



# Revision of two species of Sinopotamon Bott, 1967 (Crustacea, Brachyura, Potamidae) endemic to China: a new combination and a new synonym

Ruxiao Wang<sup>1</sup>, Da Pan<sup>1</sup>, Hui Chen<sup>1</sup>, Binqing Zhu<sup>2,3</sup>, Hongying Sun<sup>1</sup>

I Jiangsu Key Laboratory for Biodiversity and Biotechnology, College of Life Sciences, Nanjing Normal University, 1 Wenyuan Rd, Nanjing 210023, China 2 Nanjing Institute of Environmental Sciences, Ministry of Ecology and Environment, Nanjing 210042, China 3 Research Station for Ecological Environment of Wuyi Mountains/Biodiversity Comprehensive Observation Station for Wuyi Mountains/State Environmental Protection Key Laboratory on Biosafety, Nanping, 353000, China

Corresponding authors: Hongying Sun (sunhongying@njnu.edu.cn), Da Pan (dapan@njnu.edu.cn)

Academic editor: Célio Magalhães | Received 13 April 2022 | Accepted 15 June 2022 | Published 14 July 2022

https://zoobank.org/2308C47E-483A-460A-B156-1A4D8DE50C28

**Citation:** Wang R, Pan D, Chen H, Zhu B, Sun H (2022) Revision of two species of *Sinopotamon* Bott, 1967 (Crustacea, Brachyura, Potamidae) endemic to China: a new combination and a new synonym. ZooKeys 1112: 179–197. https://doi.org/10.3897/zookeys.1112.85278

#### **Abstract**

The systematics of two problematic potamid species, *Sinopotamon koatenense* (Rathbun, 1904) and *Sinopotamon wuyiensis* Li, Lin, Cheng & Tang, 1985, both originally described from the Wuyi Mountains are resolved in this study. *Sinopotamon koatenense* is transferred to the genus *Huananpotamon* Dai & Ng, 1994, as the new combination *Huananpotamon koatenense* comb. nov. The new combination differs from its congeners in the form of the carapace, male pleon, male first gonopod, and vulvae. Phylogenetic analyses based on mitochondrial 16S rDNA sequences support the identification of *Huananpotamon koatenense* comb. nov. and a redescription is also provided. In addition, *S. wuyiensis* is confirmed as a junior synonym of *Sinopotamon fukienense* Dai & Chen, 1979 based on morphological similarities and phylogenetic lineages.

#### Keywords

16S rDNA, *Huananpotamon koatenense* comb. nov., *Sinopotamon wuyiensis*, systematics, taxonomy, Wuyishan National Park

## Introduction

Guadun, also spelled Koaten or Kuatun, is located in Tongmu Village in the Wuyi Mountains, which has been under administration of the Wuyishan National Park since 2021; it is a well-known type locality for several herpetofauna species, e.g., *Boulenophrys kuatunensis* (Pope, 1929), *Opisthotropis kuatunensis* Pope, 1928. As early as 1873, Father Armand David collected a female freshwater crab specimen from Guadun, and preserved it at the Muséum national d'Histoire naturelle, Paris (MNHN). Rathbun (1904) recognized the specimen as a new species, *Potamon (Potamon) koatenensis* Rathbun, 1904. With the establishment of the genus *Sinopotamon* Bott, 1967, *Potamon (P.) koatenensis* was then transferred to *Sinopotamon*, but had been regarded as a synonym of *Sinopotamon denticulatum* (A. Milne-Edwards, 1853) (Bott 1970). Although Ng et al. (2008) considered *Sinopotamon koatenense* as a valid species, Chu et al. (2018) synonymized *S. koatenense* with *Sinopotamon fukienense* Dai & Chen, 1979. To date, the taxonomic status and validity of *S. koatenense* has been debated for more than a century.

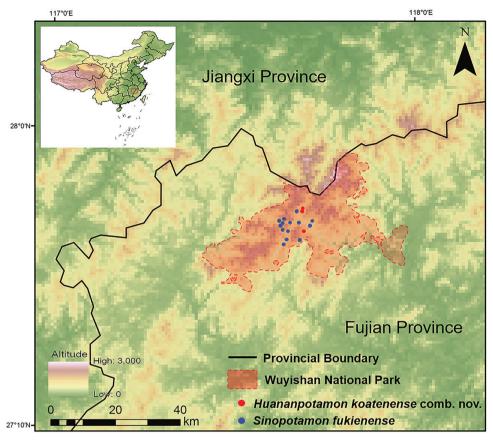
Sinopotamon wuyiensis Li, Lin, Cheng & Tang, 1985, was described from the Sangang, Tongmu Village in the Wuyi Mountains and has not been reported since its original description. Notably, except for Chu et al. (2018), neither Dai (1999) nor Ng et al. (2008) had listed the species in their species checklists. Consequently, the validity of *S. wuyiensis* has also been unconfirmed for nearly 40 years.

Recently, we conducted comprehensive surveys of freshwater crabs in the Wuyishan National Park and collected several specimens from the type localities of the abovementioned species (Fig. 1). Our morphological comparisons with the holotype of *S. koatenense* indicate that it was erroneously placed in *Sinopotamon*, and should be transferred to *Huananpotamon* Dai & Ng, 1994, as *Huananpotamon koatenense* comb. nov. Molecular phylogenetic analyses and detailed morphological comparisons with other *Huananpotamon* species indicate that the new combination as a species of *Huananpotamon*. The new combination is redescribed in the present study. In addition, *S. wuyiensis* is revised herein, suggesting that it is in fact a junior synonym of *S. fukienense*.

#### Material and methods

We conducted three comprehensive surveys of freshwater crabs at the Wuyishan National Park in March, November, and December of 2021. Specimens were collected by hand, preserved in 95% ethanol, and deposited in the freshwater crab collection of Jiangsu Key Laboratory for Biodiversity and Biotechnology, College of Life Sciences, Nanjing Normal University (NNU), Nanjing, China. Measurements, in millimeters, are of carapace width and length, respectively. The abbreviations used in the text and figures are as follows: G1, male first gonopod; G2, male second gonopod; and a.s.l., above sea level. The terminology used here follows that of Dai (1999) and Davie et al. (2015).

Shih et al. (2016) proposed that *Sinopotamon* Bott, 1967 be split into two genera, *Sinopotamon* s. str. and *Longpotamon* Shih, Huang & Ng, 2016. The authors, however, indicate that *Sinopotamon* s. str. and *Longpotamon* each comprise two lineages based on



**Figure 1.** Localities of the sampling sites for *Huananpotamon koatenense* and *Sinopotamon fukienense* in the Wuyishan National Park, Fujian.

extensive taxonomic sampling and comprehensive multi-locus phylogeny, and that another two new genera should be established. Therefore, until such taxonomic actions are made, we follow the classical taxonomic system proposed by Bott (1967) for *Sinopotamon*.

