Bottapotamon* is primarily distributed in the Wuyi Mountain
386 Range area; *B. luxiense* sp. n., *B. youxiense*, *B. nanan*, *B. engelhardti* and *B. yonganense* are
387 restricted within an area east of the Wuyi Mountain Range (Fig. 1). There is no record of any of
388 these five species in Jiangxi, despite extensive surveys of this area by the authors and their
389 colleagues over many years (Dai, 1999; Shi, 2012). The altitude of the Wuyi Mountain Range is
390 clearly high enough to prevent these species from reaching Jiangxi. *B. fukiense* occurs on both
391 sides of the Wuyi Mountain Range (Fujian and Jiangxi Provinces) and is able to disperse across
392 these mountains. The divergence time of *B. fukiense* and *B. youxiense* is 1.96 mya (95%
393 confidence interval =2.65-1.31 mya) (Fig. 12), which agrees well with records of the Pacific
394 plate and Indian plate extrusion in the Neogene-Quaternary (1.64-23.3 mya) (Li, 1984).
395 Therefore, these geological events may explain the distribution pattern of the genus
396 *Bottapotamon* in the Wuyi Mountain Range. The ancestor of the genus *Bottapotamon* originated
397 in an area close to the Wuyi Mountains, as the Wuyi Mountain Formation and smaller-scale
398 mountain deformations resulted in sufficient geographic barriers to isolate populations; thus, the
399 two species-groups were separated by the Wuyi Mountains.

400 In the Nanling mountain range, unique karst formation and the south Asian subtropical
401 humid monsoon climate conditions provide a good living environment for all types of wildlife,
402 including freshwater crabs. However, only one species of the genus *Bottapotamon*, *B.*
403 *lingchuanense*, was isolated in this area, and there is an 830 km gap between *B. lingchuanense*
404 and other species distributed within the Wuyi Mountain Range (Fig. 1), which has always been
405 the focus of researches on the genus *Bottapotamon*. This study reports two new species of genus
406 *Bottapotamon*, *B. chenzhouense* sp. n., Which was first discovered in Chenzhou City, Hunan
407 Province, in south of Luoxiao Mountains, and *B. luxiense* sp. n., which is distributed in north of
408 the Luoxiao Mountains (Fig. 1). Divergence time estimation results suggested that *B.*
409 *chenzhouense* sp. n., *B. luxiense* sp. n., and *B. lingchuanense* were isolated at almost the same
410 time (*B. luxiense* sp. n. diverged 1.90 mya, and *B. lingchuanense* and *B. chenzhouense* sp. n.
411 diverged at 1.51 mya) (Fig. 12). The authors speculated that the Luoxiao Mountains
412 continuously rose due to neotectonic movement and gradually became the Xiangjiang River and
413 Ganjiang River watershed (Wang, 1998). The ancestors of the genus *Bottapotamon* occurred on
414 both sides of the Luoxiao Mountains during the mountains formation process, and under the
415 influence of karst landforms and the Danxia landform, gradually isolated *B. luxiense* sp. n., *B.*
416 *chenzhouense* sp. n and *B. lingchuanense* was gradually isolated. In addition, the climatic
417 conditions in this area are ideal for *Bottapotamon*. The authors speculate that many new species
418 of the genus *Bottapotamon* are likely to exist in the region from the Wuyi Mountain Ranges to
419 the Nanling Mountain Range.

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422 Conclusions

423 *B. chenzhouense* sp. n. and *B. luxiense* sp. n., two new species from the Luoxiao Mountains
 424 were reported in this paper. These two new species compensated for the geographical gap in the
 425 genus *Bottapotamon*, and confirm the independence and intra- and interspecific relationships of
 426 genus *Bottapotamon*. Combined with estimates of divergence times, this paper suggests that the
 427 genus *Bottapotamon* was formed at 3.49-1.08 mya. Molecular evidence further supports the
 428 scientific hypothesis of the authors that genus *Bottapotamon* originated on both sides of the
 429 Wuyi Mountains and Luoxiao Mountains. In the geological area where the genus *Bottapotamon*
 430 is distributed, the Wuyi Mountains gradually formed offshore and inland of southeastern China
 431 by the compression of the Pacific plate and the Indian plate in the Neogene-Quaternary, and the
 432 Luoxiao Mountains formed continuously in the north-south direction because of neotectonic
 433 movement. Thus, the geographical distribution patterns of the genus *Bottapotamon* formed
 434 gradually.

