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New psychropotid species (Echinodermata Holothuroidea Elasipodida) of the Western Pacific with phylogenetic analyses

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1 New psychropotid species (Echinodermata: Holothuroidea: 2 Elasipodida) of the Western Pacific with phylogenetic analysis

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12 Abstract

13 Holothurians of the family Psychropotidae are widely distributed, but remain least
14 studied deep-sea holothurians. On an expedition in the Western Pacific, six
15 psychropotid specimens were collected by the Jiaolong human operated vehicle (HOV).
16 Through morphological examination, four of them were identified as a new species,
17 *Benthodytes jiaolongi* **sp. nov.**, which was characterized as having minute papillae,
18 narrow brims, and terminal anus; the ossicles were either rods or primary crosses. The
19 remaining two specimens were identified as *Psychropotes verrucicaudatus* Xiao, Gong,
20 Kou & Li, 2019, which was first recorded at the Kyushu-Palau Ridge. The phylogenetic
21 analysis confirmed the classification status of *B. jiaolongi* and *P. verrucicaudatus*, and
22 indicated a paraphyletic relationship within the genus *Benthodytes*. The new species
23 clustered with *Benthodytes sanguinolenta* and was separated from the clade containing
24 the other *Benthodytes* species.

25

26 Key words

27 *Benthodytes*, holothurians, new record, paraphyletic group, *Psychropotes*, Western
28 Pacific

29

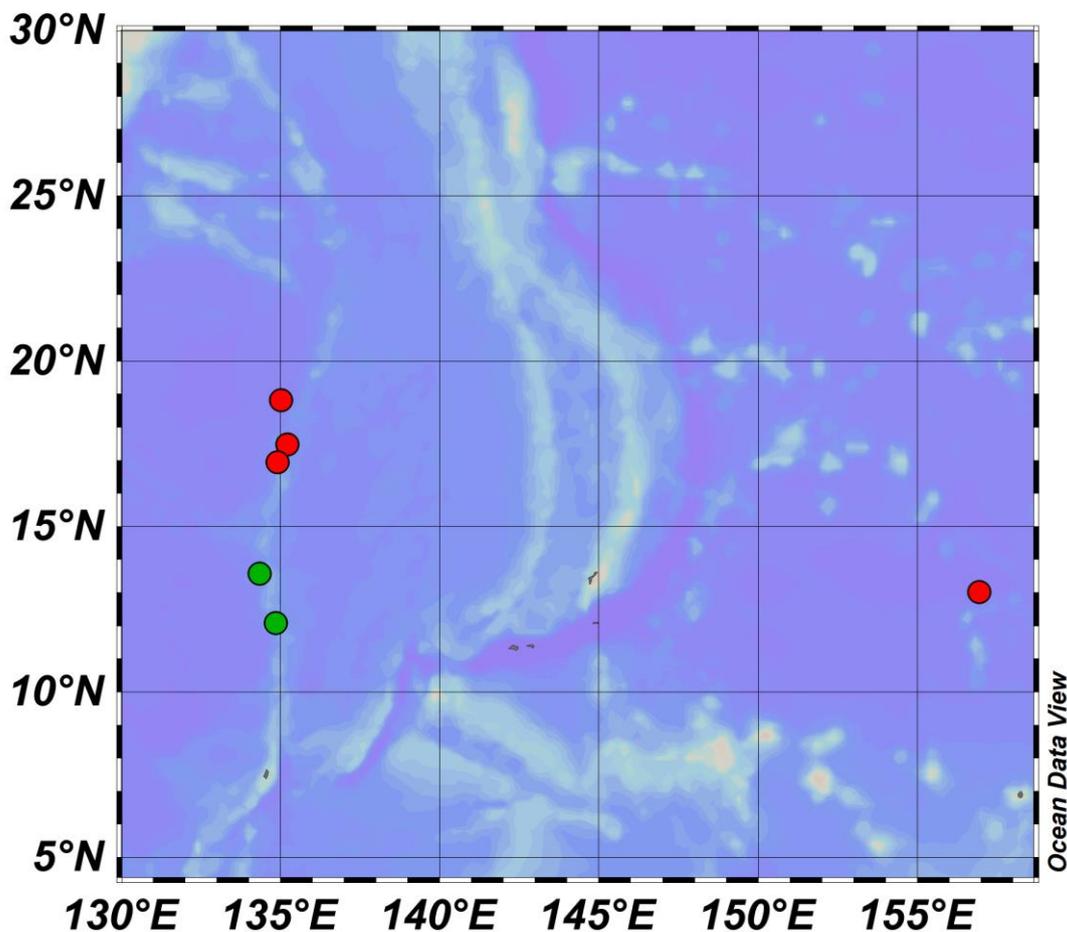
30 Introduction

31

32 Holothurians of the family Psychropotidae (Elasipodida) were first identified by
33 Théel (1882) who defined four genera of the deep-sea sea cucumbers discovered on the
34 H.M.S. Challenger Expedition. Subsequently, Hérouard (1909) and Vinogradov (1969)
35 erected *Triconus* Hérouard and *Nectothuria* Belyaev and Vinogradov, which were later
36 regarded as synonyms of *Psychropotes* by Hansen (1975). Psychropotidae comprises
37 three genera and 37 species. Hansen (1975) distinguished the three genera by the
38 presence or absence of an unpaired dorsal appendage, the position of the anus, and the
39 presence or absence of circum-oral (or post-oral) papillae. Although, taxonomists have

40 long worked on this family, Psychropotidae are still the least studied deep-sea
 41 holothurians. Thus, the phylogenetic relationships within Psychropotidae remain
 42 unclear.

43 An expedition of the Jiaolong HOV concentrated on furthering our understanding
 44 of the biodiversity, connectivity, and conservation value of the Western Pacific. During
 45 sampling, six specimens of Psychropotidae were collected from seamounts on the
 46 Kyushu-Palau Ridge and Weijia Guyot (Fig. 1). Based on an analysis of the external
 47 morphological characters and ossicles, we identified four specimens as a new species
 48 (*Benthodytes jiaolongi* **sp.nov.**) and the other two as new records *Psychropotes*
 49 *verrucicaudatus* Xiao, Gong, Kou & Li, 2019.



50
 51 **Figure 1.** Red dots show the location of *Benthodytes jiaolongi* **sp.nov.** and green dots
 52 indicated *Psychropotes verrucicaudatus*.

53

54 **Materials and Methods**

55 *Sampling and morphological observations*

56 The samples described in the present study were collected by the Jiaolong HOV at
 57 a depth of 2408-2602 m, from the Kyushu-Palau Ridge and Weijia Guyot. Before
 58 preservation, a Canon EOS 5DII camera (Canon Inc., Tokyo, Japan) was used to take
 59 photographs of the specimens on board the ship. Then, a piece of tissue was cut from

60 all specimens and frozen at -20 °C for DNA extraction. Finally, the specimens were
 61 fixed in 10% seawater formalin or 99% alcohol and deposited at the Repository of
 62 Second Institute of Oceanography (RSIO). Sodium hypochlorite was used to dissolve
 63 some body tissues (tentacles, dorsum, ventrum, brim, dorsal warts and gonads), and
 64 ossicles were rinsed five times with purified water. The ossicles were observed using a
 65 scanning electron microscope (TM 1000; Hitachi, Ltd., Tokyo, Japan).

