

One becomes three: An integrative morphological and molecular analysis of the windowpane oyster *Placuna* (Bivalvia: Pectinida) reveals new species

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Funding information

Hong Kong Offshore LNG Terminal Project, Grant/Award Number: MCEF22003; Lantau Conservation Fund, Grant/Award Number: RE-2020-22

Abstract

For decades, many marine animals have been considered to exhibit cosmopolitan or transoceanic distribution. This situation is prevalent in Asia, where many species were collected and named by American or European experts in the 1700s to early 1900s. Using the windowpane oysters *Placuna*—a small genus of bivalves with five recognized species—we show that careful analysis is required to reassess the validity of these species. Currently, only two species of *Placuna* (*P. placenta* and *P. ephippium*) widely reported in the Indo-Pacific region have been recorded from Chinese coastal waters. Here, we described two new species of *Placuna* from China. *Placuna vitream* sp. nov. can be distinguished from *P. placenta* by its larger ridge angle. Phylogenetic analysis using five gene fragments fully supported that *P. vitream* sp. nov. is a sister to the specimen from Singapore identified as *P. placenta* and more distant from other *Placuna* species with available molecular data. Besides, based on subfossil shells, we describe *Placuna aestuaria* sp. nov. that differs from its congeneric species by its broad hinge, medium ridge angle, and nearly straight ridges. Finally, we suggest a combination of hinge structure and ridge angle that can be used for identifying *Placuna* species and preparing a key to this genus. Our findings of two new species expand the diversity of *Placuna* and prompt reassessment of the many presumably widely distributed marine species in Asia.

KEYWORDS

Bivalvia, capiz shell, phylogeny, Placunidae, windowpane oyster

TAXONOMY CLASSIFICATION

Biodiversity ecology, Phylogenetics, Taxonomy, Zoology

1 | INTRODUCTION

Many marine species have been considered to exhibit transoceanic or even cosmopolitan distribution (Hutchings & Kupriyanova, 2018; Knowlton, 1993). This situation is especially prevalent in the

Asia-Pacific where many species were collected and named by American or European experts in the 1700s to early 1900s (Hutchings & Kupriyanova, 2018). The extensive distribution was attributed to the high uniformity of the marine water, the absence of physical barriers for larval dispersal, and the extended

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larval dispersal period observed in certain taxa (Hansen, 1980; Scheltema, 1971; Schulze et al., 2012). Reassessment of such widely distributed species, however, has shown mixed results. While some have discovered morphologically similar cryptic species such as those from chemosynthetic habitats (Bickford et al., 2007; Hutchings & Kupriyanova, 2018; Pérez-Portela et al., 2013; Wang et al., 2020), others have confirmed their wide distribution patterns (McCowin et al., 2019; Thomas et al., 2020). Nevertheless, a significant number of marine species have not undergone a thorough evaluation of their species identity and distribution, impeding our comprehension of diversity and biogeographical patterns.

The family Placunidae Rafinesque, 1815, commonly called windowpane oysters, windowpane shells, or capiz shells, serve as a compelling case for investigating species identity and distribution patterns. This family, classified in the order Pectinida Gray, 1854, inhabits predominantly the coastal waters of the Indo-Pacific. The family is monogeneric, with *Placuna* Lightfoot, 1786 as the only genus, and it contains five extant species (*P. ehippium* Retzius, 1788, *P. lincollii* Gray, 1849, *P. lobata* Sowerby, 1871, *P. placenta* Linnaeus, 1758, and *P. quadrangula* Retzius, 1788) and two fossil species (*P. mandirantjanensis* Martin, 1909 and *P. pseudoplacenta* Martin, 1909), all named between 1758 and 1871 (MolluscaBase, 2024). The type species *P. placenta*, characterized by semitransparent shells, has high commercial value: it is an edible species and its shells are widely used for crafting ornamental wares and traditional windowpanes (Gallardo et al., 1995; Rustia et al., 2023). This species is widely reported from the northern to the eastern Indian Ocean and the western to southern Pacific Ocean (MolluscaBase, 2024). In China, *P. placenta* has been reported from the intertidal zone of the northern South China Sea and the southern East China Sea (Li et al., 2019; Liu, 2008). However, our preliminary analysis of the mitochondrial cytochrome c oxidase subunit I (*cox1*) of "*P. placenta*" specimens from the Chinese coastal waters revealed Kimura 2-parameter (K2P) genetic distances >11% with a *P. placenta* sample collected from Singapore (Bieler et al., 2014). These K2P distances are much larger than the variations typically considered intraspecific for bivalves (i.e., 2.0%) (Lin et al., 2022; Yu & Li, 2012). In addition, we collected several shells of 8000–6000 years old from Hong Kong (WWF Hong Kong, 2013), which appear to come from an extinct species since we cannot find any living windowpane