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437 ADDITIONAL INFORMATION AND DECLARATIONS

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444 Data Availability

445 Regarding data availability: all specimens in this study are housed in the permanent
 446 collections at the Department of Parasitology, Medical College of Nanchang University (NCU
 447 MCP), and the raw DNA data are included in the supplemental files.

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450 2) please be consistent with the format of your reference, some journals are in
 451 abbreviated form and some are in full.

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454 References

- 455 **Bott, R. 1967.** Potamoniden aus Ost-Asien (*Parapotamon* de Man, *Sinopotamon* N.
 456 gen. *Candidiopotamon* n. gen., *Geothelphusa* Stimpson) (Crustacea, Decapoda).
 457 *Senckenbergiana biol.*, Frankfurt, 48(3): 203-220, pls. 7-10, Figs. 1-13.
- 458 **Cheng YZ, JX Lin & XQ Luo. 1993.** A new species of crab of the genus
 459 *Malayopotamon* (Decapoda: Isolapotamidae). *Acta Zootax. Sinica*, 18(4): 412-
 460 416.

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- 465 **Dai AY, GX Chen, YZ Song, PF Fan, YG Lin & YQ Zeng. 1979.** On New Species of
466 Freshwater Crabs Harbouring Metacercariae of Lung Flukes. *Acta Zootax.*
467 *Sinica*, 4(2): 122-121, 1 pl.
- 468 **Türkay M & Dai AY. 1997.** Review of the Chinese freshwater crabs previously placed in
469 the genus *Malayopotamon* Bott, 1968 (Crustacea : Decapoda : Brachyura :
470 Potamidae). *The Raffles bulletin of zoology*. 45: 189-207.
- 471 **Cheng YZ, Lin GH, and Li YS. 2010.** Two New Species of Freshwater Crabs
472 (Decapoda : Potamidae) Serving as Intermediate Hosts of *Paragonimus* in
473 Fujian, China. *Chinese Journal of Parasitology and Parasitic Diseases* 28:241-
474 245.
- 475 **Zhou X, Zhu C, and Naruse T. 2008.** *Bottapotamon nanan*, a New Species of
476 Freshwater Crab (Decapoda, Brachyura, Potamidae) from Fujian Province,
477 China. *Crustaceana* 81:1389-1396.
- 478 **Daniels SR, Gouws G, Stewart BA, Coke M. 2003.** Molecular and morphometric
479 data demonstrate the presence of cryptic lineages among freshwater crabs
480 (Decapoda: Potamonautidae: Potamonautes) from the Drakensberg Mountains,
481 South Africa. *Biol. J. Linn. Soc.* 78, 129–147.
- 482 **Yeo DCJ, Ng PKL, Cumberlidge N, Magalhães C, Daniels SR, Campos MR. 2008.**
483 Global diversity of crabs (Crustacea: Decapoda: Brachyura) in freshwater.
484 *Hydrobiologia*. 595, 275–286.
- 485 **Shih HT, Hung HC, Schubart CD, Chen CA, Chang HW. 2006.** Intraspecific genetic
486 diversity of the endemic freshwater crab *Candidiopotamon rathbunae*
487 (Decapoda, Brachyura, Potamidae) reflects five million years of geological history
488 of Taiwan. *J. iogeogr.* 33, 980–989.
- 489 **Yeo DCJ, Shih HT, Meier R, Ng PKL. 2007.** Phylogeny and biogeography of the
490 freshwater crab genus *Johora* (Crustacea: Brachyura: Potamidae) from the
491 Malay Peninsula, and the origins of its insular fauna. *Zool. Scr.* 36, 255–269.
- 492 **Dai AY. 1999.** Fauna Sinica Arthropoda: Crustacea: Malacostraca: Decapoda:
493 Parathelphusidae, Potamidae. *Science Press, Beijing* (in Chinese).
- 494 **Gong HL, Zhuang WY, Liao WB. 2016.** Comprehensive scientific investigation of
495 biodiversity in Luojing Mountain area. *Chinese scientific and technological*
496 *achievements*. 7(22):9-10.
- 497 **Yi MC. 1996.** Cenozoic para-orogenic movement in China. *Acta Geol.* 17, 249 - 255 (in
498 Chinese).
- 499 **Zhou XM, Li WX. 2000.** Origin of late Mesozoic igneous rocks in Southeastern China:
500 implications for lithosphere subduction and underplating of mafic magmas.
501 *Tectonophysics*. 326: 269-287.
- 502 **Crandall KA, Fitzpatrick JF, and Faith D. 1996.** Crayfish Molecular Systematics:
503 Using a Combination of Procedures to Estimate Phylogeny. *Systematic Biology*
504 45:1-26.
- 505 **Folmer O, Black M, Hoeh W, Lutz R, and Vrijenhoek R. 1994.** DNA primers for
506 amplification of mitochondrial cytochrome c oxidase subunit I from diverse
507 metazoan invertebrates. *Mol Mar Biol Biotechnol* 3:294-299.
- 508 **Colgan DJ, Mclauchlan A, Wilson GDF, Livingston SP, Edgecombe GD,**
509 **Macaranas J, Cassis G, and Gray MR. 1998.** Histone H3 and U2 snRNA DNA