Total genomic DNA was extracted from gill tissue using the Trelief<sup>TM</sup> Animal Genomic DNA kit (Tsingke). 16S rDNA sequences were amplified using the primers 1471 and 1472 (Crandall and Fitzpatrick 1996). The protocol as follows: initial denaturation for 3 min at 95 °C, 35 cycles of denaturation for 15 s at 95 °C, annealing for 15 s at 48 °C, extension for 15 s at 72 °C, and final extension for 5 min at 72 °C. PCR products were sequenced on an ABI 3730 automatic sequencer. Newly generated sequences were deposited in GenBank (Table 1).

A total of 43 sequences from the ingroup and outgroup taxa were used in the phylogenetic analyses, including 33 downloaded sequences (Table 1). Pan et al. (2022) showed that the genus *Huananpotamon* nested within the "South China-adjacent Islands" clade; therefore, closely related genera from the same clade, in particular the taxa of closely related genera from Fujian, Guangdong, and Jiangxi, and some others from adjacent islands, were included for the present phylogenetic analyses (Table 1). For

**Table 1.** The 16S rDNA sequences used in the phylogenetic analysis.

Species	Locality	Accession No.	Reference
Ingroup taxa			
Apotamonautes hainanensis bawanglingensis	Haikou, Hainan	MN737137	Pan et al. 2022
Apotamonautes hainanensis hainanensis	Lean, Hainan	AB428459	Shih et al. 2009
Bottapotamon lingchuanense	Gongcheng, Guangxi	NC049011	Pan et al. 2022
Candidiopotamon okinawense	Ryukyus, Okinawa	AB208627	Shih et al. 2006
Chinapotamon depressum	Mashan, Guangxi	MZ350918	Pan et al. 2022
Geothelphusa dehaani	Chiba, Japan	AB290630	Yeo et al. 2007
Geothelphusa albogilva	Pingdong, Taiwan	MZ350921	Pan et al. 2022
Hainanpotamon daiae	Lingshu, Hainan	MZ350922	Pan et al. 2022
Hainanpotamon fuchengense	Haikou, Hainan	AB428461	Shih et al. 2009
Huananpotamon angulatum	Fuzhou, Fujian	AB428454	Shih et al. 2009
Huananpotamon chongrenense	Chongren, Jiangxi	ON454341	This study
Huananpotamon guixiense	Guixi, Jiangxi	ON505056	This study
Huananpotamon koatenense	Wuyishan, Fujian	ON505057	This study
Huananpotamon koatenense	Wuyishan, Fujian	ON505058	This study
Huananpotamon lichuanense	Lichuan, Jiangxi	MN737141	Pan et al. 2022
Huananpotamon lini	Songxi, Fujian	ON454342	This study
Huananpotamon medium	Nancheng, Jiangxi	ON454342	This study
Huananpotamon sheni	Shaowu, Fujian	ON505059	This study
Huananpotamon yiyangense	Yiyang, Jiangxi	ON505060	This study
Minpotamon sp.	Longhai, Fujian	ON454347	This study
Minpotamon kityang	Jieyang, Guangdong	MN253481	Mao et al. 2020
Minpotamon nasicum	Longhai, Fujian	AB428450	Shih et al. 2009
Nanhaipotamon hongkongense	Hong Kong, China	AB212869	Shih et al. 2005
Nanhaipotamon nanriense	Putian, Fujian	AB212868	Shih et al. 2005
Nanhaipotamon pinghense	Heping, Guangdong	AB433553	Shih et al. 2011
Nanhaipotamon pingyuanense	Pingyuan,Guangdong	AB265237	Shih et al. 2006
Nanhaipotamon wupingense	Wuping, Fujian	AB470496	Shih et al. 2011
Nanhaipotamon yongchuense	Yongchun, Fujian	AB433546	Shih et al. 2011
Neotiwaripotamon jianfengense	Changjiang, Hainan	MZ350933	Pan et al. 2022
Neotiwaripotamon whiteheadi	Qiongzhong, Hainan	MZ350934	Pan et al. 2022
Qianguimon elongatum	Guangxi, China	MG709240	Huang 2018
Sinolapotamon patellifer	Lingchuan, Guangxi	MZ350948	Pan et al. 2022
Outgroup taxa			
Acantiapotamon inflatum	Xuanen, Hubei	MZ350902	Pan et al. 2022
Aiyunamon lushuiense	Lushui, Yunnan	MZ350919	Pan et al. 2022
Indochinamon jianchuanense	Jianchuan, Yunnan	MZ350928	Pan et al. 2022
Latopotamon obtortum	Shuicheng, Guizhou	MZ350930	Pan et al. 2022
Parapotamonoides endymion	Kunming, Yunnan	MZ350935	Pan et al. 2022
Potamiscus yiwuensis	Yunnan, China	AB428476	Shih et al. 2009
Sinopotamon fukienense	Wuyishan, Fujian	ON505061	This study
Sinopotamon fukienense	Shaowu, Fujian	KT586120	Ji et al. 2016
Tenuilapotamon joshuiense	Leishan, Guizhou	MZ350951	Pan et al. 2022
Tenuilapotamon latilum latilum	Xianfeng, Hubei	MN737132	Pan et al. 2022
Vadosapotamon sheni	Rongjing, Sichuan	MZ350958	Pan et al. 2022

the genus *Huananpotamon*, eight species from the adjacent area of the Wuyishan National Park were included, of which four species are morphologically most similar to *H. koatenense* (Table 1; Fig. 9). Ten species belonging to the closely related "Central China" and "Indochina-Southwest China" clades (Pan et al. 2022) were used as outgroups. Among them, two specimens of *S. fukienense*, collected from Guadun (Koaten) and Shaowu of Fujian Province were also included in our phylogenetic analyses (Table 1).

All sequences were aligned using MAFFT 7.487 (Katoh and Standley 2013) based on the G-INS-I method. The best-fit model was selected using jModelTest 2.1.10 (Darriba et al. 2012). Maximum likelihood (ML) analysis was implemented in IQ-TREE 1.6.12 (Nguyen et al. 2015) using the ultrafast bootstrapping approach (Minh et al. 2013) with 1000 replicates. MrBayes 3.2.7a (Ronquist et al. 2012) was used to perform Bayesian inference (BI) analysis. Two independent runs were carried out. In each run, four MCMC chains were run for 30 million generations, with every 100 generations being sampled. Convergence of two runs was checked using Tracer 1.6 (Rambaut et al. 2014). All effective sampling sizes (ESS) values were more than 200. The first 25% of samples from the MCMC chain were discarded as burn-in. The pairwise estimates of Kimura 2-parameter (K2P) distances (Kimura 1980) were calculated using MEGA X (Kumar et al. 2018).

## **Results**

**Taxonomy** 

Family Potamidae Ortmann, 1896 Subfamily Potamiscinae Bott, 1970 Genus *Sinopotamon* Bott, 1967

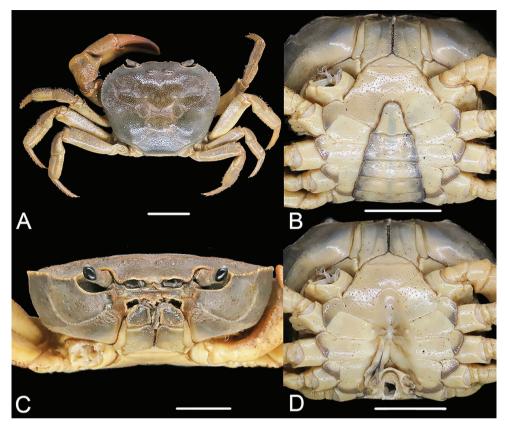
Sinopotamon fukienense Dai & Chen, 1979

Figs 2, 3

Sinopotamon fukienense Dai & Chen, 1979: 247, fig. 2(1)–(3); Dai 1999: 259, pl. 17 (7), fig. 136.