66
 67 ***PCR amplification and phylogenetic analysis***

68 Total genomic DNA was extracted from 100 mg of muscle tissue using a DNeasy
 69 Blood & Tissue Kit (QIAGEN, Hilden, Germany) according to the manufacturer's
 70 instructions. Two partial mitochondrial genes, 16S rRNA and COI, were amplified
 71 using primers 16S-arL/brH and COI-ef/er (Miller et al. 2017). The PCR reactions were
 72 performed using a 50-μL reagent mix, containing 25 μL 2× Phanta Max Master Mix
 73 (Vazyme, Biotech Co., Ltd., Nanjing, China), 20-μL DNase free ddH₂O, 2-μL of each
 74 primer, and 1-μL template DNA, as suggested by the manufacturer. The PCR
 75 amplification procedure is shown in Table 1. PCR products were confirmed by 1.5%
 76 agarose gel electrophoresis, and purified using an OMEGA PCR kit (Omega, Biotek,
 77 Norcross). The purified PCR products were sequenced on an ABI 3730XL sequencer
 78 (Sangon, Biotech Co., Ltd., Shanghai). Sequence data were edited with Geneious
 79 R6.1.6 (Kearse et al. 2012) and deposited in GenBank (Table 2).

80
 81 **Table 1.** The PCR amplification procedures.

Primer	Sequence 5'→ 3'	PCR procedure
COI-ef	ATAATGATAGGAGGRTTTGG	Pre denaturation:95 °C for 3 min 40 cycles: Denaturation:95 °C for 40 s Annealing: 45 °C for 40 s Extension: 72 °C for 50 s
COI-er	GCTCGTGTRTCTACRTCCAT	
16S-arL	CGCCGTTTATCAAAAACAT	Pre denaturation:95 °C for 3 min 35 cycles: Denaturation:95 °C for 40 s Annealing: 50 °C for 40 s Extension: 68 °C for 50 s
16S-brH	CCGGTCTGAACTCAGATCACG	

82
 83 For a more comprehensive phylogenetic analysis, we not only used the sequences
 84 of Psychropotidae but also used the mitochondrial sequences of Elpidiidae and two
 85 species of Molpadiidae as the outgroup (Table 2). Twenty-five COI and 18 16S
 86 sequences were aligned using MAFFT 7 (Katoh and Standley 2013) using the E-INS-I
 87 strategy. The 16S and COI alignments were concatenated into the dataset (COI/16S =
 88 687/578 bp), while the alignment gaps and missing data were represented as '-' and '?'.
 89 Maximum likelihood (ML) and Bayesian inference (BI) analyses were performed using
 90 the concatenated dataset. JModelTest 2.1.10 (Darriba et al. 2012) was used to find the
 91 best-fit model from 88 competing models using Akaike information criterion (AIC)

92 calculations. In each case, GTR+I+G was the best-fit model for BI analyses. MrBayes
 93 3.2 (Huelsenbeck and Ronquist 2001) was used to conduct BI analyses. Markov Chain
 94 Monte Carlo (MCMC) iterations were run for 1, 000, 000 generations with sampling
 95 every 100 generations. The first 25% of trees were discarded as burn-in, and the
 96 consensus trees were summarized in 75% majority-rule trees. RAxML GUI 1.5
 97 (Silvestro and Michalak 2012; Stamatakis 2014) was used to perform ML analysis with
 98 the GTR+GAMMA+I substitution model for 1000 bootstraps, as recommended by
 99 Miller et al. (2017).

100

101 **Table 2.** The gene sequences of the specimens used in this study.

Family	species	GenBank accession number	
		16S	COI
Psychropotidae	<i>Benthodytes manusensis</i>	MH627223.1	MH627222.1
	<i>Benthodytes sanguinolenta</i>		HM196507.1
	<i>Benthodytes marianensis</i>	MH049433.1	MH049435.1
	<i>Benthodytes jiaolongi sp. nov.</i>	MW992746	MW990356
	<i>Benthodytes jiaolongi sp. nov.</i>	MW992747	MW990357
	<i>Psycheotrepes exigual</i>		KX874392.1
	<i>Psychropotes longicauda</i>	DQ777099.1	KU987469.1
	<i>Psychropotes moskalevi</i>	MN310400.1	MN313655.1
	<i>Psychropotes raripes</i>	MN310403.1	MN313656.1
	<i>Psychropotes verrucicaudatus</i>	MW992749	MW980089
	<i>Psychropotes verrucicaudatus</i>	MW992748	MW980088
	Elpidiidae	<i>Peniagone diaphana</i>	KX856725.1
<i>Peniagone incerta</i>			HM196402.1
<i>Peniagone sp. AKM-2016</i>		KX856726.1	KX874385.1
<i>Peniagone vignoni</i>			HM196381.1
<i>Elpidia glacialis</i>			HM196413.1
<i>Amperima robusta</i>		KX856728.1	KX874381.1
<i>Protelpidia murrayi</i>		KX856727.1	KX874382.1
<i>Scotoplanes sp. TT_2017</i>			LC230158.1
Laetmogone	<i>Laetmogone wyvillethomsoni</i>		HM196504.1
	<i>Pannychia moseleyi</i>	KX856731.1	KX874380.1
	<i>Benthogone abstrusa</i>	KX856733.1	KX874374.1
	<i>Enypniastes eximia</i>	KX856730.1	KX874383.1
Molpadiidae	<i>Molpadia arenicola</i>	KX856741.1	KX874344.1
	<i>Molpadia musculus AKM_2016</i>	KX856739.1	KX874386.1

102

103 **Results and Discussion**

104 *Morphological observations*

105 Order Elaspodida Théel, 1882

106 Suborder Psychropotina Hansen, 1975

107 Family Psychropotidae Théel, 1882

108 Genus *Benthodytes* Théel, 1882

109

110 **Diagnosis** (according to Hansen, 1975)

111 Auns dorsal. Unpaired dorsal appendages absent. Circum-oral (or post-oral) papillae
112 present. Tentacles soft, pliable, and retractile.

113

114 *Benthodytes jiaolongi* sp. nov.

115 (Figs 2-5)

116

117 **Material examined.** Catalog numbers: RSIO3710601, RSIO590504, RSIO590506 and
118 RSIO6017101, four adult specimens, collection numbers: DY37-JL106-B01 (156.947°
119 E, 13.017° N), DY59-ROV05-B05 (134.916° E, 16.916° N), DY59-ROV05-B05
120 (134.916° E, 16.933° N) and DY60-JL180-B04 (134.911° E, 16.935° N).