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512 sequences and arthropod molecular evolution. *Australian Journal of Zoology*
513 46:419-437.

514 **Tamura K, Stecher G, Peterson D, Filipinski A, and Kumar S. 2013.** MEGA6:
515 Molecular Evolutionary Genetics Analysis Version 6.0. *Molecular Biology and*
516 *Evolution* 30:2725-2729.

517 **Trifinopoulos J, Nguyen LT, Haeseler Av, and Minh BQ. 2016.** W-IQ-TREE: a fast
518 online phylogenetic tool for maximum likelihood analysis. *Nucleic Acids Research*
519 44:W232-W235.

520 **Ronquist F, and Huelsenbeck JP. 2003.** MrBayes 3: Bayesian phylogenetic inference
521 under mixed models. *Bioinformatics* 19:1572-1574.

522 **Shih HT, Yeo DCJ, and Ng PKL. 2010.** The collision of the Indian plate with Asia:
523 molecular evidence for its impact on the phylogeny of freshwater crabs
524 (Brachyura: Potamidae). *Journal of Biogeography* 36:703-719.

525 **Jia XN, Xu SX, Bai J, Wang YF, Nie ZH, Zhu CC, Wang Y, Cai YX, Zou JX, and**
526 **Zhou XM. 2018.** The complete mitochondrial genome of *Somanniathelphusa*
527 *boyangensis* and phylogenetic analysis of Genus *Somanniathelphusa*
528 (Crustacea: Decapoda: Parathelphusidae). *Plos One* 13:e0192601-.

529 **Li ZZ. 1984.** THE Origin And Morphological Characteristics Of The Wuyi Mountain,
530 Fujian Province. *Journal of Nanjing University(Natural Sciences)*.

531 **Shi LB, Zhang XY, Zou JX, Wang Y, Li DR, Zhu CC, and Zhou XM. 2012.** Distribution
532 pattern of the freshwater crabs among Wuyi Mountains. *Journal of Nanchang*
533 *University (Natural Science)* 36:556-561.

534 **Wang CL. 1998.** Formation Of Luxiao Mountains And Development Of Its Danxia Land
535 Feature. *Journal of Xiangtan Normal University*:110-115.