Sinopotamon wuyiensis Li, Lin, Cheng & Tang, 1985: 144, figs 1-6 [synonymy].

**Material examined.** 2 males, 23.2 × 29.0 mm, NNU11B-211121SF1; 22.0 × 27.6 mm, NNU 11B-211121SF2, Guadun, Wuyishan National Park, Fujian, coll. Caixin Liu, Kangqin Zhang, Ruxiao Wang & Hongying Sun, 21 November 2021. 3 female, 17.5 × 22.3 mm, NNU 11B-211121SF3; 19.5 × 25.4 mm, NNU 11B-211121SF4; 20.5 × 26.1 mm, NNU 11B-211121SF5, same data as above. 1 male, 22.5 × 28.1 mm, NNU 11B-211217SF6, Guadun, Wuyishan National Park, Fujian, coll. Hui Chen, Yunlong Sun, Ruxiao Wang & Hongying Sun, 17 December 2021. 2 males, 17.9 × 22.2 mm, NNU 11B-211217SF7; 17.6 × 22.3 mm, NNU 11B-211217SF8, Tongmuguan, Wuyishan National Park, Fujian, same data as above. 1 female, 16.8 × 21.0 mm, NNU 11B-211217SF9, same data as above. 4 females, 17.9 × 23.7 mm, NNU 11B-21121SF10, Dazhulan, Wuyishan National Park, Fujian, coll. Caixin Liu, Kangqin Zhang, Ruxiao Wang & Hongying Sun, 21 November 2021. 3 females, 22.3 × 28.0 mm, NNU 11B-211215SF11; 17.0 × 21.2 mm, NNU 11B-211215SF12; 26.3 × 33.4 mm, NNU 11B-211215SF13, same data as above, 15 December 2021. 2 males, 22.5 × 29.0 mm, NNU 11B-211219SF14; 21.3 × 27.2 mm, NNU 11B-211219SF15, Longdu, Wuyishan National Park, Fujian, coll. Hui Chen, Yunlong Sun,

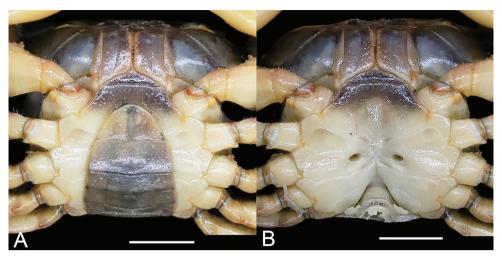


**Figure 2.** *Sinopotamon fukienense*, male,  $23.2 \times 29.0$  mm, NNU11B-211121SF1 **A** dorsal overall view ventral **B** view of anterior thoracic sternum and pleon **C** frontal view **D** ventral view of sterno-pleonal cavity with right G1 in situ (left G1 removed). Scale bars: 1.0 cm.

Ruxiao Wang & Hongying Sun, 19 December 2021. 1 female, 21.8 × 27.7 mm, NNU 11B-211219SF16, same data as above. 1 male, 19.6 × 24.9 mm, NNU 11B-211218SF17, Qiaoxia, Wuyishan National Park, Fujian, coll. Hui Chen, Yunlong Sun, Ruxiao Wang & Hongying Sun, 18 December 2021. 1 female, 17.9 × 22.8 mm, NNU 11B-211218SF18, same data as above. 1 female, 35.5 × 43.4 mm, NNU 11B-211218SF19, Jiangdun, Wuyishan National Park, Fujian, same data as above.

**Distribution and habitat.** Jiangxi Province, and Minjiang basin in Fujian Province. The crabs were found under rocks in mountain streams.

**Remarks.** Sinopotamon wuyiensis was considered a new species based on its distinct morphological characters in both male and female specimens, especially in the male specimen having a pair of vulvae on sternite 6, which is not seen in any known species of Sinopotamon. Unfortunately, the type material of S. wuyiensis, which was originally deposited in Fujian Research Institute of Parasite Disease, Fuzhou, Fujian Province, is now lost (Youzhu Cheng, pers. comm.). Therefore, to investigate the status and validity of S. wuyiensis, we conducted comprehensive collections in Tongmu Village but found no record of S. wuyiensis in the area. Instead, we collected 21 new specimens from 14



**Figure 3.** *Sinopotamon fukienense*, female, 22.5 × 29.0 mm, NNU 11B-211219SF14 **A** ventral view of pleon **B** ventral view showing female vulvae. Scale bars: 1.0 cm.

localities within the area (Fig. 1), all of which were identified as *S. fukienense* by morphological comparison (Figs 2, 3). According to the original description of *S. wuyiensis*, apart from the unique male sternite 6, *S. wuyiensis* is identical with *S. fukienense* in most characters, including outer margin of external orbital angle lined with 5–7 granules (Fig. 2A; cf. Dai and Chen 1979; Li et al. 1985; Dai 1999); anterolateral margin ~ 2× as long as external orbital outer margin (Fig. 2A; cf. Dai and Chen 1979; Li et al. 1985; Dai 1999); G1 terminal segment tapering, tip directed outwards (Fig. 2D; cf. Dai and Chen 1979: fig. 2(1), (2); Li et al. 1985: figs 1, 2; Dai 1999: fig. 136(3), (4)), not reaching suture sternites 5/6 (Fig. 2D; cf. Dai and Chen 1979: fig. 2(1), (2); Li et al. 1985: figs 1, 2; Dai 1999: fig. 136(3), (4)), subterminal segment to terminal segment ratio ~ 2.5 (Fig. 2D; cf. Dai and Chen 1979; Li et al. 1985; Dai 1999); female pleon subovate, vulvae ovate (Fig. 3; cf. Dai and Chen 1979; Li et al. 1985: figs 5, 6; Dai 1999: fig. 136(7), (8)). Therefore, it is most likely that the holotype of *S. wuyiensis* is a rare gynandromorphic specimen (Youzhu Cheng, pers. comm.), rather than a representative of a new species. Therefore, *S. wuyiensis* is herein synonymized with *S. fukienense*.

# Genus Huananpotamon Dai & Ng, 1994

*Huananpotamon koatenense* (Rathbun, 1904) comb. nov. Figs 4–6, 7A, 8A

Patamon (Potamon) koatenensis Rathbun, 1904: 308–309, pl. 13, fig. 3. Sinopotamon koatenense (Rathbun, 1904): Ng et al. 2008: 166.