121

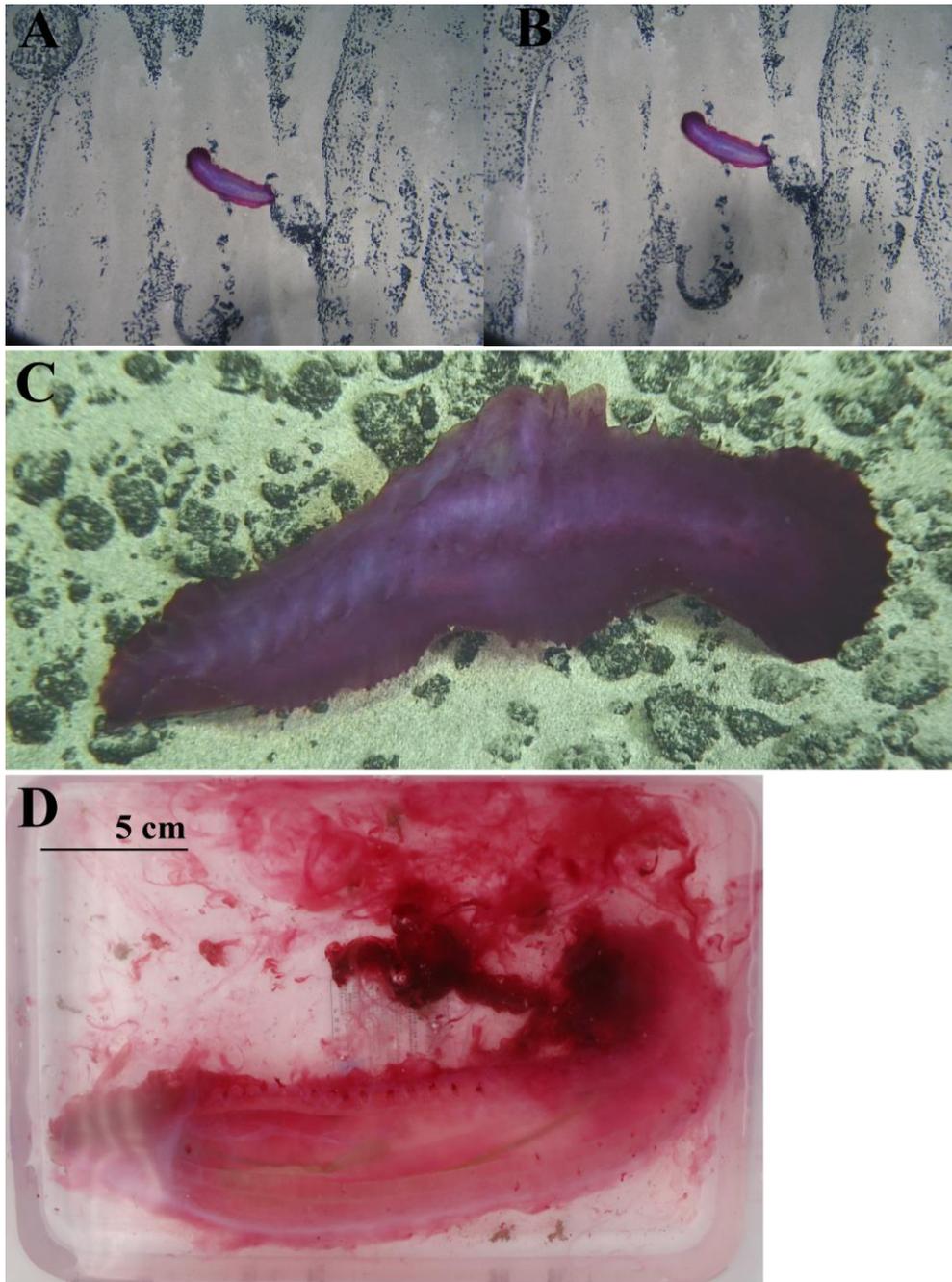
122 **Diagnosis**

123 Body elongated and subcylindrical when fixed. Skin red with violet, thin and soft.
124 No obvious large papillae arranged on the dorsal surface. Some minute papillae, conical
125 with tips, on the anterior dorsum. Brim narrow, thin, flattened. Mouth ventral, anus
126 terminal. Eighteen tentacles, retracted after fixing; circum oral papillae present. Dorsal
127 ossicles include rods and primary crosses with four arms. Rods present in tentacles.
128 Ossicles of ventrum not observed.

129

130 **Description of holotype (RSIO6017101)**

131 The length of the specimen was approximately 25 cm before being preserved in
132 10% seawater formalin. Color was violet in life (Fig. 2C); the skin was transparent, thin,
133 soft, and gelatinous after fixing. Brim retracted less than 0.7 cm in width. The dorsal
134 papillae did not develop well, but approximately nineteen pairs of minute papillae were
135 placed closely in two bands along the anterior dorsal radii. Another four single minute
136 papillae were located on the posterior dorsal edge. Midventral tube feet with
137 degeneration arranged in two rows, approximately 28 pairs. Mouth ventral, circum oral
138 papillae present, anus terminal. After the specimen arrived at the sea surface, autolysis
139 began; therefore, tentacles could not be identified (Fig. 2D). Few ossicles were
140 observed. Dorsal ossicles contained primary crosses with spiny arms and spinous rods
141 (Fig. 4A–F). The rods were approximately 400 µm long and the arms of the crosses
142 were approximately 200 µm in length. Rods were presented in tentacles and the length
143 was approximately 400–500 µm (Fig. 4G–J). Ossicles were not be found in the other
144 body parts.