oysters in that area. These live specimens and subfossil shells appear distinct from each other and the recognized species of the genus.

Therefore, this study aims to characterize the two new species of *Placuna* from China based on a rigorous molecular phylogenetic framework and morphological analysis. We describe the morphology of the two new species of *Placuna* and construct an identification key to all species of the genus. For species with soft tissues available, we amplified five gene fragments and conducted a phylogenetic study of *Placuna* spp. Our results enrich the genetic information of Placunidae and prompt reevaluation of marine bivalve species that are considered widely distributed (Jackson et al., 2015).

2 | MATERIALS AND METHODS

2.1 | Sample collection

Type specimens of *P. vitream* sp. nov. were collected from the intertidal zone of Xincun Port (18°24.55' N, 109°58.49' E), Sanya, Hainan Island, China, in November 2023. The type specimens of *P. aestuaria* sp. nov. (empty shells only) were collected from the Mai Po Nature Reserve (22°29.03' N, 114°01.56' E), Hong Kong, in July 2023, buried in the unearthed mud located in the Deep Bay area. The adductor muscles of fresh specimens were preserved using 100% ethanol for DNA extraction, and all shells were cleaned and kept in room temperature (Table 1).

2.2 | Other materials studied

More specimens of *P. vitream* sp. nov. were purchased from the fisherman in Dongmen Market, Haikou, Hainan Island, China, in May 2023, while an adductor muscle sample of *P. vitream* sp. nov. was collected from Xiajin Bay (24°30.48' N, 118°12.25' E), Xiamen, Fujian, China, in January 2023. The specimens of *P. ehippium* were collected from the same location as the *P. vitream* sp. nov. type specimens. The samples of *P. quadrangula* were collected from the intertidal zone of Bilangbilangan Island (10°14.49' N, 124°27.14' E), Philippines (Table 1).

TABLE 1 Specimens used in this study.

| Species | Specimen ID | Condition | Date | Location |
|---------------------------------|-------------------|---------------------------------------|---------|------------------------------------|
| <i>Placuna vitream</i> sp. nov. | TMBC031019-031023 | Complete individuals | 11/2023 | Sanya, Hainan, China |
| | TMBC031024-031036 | Complete individuals | 05/2023 | Haikou, Hainan, China |
| | TMBC031037 | Adductor muscle | 01/2023 | Xiamen, Fujian, China |
| <i>P. aestuaria</i> sp. nov. | TMBC031038-031057 | Subfossil paired shells | 07/2023 | Hong Kong, China |
| <i>P. ehippium</i> | TMBC031058 | Left shell | 11/2023 | Sanya, Hainan, China |
| | TMBC031059-031060 | Paired shell | 12/2023 | Sanya, Hainan, China |
| | TMBC031061-031063 | Complete individual | 12/2023 | Sanya, Hainan, China |
| <i>P. quadrangula</i> | TMBC031064-031065 | Paired shells | Unknown | Mactan Island, Philippines |
| | TMBC031066 | Shells with air-dried adductor muscle | Unknown | Bilangbilangan Island, Philippines |

2.3 | Morphological measurement and photography

The *Placuna* shells were measured using a vernier caliper to determine the shell length (L), height (H), hinge length (HL), hinge height (HH), anterior hinge length (AHL), anterior ridges length (ARL), posterior ridges length (PRL), scar length (SL), anterior length (AL), dorsal height (DH), and ridge angle (RA) (Figure 1). The specimens were photographed using an EOS 5D Mark IV camera (Canon, Japan), and their details were observed using a digital Stereo Microscope MZ1270i Imaging System (Nikon, Japan). The type specimens used in this study were deposited in the Tropical Marine Biodiversity Collections of the South China Sea (TMBC), Chinese Academy of Sciences, Guangzhou, China.