**Type material.** *Potamon* (*P.*) *koatenensis*, *holotype*, 1 female, 18.4 × 23.4 mm, MNHN-IU-2014-23011 (citation: http://coldb.mnhn.fr/catalognumber/mnhn/

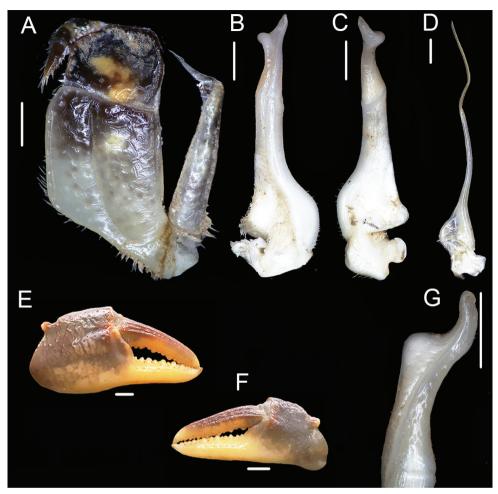
iu/2014-23011), 1200 m a.s.l., Koaten, Western Fujian Province, China, coll. Armand David, October 1873.

Other material. 1 male 15.8 × 19.2 mm, NNU 16C-211220HK1, 27°42.40'N, 117°41.47'E, 890 m a.s.l., Guwangkeng, Tongmu Village, Wuyishan National Park, Fujian Province, China, coll. Hui Chen, Ruxiao Wang, Yunlong Sun & Hongying Sun, 20 December 2021. 2 females, 19.7 × 25.4 mm, NNU 16C-211220HK2; 14.9 × 18.8 mm, NNU 16C-211220HK3, same collection data as above. 2 females, 17.0 × 21.6 mm, NNU 16C-211220HK4; 14.3 × 17.6 mm, NNU 16C-211220HK5, 27°45.77'N, 117°41.23'E, 854 m a.s.l., Qiaoxia, Tongmu Village, Wuyishan National Park, Fujian Province, China, coll. Hui Chen, Ruxiao Wang, Yunlong Sun & Hongying Sun, 17 December 2021. 1 female, 13.6 × 16.8 mm, NNU 16C-211220HK6, 27°46.15'N, 117°41.28'E, 893 m a.s.l., Jiangdun, Tongmu Village, Wuyishan National Park, Fujian Province, China, coll. Hui Chen, Ruxiao Wang, Yunlong Sun & Hongying Sun, 18 December 2021.

**Comparative material.** *Huananpotamon angulatum* (Dai, Chen, Song, Fan, Lin & Zeng, 1979), male, 14.3 × 18.9 mm, NNU 11B-21413HA1, Jianou, Fujian, 1 April 2021; *Huananpotamon lichuanense* Dai, Zhou & Peng, 1995, male, 12.0 × 14.4 mm, NNU 11B-21320HL1, Lichuan, Jiangxi, 20 March 2021; *Huananpotamon* 



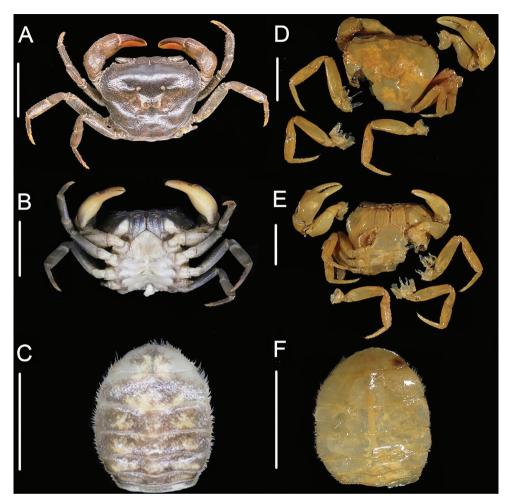
**Figure 4.** *Huananpotamon koatenense*, male, 15.8 × 19.2 mm NNU 16C-211220HK1 **A** dorsal view **B** ventral view **C** frontal view **D** ventral view showing sterno-pleonal cavity with right G1 in situ. Scale bars: 1.0 cm.



**Figure 5.** *Huananpotamon koatenense*, male, 15.8 × 19.2 mm, NNU 16C-211220HK1 **A** left third maxilliped **B** left G1 (dorsal view) **C** left G1 (ventral view) **D** left G2 **E** male right cheliped **F** male left cheliped **G** left G1 terminal segment (lateral view). Scale bars: 1.0 mm.

*lini* Cheng & Li, 2008, male, 16.5 × 18.8 mm, NNU 11B-211225HL1, Songxi, Fujian, 25 December 2021; *Huananpotamon yiyangense* Dai, Zhou & Peng, 1995, male, 13.7 × 16.3 mm, NNU 11B-21415HY1, Yiyang, Jiangxi, 15 April 2021.

**Diagnosis.** Carapace broader than long, dorsal surface slightly convex, finely pitted (Figs 4A, 6A, D); frontal margin distinctively bilobed, separated medially by shallow concavity (Figs 4A, 6A, D); epigastric cristae prominent, separated medially by distinct Y-shaped furrow (Figs 4A, 6A, D); postorbital cristae distinct, slightly rugose, confluent with epibranchial teeth (Figs 4A, 6A, D); anterolateral region convex, with rugose (Figs 4A, C, 6A, D). Third maxilliped exopod with flagellum (Figs 4A, 6A). G1 slender; tip part of terminal segment expanded, inner-distal angle prominent, outer-distal angle elongated, dagger-shaped (Figs 5B, G, 8A). G2 slender, terminal segment



**Figure 6.** Female specimens compared in this study: *Huananpotamon koatenense*, 19.7 × 25.4 mm, NNU 16C-211220HK2 **A** dorsal view **B** thoracic sternum and vulvae **C** pleon. *H. koatenense*, holotype, 18.4 × 23.4 mm, MNHN-IU-2014-23011 **D** dorsal view **E** thoracic sternum and vulvae **F** pleon. Scale bars: 1.0 cm.

long (Fig. 5D). Vulvae without operculum, not reaching suture of sternites 5/6 anteriorly, closely spaced from one another, opening inwards (Fig. 6B).

**Description.** Carapace broader than long, regions distinct, dorsal surface gently convex, finely pitted (Figs 4A, 6A, D). Frontal margin distinctively bilobed, divided into two broad lobes, separated by shallow concavity, margin of each lobe gently convex (Figs 4A, 6A, D). Epigastric cristae prominent, separated medially by distinct Y-shaped furrow extending to frontal region (Figs 4A, 6A, D); postorbital cristae distinct, slightly rugose, confluent with epibranchial teeth (Figs 4A, 6A, D). Cervical grooves distinct, deep (Figs 4A, 6A, D); H-shaped groove shallow but distinct (Figs 4A, 6A, D); anterolateral region convex with weak rugae; posterolateral surface smooth, with oblique striae; posterolateral margins converging posteriorly (Figs 4A, 6A, D). External orbital angle distinct, sharp, triangular, outer



**Figure 7.** *Huananpotamon koatenense* **A** color in life **B** natural habitat. Photographs by Hongying Sun, 20 December 2021, Tongmu Village, Wuyishan National Park, Fujian Province, China.

margin longer in length to inner margin, outer margin lined with small granules (Figs 4A, 6A, D). Epibranchial tooth granular, clearly demarcated from external orbital tooth by small gap (Figs 4A, 6A, D). Anterolateral margin convex, lined with 15–18 granules (Figs 4A, C, 6A, D). Orbits ovate, large; supraorbital, infraorbital margins cristate, lined with numerous inconspicuous granules (Fig. 4C). Epistome posterior margin with median lobe broadly triangular, lateral margins almost straight (Fig. 4C).