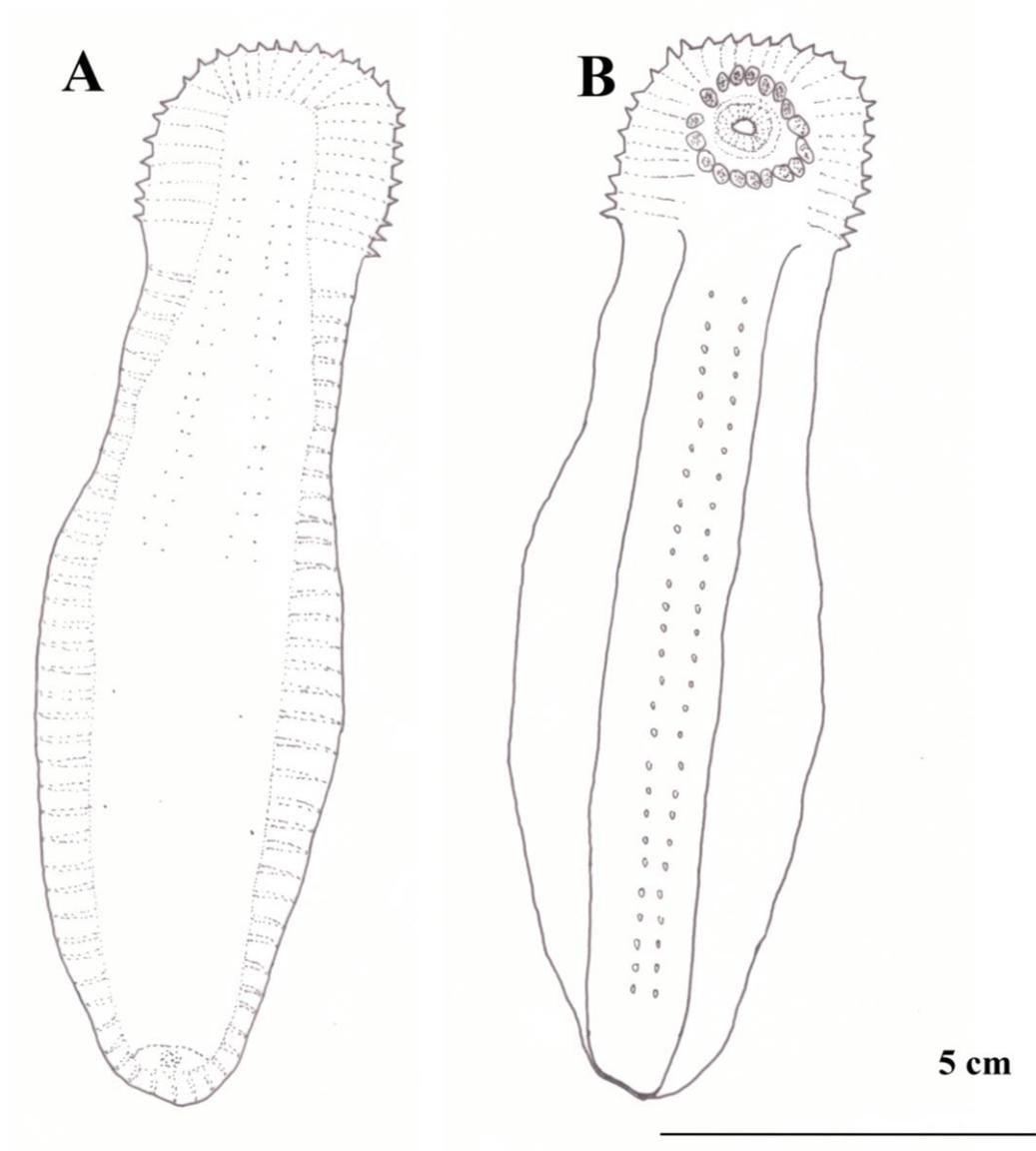


145

146 **Figure 2.** A-B: *Benthodytes jiaolongi* **sp.nov.** in situ on Weijia Guyot (RSIO3710601).

147 C: *Benthodytes jiaolongi* **sp.nov.** in situ on Kyushu-Palau Ridge (RSIO6017101). D:

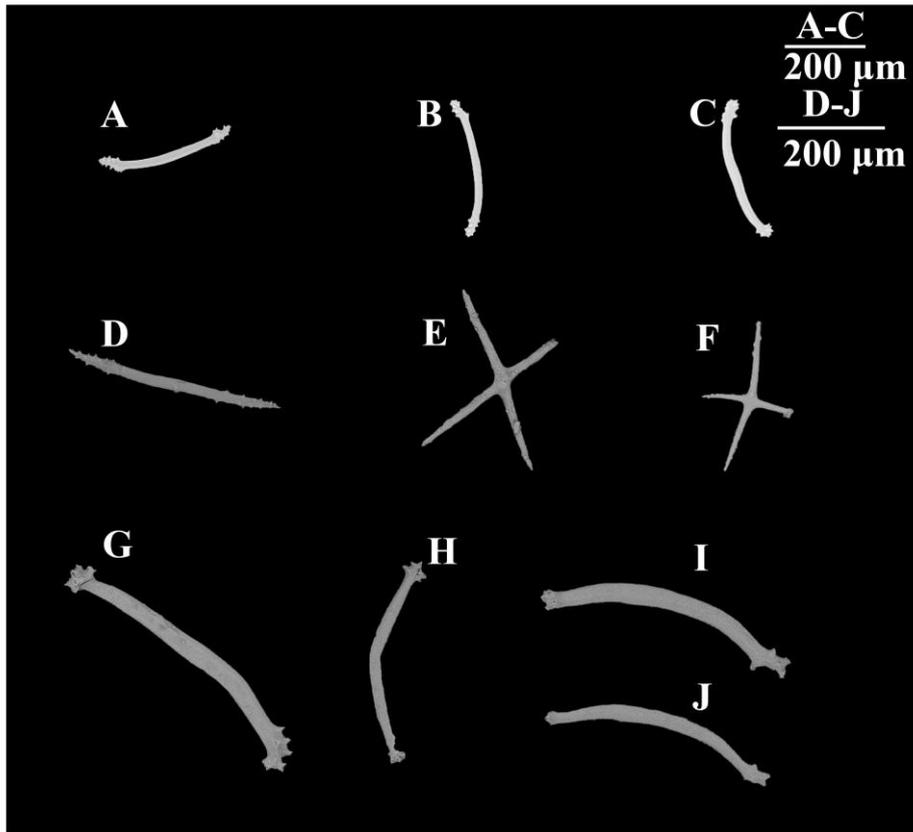
148 Specimen (RSIO6017101) before preservation in 10% seawater formalin.



149

150

Figure 3. *Benthodytes jiaolongi* sp.nov. A. Dorsal view. B: Ventral view.

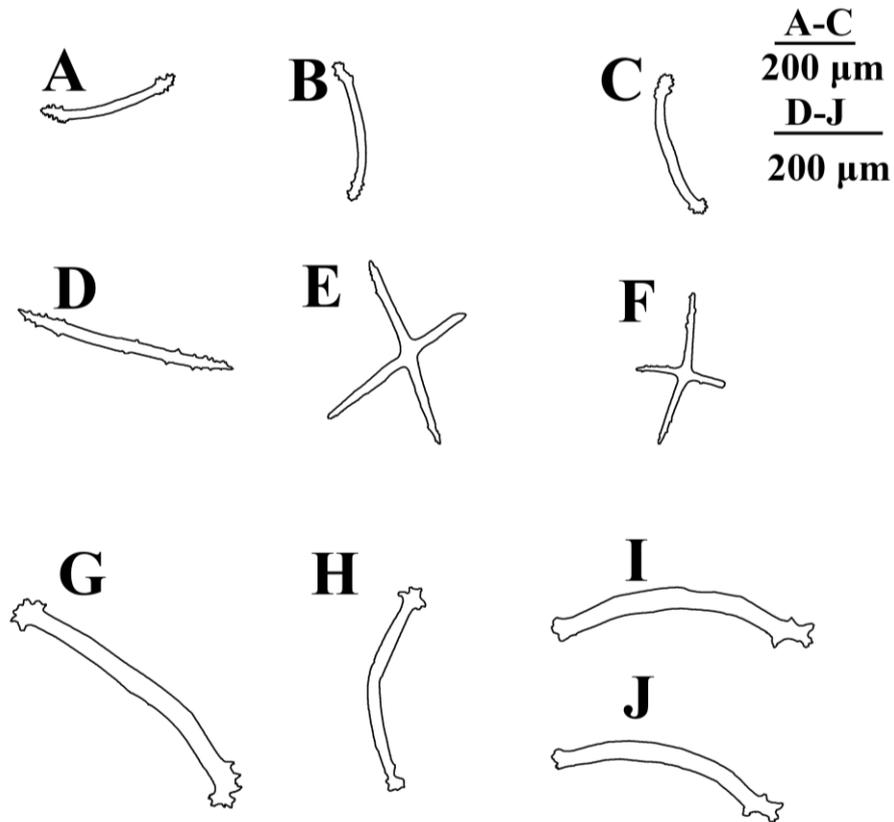


151

152 **Figure 4.** A-C: Scanning electron micrographs of dorsal body wall ossicles from

153 *Benthodytes. jiaolongi*, RSIO590506. D-F: Dorsal body wall ossicles from

154 *Benthodytes.jiaolongi*, RSIO6017101. G-J: Ossicles of tentacles.



155

156 **Figure 5.** *Benthodytes jiaolongi* **sp.nov.** ossicles from, A-F: dorsal body wall, G-J:
157 tentacles.

158

159 **Description of paratype.** RSIO3710601. The specimen was approximately 22 cm in
160 length and 5 cm wide at the maximum point. Color was red-violet at the seabed (Fig.
161 2A, B); after arriving at the sea surface, it became pale violet, and the skin appeared
162 transparent, with a white color after preservation in 10% seawater formalin for 5 years.
163 The normal paired dorsal papillae were absent, and the minute papillae could not be
164 distinguished. Owing to the long-term preservation, the quantity of the midventral tube
165 feet could not be determined, but the tube feet were arranged in two rows; the brim
166 could not be distinguished. Mouth ventral, with circum-oral papillae, anus terminal.
167 Eighteen tentacles, were retracted to the stalk. Ossicles were not observed in the
168 specimen.

169

170 RSIO590504. The specimen was approximately 22 cm in length before being
171 preserved in 10% seawater formalin. Color was red-violet on the deck and the skin
172 appeared transparent, with a white color after preservation in 10% seawater formalin.
173 During sampling, the specimen was accidentally stained with sponge, meaning that the
174 tentacles could not be determined and the dorsal tips could not be distinguished.
175 Midventral tube feet were identified but the quantity could not be determined. Mouth
176 ventral, anus terminal. Ossicles were not observed in the specimen.

176

177 RSIO590506. Specimen was approximately 13 cm in length before being preserved
in 99% alcohol and was heavily damaged. Color was red-violet after arriving at the sea

178 surface, and the skin appeared transparent. The specimen was stained with sponge as
179 was RSIO590504 and many external character could not be distinguished. Mouth
180 ventral, anus terminal. Few rods were observed on the dorsal body (Fig. 4A-C). Rods
181 were approximately 400 μm and the spine was terminal. Ossicles were not observed in
182 other the body wall.

183

184 **Etymology.** The name is derived from the first Chinese HOV 'Jiaolong'.

185

186 **Type locality.** Kyushu-Palau Ridge, tropical Western Pacific. Depth: 2453–2692 m,
187 muddy sediments, July 20, 2020 and January 21, 2021, Ruiyan Zhang, Bo Lu.

188

189 **Distribution.** Known from Weijia Guyot and Kyushu-Palau Ridge.

190

191 **Remarks.** According to the taxonomy of Hansen (1975), *Benthodytes jiaolongi* **sp. nov.**
192 was close to *Benthodytes sanguinolenta* Théel, 1882 and *Benthodytes typica* Théel,
193 1882.

194 *B. typica* was described by Théel in 1882 based on the Challenger Expedition, and
195 was the type species of *Benthodytes*. The original description indicated approximately
196 eight, minute, retractile processes located on each of the dorsal ambulacra and
197 unbranched spinose calcareous spicula scattered on the integument. Hansen (1975) re-
198 examined *B. typica* and reported that the specimens showed considerable variation. The
199 specimens from station 575 showed three or four pairs of minute dorsal papillae placed
200 along the anterior part of the radii, and rods were scattered in the ventrum, dorsum, and
201 tentacles. The specimens from the Bay of Bengal usually possessed three pairs of dorsal
202 papillae on the anterior body, but deposits were not found. Most specimens from the
203 Mozambique Channel did not show the the papillae and six or seven pairs of minute
204 dorsal papillae were presented in the specimens from the northern Indian Ocean.

205 *Benthodytes papillifera* Théel, 1882 was identified based on 13 specimens taken
206 from three Pacific Challenger stations. Théel described this species as being similar to
207 *B. sanguinolenta* based on the tentacles and pedicels. Some minute papillae, considered
208 as prominences of the integument, covered the dorsal surface and each papilla contained
209 one or more spicula. Hansen (1975) re-examined specimens from each of the stations
210 and proposed that the variation in *B. papillifera* represented the geographic variation of
211 *B. typica*. In the original description of *Benthodytes glutinosa*, Perrier (1896) indicated
212 that the differences from *B. typica* were the more elongated shape and the complete
213 absence of dorsal papillae. Hansen (1975) considered this species to be a synonym of
214 *B. typica*.

215 In general, the morphological features of *B. typical* can be summarized as: 3–7
216 pairs of minute papillae arranged on the dorsal surface, and rods scattered on the body
217 integument and tentacles. *B. jiaolongi* differed from *B. typica* in its arrangement and
218 number of dorsal papillae and by the ossicles. The dorsal minute papillae of *B. jiaolongi*
219 were arranged in two bands along the anterior dorsal ambulacra, and those of *B. typica*
220 were arranged in a row with 3–7 pairs of papillae. The ossicles of *B. jiaolongi* were
221 only presented in the tentacles and dorsum with rods and primary crosses, but *B. typica*

222 only possessed rods scattered on the ventrum, dorsum and tentacles.

223 The characteristics of *B. sanguinolenta* as described by Théel (1882) were many
 224 minute retractile processes scattered on the dorsal surface and dissolved calcareous
 225 deposits. According to a re-examination by Hansen (1975), the dorsal minute papillae
 226 were arranged in two radial bands, and the rods only presented on the midventral tube
 227 feet and tentacle discs within specimens from station 663. Rogacheva et al. (2009)
 228 recorded *B. sanguinolenta* and the main characteristics can be described as: minute
 229 dorsal papillae arranged in two bands or between the two bands; approximately 1–4
 230 papillae placed in a band, narrowing to one or two papillae at the posterior end; ossicles
 231 were not found. The differences in the characteristics between the new species *B.*
 232 *jiaolongi* and *B. sanguinolenta* can be listed as follows: (1) The dorsal papillae of *B.*
 233 *sanguinolenta* are arranged into two bands, whereas those of the new species were
 234 arranged in two rows on the anterior dorsal ambulacra. (2) The ossicles of the new
 235 species were only presented on the tentacles and dorsum with rods and primary crosses,
 236 whereas the ossicles were not found in *B. sanguinolenta*.

237

238 Genus *Psychropotes* Théel, 1882

239

240 *Psychropotes verrucicaudatus*

241 (Figs 6-8)

242

243 *Psychropotes verrucicaudatus* Xiao, Gong, Kou & Li, 2019:421-430.