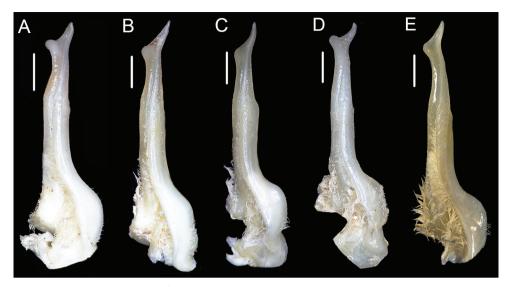
Ischium of third maxillipeds trapezoidal, length ~ 1.3× width, with distinct median oblique groove; merus subquadrate, length ~ 0.8× width (Figs 5A, 6B, E); exopod slender, reaching proximal one-third of merus length, with flagellum (Fig. 5A).

Chelipeds slightly asymmetrical (Figs 4A, 6A, D). Merus margins crenulated (Fig. 4A). Carpus surface wrinkled with sharp spine on inner margin, spinule at base (Fig. 4A). Major cheliped palm length ~ 1.4× height, surface rugose; fingers of major cheliped slightly curved, outer surface with rows of pits; dactylus 1.1× as long as palm length. Occlusal margin of both fingers lined with 11–15 irregular small teeth, forming small gape when fingers closed (Fig. 5E, F).

Ambulatory legs slender, surfaces and margins with scattered short setae (Figs 4A, 6A, D); second ambulatory leg longest, merus  $\sim 1.5 \times$  as long as dactylus (Figs 4A, 6A, D).

Thoracic sternum surface generally smooth, weakly pitted (Figs 4B, D, 6B, E); sternites 1, 2 fused, forming triangular structure, separated from sternite 3 by conspicuous suture; sternites 3, 4 completely fused without obvious median suture (Figs 4B, 6B, E). Male sterno-pleonal cavity reaching to imaginary line connecting median part of cheliped coxae (Fig. 4B, D). Male pleonal locking tubercle on median part of sternite 5 (Fig. 4D).

Male pleon triangular, surface smooth (Fig. 4B, D); somites 3–6 gradually decreasing in width; somite 6 width ~  $2.0\times$  length; telson broad triangular, width ~  $1.6\times$  length, apex rounded (Fig. 4B). Female pleon ovate; somite 6 width ~  $3.4\times$  length; telson semicircular, width ~  $2.9\times$  length (Fig. 6C, F).



**Figure 8.** The ventral view of G1 **A** *Huananpotamon koatenense*, 15.8 × 19.2 mm, NNU 16C-211220HK1 **B** *Huananpotamon angulatum*, male, 14.3 × 18.9 mm, NNU 11B-21413HA1 **C** *Huananpotamon lichuanense*, male, 12.0 × 14.4 mm, NNU 11B-21320HL1 **D** *Huananpotamon lini*, male, 16.5 × 18.8 mm, NNU 11B-211225HL1 **E** *Huananpotamon yiyangense*, male, 13.7 × 16.3 mm, NNU 11B-21415HY1. Scale bars: 1.0 mm.

G1 slender, bent dorsal-ward at proximal one-thirds (Fig. 5B, C); tip well exceeding pleonal locking tubercle in situ, reaching suture between thoracic sternites 4, 5 (Fig. 4D); subterminal segment length ~ 2.1× length of terminal segment; distal part of subterminal segment almost straight; terminal segment slender, inner margin slightly concave, outer margin straight, distally expanded, inner-distal angle high, rounded, outer-distal angle elongated, dagger-shaped (Fig. 5B, C, G); groove for G2 running mid-line of ventral surface (Fig. 5B). G2 slender, longer than G1, subterminal segment length ~ 1.7× length of terminal segment (Fig. 5D). Female vulvae on thoracic sternite 6, elongate-ovate, without distinct operculum, closely spaced from one another, opening inwards (Fig. 6B).

**Live coloration.** Generally dark brown to purplish brown. The immovable finger of the chelipeds and surrounding areas are pale yellow (Fig. 7A).

**Distribution and habitat.** Tongmu Village, Wuyishan National Park, Fujian Province, China. This species lives in moist mud burrows under rocks near the small hillstream, ca 900 m a.s.l. (Fig. 7B). *Bottapotamon engelhardti* (Bott, 1967) was found to be sympatric with *H. koatenense* and appears to occupy the shallower parts of the streams and hides under rocks.

**Remarks.** Thanks to digitalization of the specimen, we are able to examine the photographs of the holotype of *S. koatenense* provided by MNHN (http://coldb.mnhn. fr/catalognumber/mnhn/iu/2014-23011). *Potamon* (*P.) koatenensis* can be distinguished from *Sinopotamon* by several characters: small sized (versus large sized in *Sinopotamon*; cf. Bott 1967; Dai 1999), external orbital angle acutely triangular (Fig. 6D) (versus external orbital angle broadly triangular in *Sinopotamon*; cf. Bott 1967; Dai 1999), maxilliped

3 ischium relatively broad (Fig. 6E) (versus maxilliped 3 ischium relatively narrow in Sinopotamon; cf. Bott 1967; Dai 1999; fig. 139(1)), ambulatory legs slender (Fig. 6D, E) (versus ambulatory legs stout in *Sinopotamon*; cf. Bott 1967; Dai 1999), male sternopleonal cavity relatively narrow (Fig. 6E) (versus male sterno-pleonal cavity relatively broad in Sinopotamon; cf. Bott 1967; Dai 1999), female pleon broadly ovate (Fig. 6F) (versus female pleon ovate in *Sinopotamon*; cf. Bott 1967; Dai 1999), vulvae relatively small and narrow (Fig. 6E) (versus vulvae relatively large and wide in Sinopotamon; cf. Bott 1967; Dai 1999). In contrast, the holotype of S. koatenense conforms well to the genus diagnosis for Huananpotamon: small sized (cf. Dai and Ng 1994; Dai 1999), postorbital region narrow (Fig. 6A, D; cf. Dai and Ng 1994; Dai 1999: pl. 7, fig. 7), external orbital angle acutely triangular (Fig. 6A, D; cf. Dai and Ng 1994; Dai 1999: pl. 7, fig. 7), maxilliped 3 ischium relatively broad (Fig. 6A, D; cf. Dai and Ng 1994: fig. 1; Dai 1999: fig. 66(1)), ambulatory legs slender (Fig. 6A, B, D, E; cf. Dai and Ng 1994; Dai 1999: pl. 7, fig. 7), male sterno-pleonal cavity relatively narrow (Fig. 6B, E; cf. Dai and Ng 1994: fig. 1; Dai 1999: fig. 66(8)), female pleon broadly ovate (Fig. 6C, F; cf. Dai and Ng 1994: fig. 1; Dai 1999: fig. 66(7)), vulvae small, ovate, not reaching the sutures of sternites 5/6 (Fig. 6B, E; cf. Dai and Ng 1994: fig. 1; Dai 1999: fig. 66(8)).

Huananpotamon koatenense is close to four other Huananpotamon species, H. angulatum (Dai, Chen, Song, Fan, Lin & Zeng, 1979), H. lichuanense Dai, Zhou & Peng, 1995, H. lini Cheng & Li, and H. yiyangense Dai, Zhou & Peng, 1995 in having slender G1 terminal segment, and elongated outer-distal angle of G1 terminal segment (Fig. 8). However, the G1 terminal segment of H. koatenense can be distinguished from its congeners by the following characters: inner margin of G1 terminal segment slightly concave (Fig. 8A) (versus inner margin of terminal segment relatively straight in H. lichuanense and H. lini; cf. Dai et al. 1995: fig. 3; Cheng et al. 2008: fig. 1; Fig. 8C, D), G1 terminal segment with inner-distal angle relatively higher, globular in shape (Fig. 8A) (versus terminal segment with inner-distal angle relatively lower, somewhat blunt in H. angulatum, H. lichuanense, and H. yiyangense; cf. Dai and Ng 1994: fig. 1; Dai et al. 1995: figs 1, 3; Fig. 8B-C, E), G1 terminal segment with outer-distal angle relatively short (Fig. 8A) (versus terminal segment outer-distal angle relatively elongated in H. angulatum, H. lichuanense, and H. yiyangense; cf. Dai and Ng 1994: fig. 1; Dai et al. 1995: figs 1, 3; Fig. 8B-C, E). Other differences existing in the external orbital teeth, epibranchial teeth, and vulva are listed in Table 2. In this study, only one male specimen was collected, so we did not have enough specimens to analyze intraspecific variations and this matter may require further investigation.

**Table 2.** Morphological differences among *Huananpotamon koatenense*, and the other four species of *Huananpotamon*, *H. chongrenense*, *H. lini*, *H. medium*, *H. obtusum*, and *H. ruijinense*.

Character	External orbital	Epibranchial	Cleft between external orbital	Vulva	
	tooth	tooth	and epibranchial teeth		
H. koatenense	Sharp (Fig. 4A)	Granular (Fig. 4A)	Shallow (Fig. 4A)	Ovate (Fig. 6B)	
H. angulatum (cf. Dai & Ng, 1994: fig. 1)	Sharp	Granular	Shallow	Ovate	
H. lichuanense (cf. Dai et al., 1995: fig. 3)	Acutely sharp	Granular	Shallow	Widely ovate	
H. lini (cf. Cheng et al., 2008: fig. 1)	Acutely sharp	Rounded	Deep, U-shaped	Widely ovate	
H. yiyangense (cf. Dai et al., 1995: fig. 1)	Acutely sharp	Rounded	Shallow	Ovate	

## Phylogenetic analyses

A total of 41 species from 21 genera were included in the present phylogenetic analyses. The phylogenetic trees were reconstructed using BI and ML and yielded similar phylogenetic topologies with some minor differences in the terminal lineages (Fig. 9). The phylogenetic relationship showed that the genus *Huananpotamon* is monophyletic and belongs to the "South China-adjacent Islands" clade (Pan et al. 2022) (Fig. 9). Huananpotamon koatenense is genetically closest to H. lichuanense in the phylogenetic tree. Except for *H. angulatum*, the interspecific K2P genetic distances between H. koatenense and its congeners range from 1.77% to 2.51% (Table 3). Although the K2P genetic distance between the new combination and H. guixiense is 1.77%, it is still larger than other interspecific distances of potamid crabs (e.g., 1.47% between Nanhaipotamon formosanum (Parisi, 1916) and Nanhaipotamon dongyinense Shih, Chen & Wang, 2005; 0.93% between Nanhaipotamon nanriense Dai, 1997 and Nanhaipotamon yongchuense Dai, 1997; cf. Shih et al. 2011; 1.47% between Minpotamon nasicum (Dai, Chen, Song, Fan, Lin & Zeng, 1979) and Minpotamon kityang Mao & Huang, 2020; cf. Mao and Huang 2020), and the two species occur far apart (~ 90 km) and can be easily distinguished by the morphological characters in G1 and external orbital teeth. Additionally, we note the smallest genetic distance of 0.27% between *H. medium* and *H. yiyangense*. It is nevertheless comparable with some other pairs of potamid species (e.g., 0.19% between Aparapotamon grahami (Rathbun, 1931) and Aparapotamon binchuanense Tan, Zhou & Zou, 2021; cf. Tan et al. 2021; 0.38% between Aparapotamon similium Dai & Chen, 1985 and Aparapotamon binchuanense Tan, Zhou & Zou, 2021; cf. Tan et al. 2021). Additionally, they can be distinguished from each other by their morphologies (Dai et al. 1995; Cheng et al. 2008). It is noteworthy that the K2P genetic distances between H. angulatum and other Huananpotamon species are relatively large, ranging from 3.80% to 6.38% (Table 3). However, H. angulatum is not morphologically or geographically distinct. Further phylogenetic study of *Huananpotamon* with more extensive sampling of geographical populations, species, and genetic markers is needed. The specimen of S. fukienense from the Tongmu Village, the type locality for S. wuyiensis, formed a monophyletic group with the specimen from Shaowu City, the type locality for S. fukienense (Fig. 9), without distinct genetic difference.

### Discussion

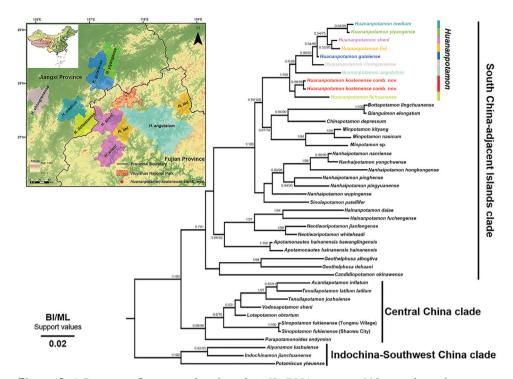
The type specimen of *Potamon (Potamon) koatenensis* is a female individual whose original type locality is recorded as Koaten (Guadun), Fujian Province, China, inhabiting an altitude of ca 1200 m (Rathbun 1904). The original description of the species is brief and minimally illustrated, and the significance of G1 in freshwater crab classification was not appreciated at that time, making correct judgments based solely on the original description difficult. Follow-up field surveys of the type locality of this species

**Table 3.** Genetic distances of K2P pairwise genetic distances based on 16S rDNA sequences among *Huananpotamon koatenense*, other *Huananpotamon* species, and five taxa from allied genera.