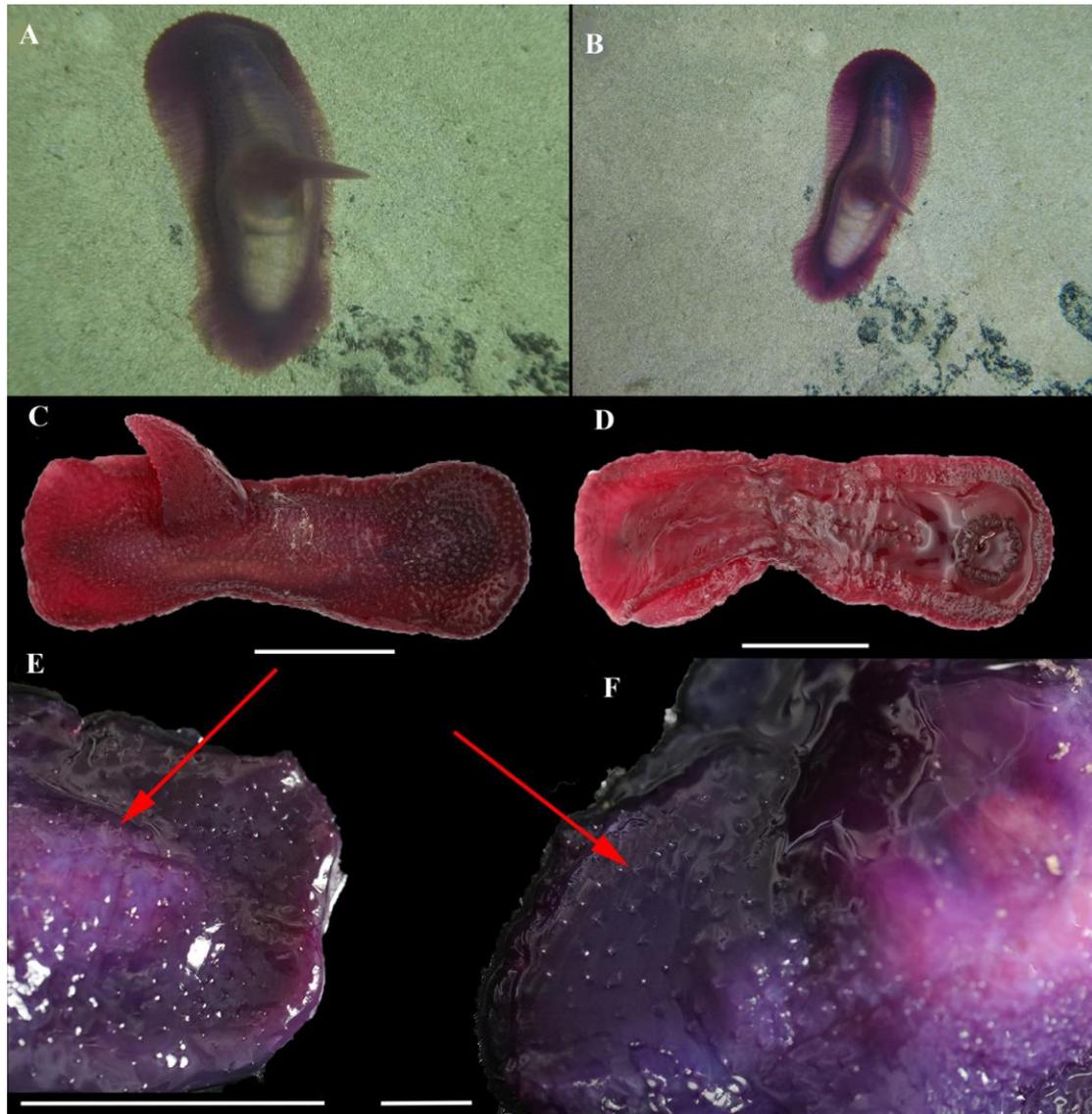
244

245 **Material examined.** Catalog numbers: RSIO6017006 and RSIO6018004, Two adult
 246 specimens, collection numbers: DY60-JL170-B05 (134.860° E, 12.079° N) and DY60-
 247 JL180-B04 (134.352° E, 13.569° N).

248

249 **Description**

250 RSIO6018004. The external morphology of the specimen resembled a barbell after
 251 collection and was approximately 20 cm in length before being preserved in 10%
 252 seawater formalin (Fig. 6C, D). Before preservation, the height of the appendage was
 253 approximately 50 mm, and the width was approximately 30 mm at the base (Fig. 6C,
 254 D). The dorsal skin was transparent with a brownish red color on the seabed and dark
 255 brown on the deck. Warts covered the dorsal skin and appendage, and the giant ossicles
 256 can be seen in the warts (Fig. 6E-F). Approximately 30 pairs of tube feet were arranged
 257 in two rows along the middle of the ventrum were degenerated. Sixteen tentacles were
 258 formed in a circle. The brim was broad and covered with warts on the dorsum.



259

260 **Figure 6.** A-B: species in situ. C-D: Specimen before preservation. Scale bar: 5 cm.

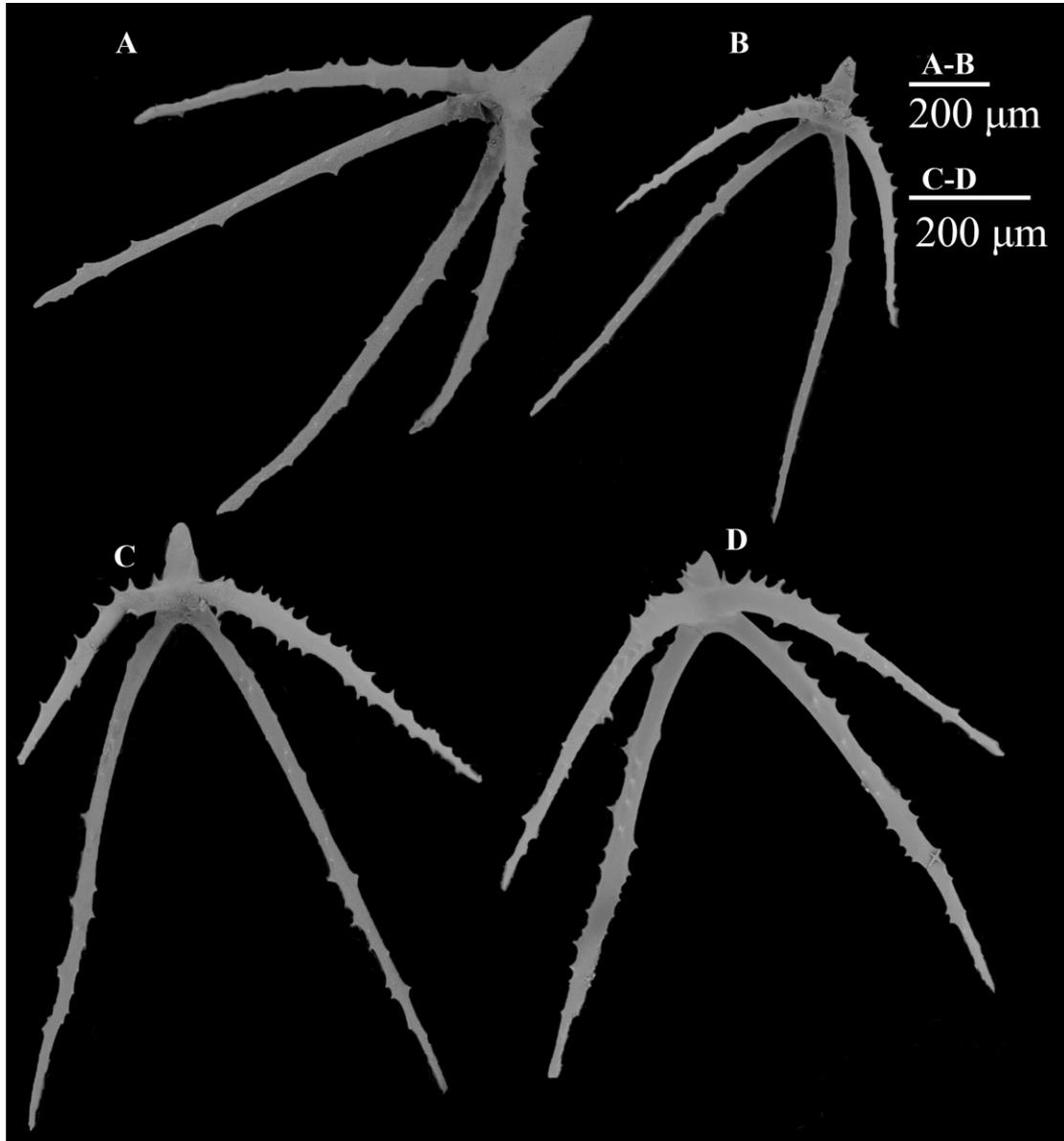
261 E-F: Red arrow points to the giant ossicles, specimen after preservation in 10%