	1	2	3	4	5	6	7	8	9	10	11	12	13	14
1. Huananpotamon angulatum														
2. Huananpotamon chongrenense	3.80%													
3. Huananpotamon guixiense	3.80%	0.92%												
4. Huananpotamon koatenense	4.16%	2.23%	1.77%											
5. Huananpotamon lichuanense	5.14%	3.51%	3.09%	2.24%										
6. Huananpotamon medium	5.31%	2.40%	2.19%	2.02%	5.08%									
7. Huananpotamon sheni	4.75%	1.34%	0.44%	2.27%	2.75%	0.89%								
8. Huananpotamon yiyangense	6.38%	2.56%	1.67%	1.78%	3.18%	0.27%	1.58%							
9. Huanpotamon lini	4.49%	1.57%	1.66%	2.51%	3.00%	1.12%	1.66%	0.88%						
10. Minpotamon nasicum	9.05%	7.27%	7.03%	5.97%	8.02%	8.84%	7.21%	9.68%	7.21%					
11. Nanhaipotamon pingyuanense	8.36%	5.63%	6.12%	5.78%	6.86%	8.92%	7.04%	8.26%	7.04%	9.21%				
12. Chinapotamon depressum	8.26%	6.18%	6.44%	6.28%	7.84%	9.14%	7.02%	9.42%	7.62%	7.52%	7.08%			
13. Qianguimon elongatum	9.53%	8.87%	8.62%	7.07%	9.97%	11.28%	8.91%	12.06%	9.22%	9.19%	10.08%	7.40%		
14. Sinolapotamon patellifer	7.49%	6.84%	6.62%	6.28%	7.31%	9.05%	7.23%	9.31%	7.23%	7.51%	6.59%	5.89%	8.37%	

was also lacking. Therefore, it is no wonder that Bott (1970), Ng et al. (2008), and Chu et al. (2018) all followed the classification of *Potamon* (*P.*) *koatenensis* as a member of *Sinopotamon*. Therefore, to confidently resolve the long-standing controversy about the taxonomic status and validity of this species, we conducted comprehensive field surveys and collected specimens of *H. koatenense* in the surrounding areas of Guadun (Koaten) (Fig. 1). As informed by senior villagers, Father Armand David bought specimens from them in a church located in Guadun (Koaten). We therefore speculate that the holotype of *H. koatenense* was probably not actually collected in Guadun. Villagers may have collected the female specimen somewhere nearby, and taken it to the church. Based on the literature and our surveys, a total of eight *Huananpotamon* species are peripatrically distributed in the adjacent regions of the Wuyishan National Park (Fig. 9). To the best of our knowledge, *H. koatenense* is the only *Huananpotamon* species distributed inside and endemic to the Wuyishan National Park.

Previously, there were three freshwater crab species, *S. koatenense*, *S. fukienense*, and *S. wuyiensis* recorded in the area of the park. Based on our comprehensive surveys and research of the crabs in this area, there are still three species, namely, *H. koatenense*, *S. fukienense*, and *B. engelhardti*. The present study sheds some light on the biodiversity of freshwater crabs in the Wuyishan National Park, Fujian Province, China.



**Figure 9.** A Bayesian inference tree based on the 16S rDNA sequence. Values at the nodes represent bootstrap (BS) and posterior probability (BPP) values for ML and BI, respectively. Only values higher than 70/0.5 (BS/BPP) are shown. Distribution areas of eight *Huananpotamon* species are shown in the upper left map using the tree colors.

# **Acknowledgements**

We are grateful to Peter K. L. Ng (National University of Singapore) for his fruitful discussions and encouragement. We also thank Yunlong Sun (College of Life Sciences, NNU) for assistance with sampling; Kangqin Zhang and Luyao Wang (College of Life Sciences, NNU) for help with lab work. Finally, we give a special thanks to the Subject editor and Chao Huang, Jongkar Grinang for greatly improving our manuscript. This project was supported by the National Natural Science Foundation of China (No. 32170454 to SHY), and the Project of Biological Resources Survey in Wuyishan National Park.

## References

Bott R (1967) Potamiden aus Ost-Asien (*Parapotamon* De Man, *Sinopotamon* n. gen., *Candidiopotamon* n. gen., *Geothelphusa* Stimpson). Senckenbergiana biologica 48: 203–220.

- Bott R (1970) Die Süßwasserkrabben von Europa, Asien, Australien und ihre Stammesgeschichte. Eine Revision der Potamoidea und Parathelphusoidea (Crustacea, Decapoda). Abhandlungen der Senckenbergischen Naturforschenden Gesellschaft 526: 1–338.
- Cheng YZ, Li LS, Lin CX, Li YS, Fang YY, Jiang DW, Huang CY, Zhou AP, Zhang X (2008) Two new species of *Huananpotamon* as the second intermediate host of *Paragonimus skriabini* (Decapoda:Potamidae). Chinese Journal of Zoonoses 24: 885–889. [in Chinese with English abstract]
- Chu KL, Ma XP, Zhang ZW, Wang PF, Lv LN, Zhao Q, Sun HY (2018) A checklist for the classification and distribution of China's freshwater crabs. Biodiversity Science 26: 274–282. [in Chinese with English summary] https://doi.org/10.17520/biods.2018062
- Crandall KA, Fitzpatrick Jr JF (1996) Crayfish molecular systematics: Using a combination of procedures to estimate phylogeny. Systematic Biology 45(1): 1–26. https://doi.org/10.1093/sysbio/45.1.1
- Dai AY (1997) A revision of freshwater crabs of the genus *Nanhaipotamon* Bott, 1968 from China (Crustacea: Decapoda: Brachyura: Potamidae). The Raffles Bulletin of Zoology 45: 209–235.
- Dai AY (1999) Fauna Sinica: Arthropoda Crustacea Malacostraca Decapoda Parathelphusidae Potamidae. Science Press, Beijing, China, 501 pp. [In Chinese with English summary]
- Dai AY, Chen GX (1979) On the freshwater crabs from Fujian Province. Acta Zoologica Sinica 25: 243–249. [in Chinese with English abstract]
- Dai AY, Chen GX (1985) A preliminary report on the freshwater crabs of Hengduan Mountains Area. Sinozoologia 3: 39–72.
- Dai AY, Ng PKL (1994) Establishment of a new genus of freshwater crab, *Huananpotamon* (Crustacea: Decapoda: Brachyura: Potamidae) from southern China. The Raffles Bulletin of Zoology 42: 657–661.
- Dai AY, Chen GX, Song YZ, Fan PF, Lin YG, Zeng YQ (1979) On new species of freshwater crabs harbouring metacercariae of lung flukes. Acta Zootaxonomica Sinica 4: 122–131. [in Chinese with English abstract]
- Dai AY, Zhou XM, Dong WP (1995) On seven new species of freshwater crab of the genus *Huananpotamon* Dai & Ng, 1994 (Crustacea: Decapoda: Brachyura: Potamidae) from Jiangxi Province, Southern China. The Raffles Bulletin of Zoology 43: 417–433.
- Darriba D, Taboada GL, Doallo R, Posada D(2012) jModelTest 2: More models, new heuristics and parallel computing. Nature Methods 9(8): e772. https://doi.org/10.1038/nmeth.2109
- Davie PJ, Guinot D, Ng PKL (2015) Systematics and classification of Brachyura. In: Castro P, Davie PJF, Guinot D, Schram FR, von Vaupel Klein JC (Eds) Treatise on zoology anatomy, taxonomy, biology. The Crustacea 9C–I. Brill, Leiden, 1049–1130. https://doi.org/10.1163/9789004190832\_021
- Huang C (2018) Revision of *Yarepotamon* Dai & Türkay, 1997 (Brachyura: Potamidae), freshwater crabs endemic to southern China, with descriptions of two new genera and four new species. Journal of Crustacean Biology 38(2): 173–189. https://doi.org/10.1093/jcbiol/rux120

- Katoh K, Standley DM (2013) MAFFT multiple sequence alignment software version 7: Improvements in performance and usability. Molecular Biology and Evolution 30(4): 772–780. https://doi.org/10.1093/molbev/mst010
- Kimura M (1980) A simple method for estimating evolutionary rate of base substitutions through comparative studies of nucleotide sequences. Journal of Molecular Evolution 16(2): 111–120. https://doi.org/10.1007/BF01731581
- Kumar S, Stecher G, Li M, Knyaz C, Tamura K (2018) MEGA X: Molecular evolutionary genetics analysis across computing platforms. Molecular Biology and Evolution 35(6): 1547–1549. https://doi.org/10.1093/molbev/msy096
- Li YS, Lin JX, Cheng YZ, Tang JH (1985) A new species of *Sinopotamon* crab harbouring metacercariae of lung flukes from the Wuyishan National Reserve, Fujian (Decapoda: Sinopotamonidae). Wuyi Science Journal 5: 141–146. [in Chinese with English abstract]
- Mao SY, Huang C (2020) Descriptions of a new species of *Minpotamon* Dai & Türkay, 1997, and a monotypic new genus of aquatic freshwater crab (Brachyura, Potamidae) from eastern Guangdong, China. Crustaceana 93(11–12): 1295–1313. https://doi.org/10.1163/15685403-bja10060
- Milne-Edwards A (1853) Mémoire sur la Familie des Oeypodiens. Annales des sciences naturelles. Zoologie et biologie animale 3: 163–228.
- Minh BQ, Nguyen MAT, Von Haeseler A (2013) Ultrafast approximation for phylogenetic bootstrap. Molecular Biology and Evolution 30(5): 1188–1195. https://doi.org/10.1093/molbev/mst024
- Ng PKL, Guinot D, Davie PJ (2008) Systema Brachyurorum: Part I. An annotated checklist of extant brachyuran crabs of the world. The Raffles Bulletin of Zoology 17: 1–286.
- Nguyen LT, Schmidt HA, von Haeseler A, Minh BQ (2015) IQ-TREE: A fast and effective stochastic algorithm for estimating maximum-likelihood phylogenies. Molecular Biology and Evolution 32(1): 268–274. https://doi.org/10.1093/molbev/msu300
- Ortmann AE (1896) Das System der Decapoden-Krebse. Zoologische Jahrbucher. Abteilung fur Systematik, Geographie und Biologie der Tiere 9: 409–453.
- Pan D, Shi BY, Du SY, Gu TY, Wang RX, Xing YH, Zhang Z, Cheng JJ, Cumberlidge N, Sun HY (2022) Mitogenome phylogeny reveals Indochina Peninsula origin and spatiotemporal diversification of freshwater crabs (Potamidae: Potamiscinae) in China. Cladistics 38(1): 1–12. https://doi.org/10.1111/cla.12475
- Parisi B (1916) I Decapodi Giapponesi del Museuo di Milano. IV. Cyclometopa. Atti Societas italiano Sciences naturelle 55: 153–190.
- Pope CH (1928) Seven new reptiles from Fukien Province, China. American Museum Novitates 320: 1–6.
- Pope CH (1929) Four new frogs from Fukien Province, China. American Museum Novitates 352: 1–5.
- Rambaut A, Suchard MA, Xie D, Drummond AJ (2014) Tracer v1.6. http://beast.bio.ed.ac.uk/Tracer
- Rathbun MJ (1904) Les crabes d'eau douce. Nouvelles Archives du Muséum d'Histoire naturelle 6: 225–312.

- Rathbun MJ (1931) New and rare Chinese crabs. Lingnan Science Journal 8: 75–104.
- Ronquist F, Teslenko M, van der Mark P, Ayres DL, Darling A, Höhna S, Arget BL, Liu L, Suchard MA, Huelsenbeck JP (2012) MrBayes 3.2: Efficient Bayesian phylogenetic inference and model choice across a large model space. Systematic Biology 61(3): 539–542. https://doi.org/10.1093/sysbio/sys029
- Shih H-T, Chen GX, Wang LM (2005) A new species of freshwater crab (Decapoda: Brachyura: Potamidae) from Dongyin Island, Matsu, Taiwan, defined by morphological and molecular characters, with notes on its biogeography. Journal of Natural History 39(31): 2901–2911. https://doi.org/10.1080/00222930500214010
- Shih H-T, Hung HC, Schubart CD, Chen CA, Chang HW (2006) Intraspecific genetic diversity of the endemic freshwater crab *Candidiopotamon rathbunae* (Decapoda, Brachyura, Potamidae) reflects five million years of geological history of Taiwan. Journal of Biogeography 20(6): 980–989. https://doi.org/10.1111/j.1365-2699.2006.01472.x
- Shih H-T, Yeo DCJ, Ng PKL (2009) The collision of the Indian Plate with Asia: molecular evidence for its impact on the phylogeny of freshwater crabs (Brachyura: Potamidae). Journal of Biogeography 36(4): 703–719. https://doi.org/10.1111/j.1365-2699.2008.02024.x
- Shih H-T, Zhou XM, Chen GX, Chien IC, Ng PKL (2011) Recent vicariant and dispersal events affecting the phylogeny and biogeography of East Asian freshwater crab genus *Nanhaipotamon* (Decapoda: Potamidae). Molecular Phylogenetics and Evolution 58(3): 427–438. https://doi.org/10.1016/j.ympev.2010.11.013
- Shih H-T, Huang C, Ng PKL (2016) A re-appraisal of the widely-distributed freshwater crab genus *Sinopotamon* Bott, 1967, from China, with establishment of a new genus (Crustacea: Decapoda: Potamidae). Zootaxa 4138(2): 309–331. https://doi.org/10.11646/zootaxa.4138.2.5
- Tan QH, Zhou XJ, Zou JX (2021) Two new species of freshwater crab of the genus *Aparapotamon* Dai & Chen, 1985 (Crustacea, Brachyura, Potamidae) from Yunnan, China. ZooKeys 1056: 149–171. https://doi.org/10.3897/zookeys.1056.63755
- Yeo DCJ, Shih H-T, Meier R, Ng PKL (2007) Phylogeny and biogeography of the freshwater crab genus *Johora* (Crustacea: Brachyura: Potamidae) from the Malay Peninsula, and the origins of its insular fauna. Zoologica Scripta 36(3): 255–269. https://doi.org/10.1111/j.1463-6409.2007.00276.x