262 seawater formalin. Scale bar: 5 cm (C, D, E) and Scale bar: 1 cm (F).

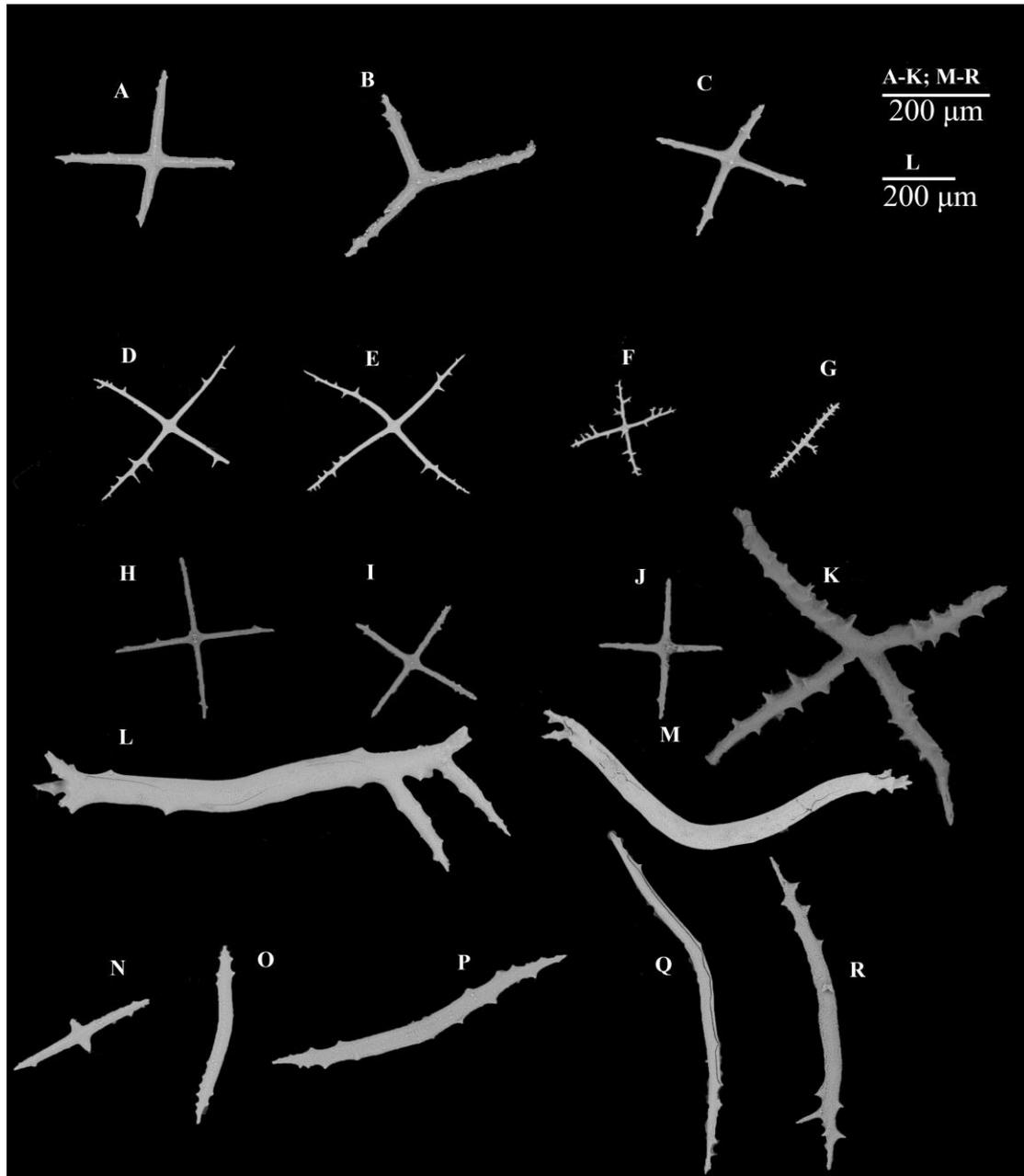
263

264 A giant cross with four arms could be seen in each wart. The arms were 800–1000
 265 μm in length, and the maximum width between the large arms was approximately 500
 266 μm . The arm flexion (presented by the length close to long arms / the length close to
 267 short arms at the intersection of the diagonal) was approximately 250 / 400 μm (Fig.
 268 7A–D). In addition, the height of the central rudimentary apophyses was approximately
 269 200–300 μm . Ventral ossicles can be divided into two types: primary cross with spiny
 270 arms and cross with three arms (Fig. 8A–C), and the length of each arm was
 271 approximately 200 μm . The primary crosses with spinous arms were present in the
 272 dorsum (Fig. 8D–F) and brim (Fig. 8H–J), and the length of the arms was up to 200 μm .
 273 Furthermore, dorsal ossicles possessed a spinous rod 170 μm in length (Fig. 8G), and
 274 large primary crosses with spiny arms were present in the brim (Fig. 8K). Ossicles of
 275 tentacles were rods with an irregular shape (Fig. 8L–R). The large rod with two

276 apophyses at the end was approximately 900–1000 μm in length; the small rod with
277 apophyses in the middle area was approximately 200 μm in length. Other rods with
278 spiny arms were 500–800 μm in length.
279



280
281 **Figure 7.** *Psychropotes verrucicaudatus*, A-D: The giant ossicles from the dorsal warts.



282

283

284 **Figure 8.** *Psychropotes verrucicaudatus* ossicles from, A-C: ventral body wall; D-G:

285 dorsal body wall; H-K: brim; L-R: tentacle.

286

287 RSIO6017005. The specimen was approximately 18 cm in length, the height of the

288 appendage was approximately 40 mm, and the width was approximately 20 mm at the

289 base. Mouth and anus ventral. The skin was transparent and of a light brown color.

290 Dorsal skin and appendage covered with warts, and the dorsum of the brim also grew

291 warts. Giant ossicles could be seen in the warts. Tentacles could not be clarified due to

292 damage, but there were more than 12. The ossicles were similar to those of

293 RSIO6018004.

294

295 **Type locality.** Jiaolong Seamount, South China Sea, western Pacific Ocean, sandy

bottom, depth 3615 m.

296

297 **Distribution.** Known from Jiaolong Seamount of South China Sea and Kyushu-Palau
298 Ridge.

299

300 **Variation.** The specimens were clearly a new record of *P. verrucicaudatus* but with
301 differences in external morphology and ossicles. *P. verrucicaudatus* was described by
302 Xiao et al. (2019). The differences between our specimens and *P. verrucicaudatus*
303 reflected intraspecific difference between growth stages. In addition, owing to the
304 preservation of the samples, Xiao et al. (2019) did not observe the complete ossicles of
305 the warts, but we were able to observe these structures in our specimens.

306

307 The intraspecific differences can be listed as follows: (1) in our specimens, the skin
308 was transparent and the color was darker than that of *P. verrucicaudatus*. (2) The width
309 of the appendage at the base was also larger than that of *P. verrucicaudatus*. (3) The
310 length of the primary crossing arms distributed in the dorsum, ventrum, and brim was
311 longer than that of *P. verrucicaudatus*. Furthermore, the spinous rod of the dorsal
312 ossicles was not present in *P. verrucicaudatus*, and the ventral body wall of the
313 specimens did not possess the tripartite ossicles. (4) Most of the ossicles of the tentacles
314 in our specimens were the same as those of *P. verrucicaudatus*, but longer.

315

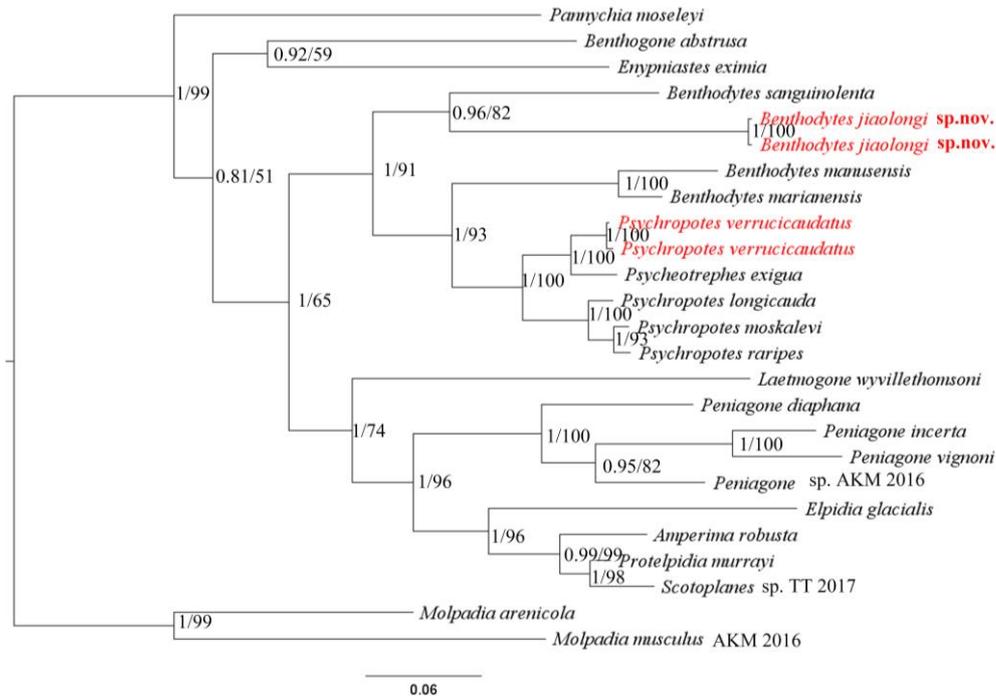
316 **Phylogenetic analyses**

317 Owing to limited genetic sequences, the phylogenetic relationships of Elasipodida
318 are not well studied. Miller et al. (2017) constructed high-level phylogenetic
319 relationships of Holothuroidea and separated Deimatidae from Elasipodida to rebuild
320 new phylogenetic relationships. The remaining families of Elasipodida included
321 Elpidiidae, Laetmogonidae, Pelagothuriidae, and Psychropotidae, but their positions
322 within Elasipodida remained unresolved. Li et al. (2018) used mitochondrial and
323 nuclear genes to perform phylogenetic analyses of Elasipodida, especially the
324 Psychropotidae, and the results showed that *Benthodytes* was a paraphyletic group of
325 Psychropotidae based on the clade of *B. sanguinolenta*.

326

327 To obtain clearer phylogenetic relationships, we concatenated 25 COI and 18 16S
328 sequences into a dataset to build the ML and BI trees. Although the genetic sequences
329 were limited, the topological structures of the ML and BI trees supported the
330 classification status of *B. jiaolongi* **sp. nov.** and *P. verrucicaudatus* (Fig. 9). The
331 phylogenetic relationships of Psychropotidae clustered into four parts and were
332 inconsistent with the traditional classification system based on Hansen (1975).
333 *Benthodytes* were paraphyletic in Psychropotidae and the new species was located in
334 the outmost clade of Psychropotidae. In addition, *Psychropotes* was a sister group to
335 *Psycheotrepes*, and the clade of *B. jiaolongi* **sp. nov.** and *B. sanguinolenta* were placed
336 in the lateral clade of the sister group. *P. verrucicaudatus* showed a paraphyletic
337 relationship with *Psychropotes* and was clustered in the clade of *Psycheotrepes exigua*.
338 Elpidiidae clustered into two clades. *Peniagone* clustered together into a supported
339 group, but *Peniagone diaphana* was a sister group to other *Peniagone* species. The

340 other four genera of Elpidiidae clustered into a group, and *Elpidia glacialis* was distant
 341 from the other three genera. *Protelpidia murrayi* and *Scotoplanes* sp. TT 2017 were
 342 closer to each other, and *Amperima robusta* was away from the two genera.
 343 Laetmogone was an obvious polyphyletic group, and *Pannychia moseleyi* was placed
 344 in the outmost clade of the other three families. *Laetmogone wyvillethomsoni* clustered
 345 with Elpidiidae and was situated outside; *Benthogone abstrusa* was clustered with
 346 *Enypniastes eximia*, but the possibility of a clade was low.



347

348 **Figure 9.** Bayesian inference (BI) and maximum likelihood (ML) trees based on
 349 concatenate sequences. The values are BI/ML at each node.

350

351 Based on the morphological and phylogenetic analyses, *B. jiaolongi* sp. nov. can
 352 be identified as a new species closely related to *B. sanguinolenta*. In addition, our
 353 specimens provided a new record of *P. verrucicaudatus* in the Western Pacific,
 354 broadening its distribution. Our results clarify that *Benthodytes* is paraphyletic and that
 355 the clade of *B. sanguinolenta* and *B. jiaolongi* sp. nov. is separated from the other
 356 species of *Benthodytes*.

357

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359

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