2.4 | DNA extraction and PCR

The genomic DNA of *P. vitream* sp. nov., *P. ephippium*, and *P. quadrangula* was extracted from the adductor muscle using the CTAB method (Stewart & Via, 1993). The genomic DNA quality was determined using agarose gel (1.0%) electrophoresis and quantified using a NanoDrop ND-1000 spectrophotometer (Thermo Scientific, USA). Three nuclear (*18S rRNA*, *28S rRNA*, and *histone H3*) and two mitochondrial (*cox1* and *16S rRNA*) maker genes were amplified using KOD One PCR Master Mix (Toyobo, Japan) following the manufacturer's

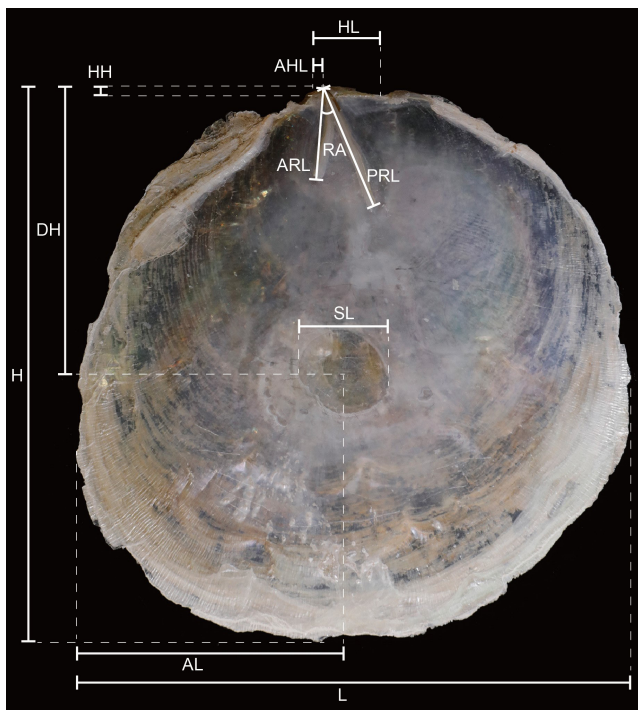


FIGURE 1 Schematic view of the *Placuna* shell. AL, anterior length; AHL, anterior hinge length; ARL, anterior ridges length; DH, dorsal height; H, height; HH, hinge height; HL, hinge length; L, length; PRL, posterior ridges length; RA, ridge angle; SL, scar length.

protocol. The following primers were used: LCO1490 and HCO2198 (5'-GGTCAACAAATCATAAAGATATTGG-3'/5'-TAAACTTCAGGTGACCAAAAAATCA-3') (Folmer et al., 1994) for cytochrome oxidase I (*cox1*), LRJ and 16SA (5'-CTCCGGTTTGAAGCTCAGATCA-3'/5'-ATGTTTGTGATAAACAGGCG-3') (Baco-Taylor, 2002; Ratnasingham & Hebert, 2007) for *16S rRNA*, F19 and R1843 (5'-ACCTGGTTGATCCTGCCA-3'/5'-GGATCCAAGCTTGATCCTTCTGCAGGTTACCTAC-3') (Elwood et al., 1985; Turbeville et al., 1994) for *18S rRNA*, D1R and LSUB (5'-ACCCGCTGAATTTAAGCATA-3'/5'-ACGAACGATTTGCACGTCAG-3') (Litaker et al., 2003; Scholin et al., 1994) for *28S rRNA*, and H3F and H3R (5'-ATGGCTCGTACCAAGCAGACGC-3'/5'-ATATCCTTGCCATATGTGAC-3') (Colgan et al., 1998) for *histone H3*. The PCR products were bidirectionally sequenced on an ABI PRISM 3730xl DNA Analyzer (Thermo Fisher Scientific). The sequences for phylogenetic analyses were assembled using SeqMan (DNASTAR).

2.5 | Phylogenetic analyses and genetic distance estimation

Phylogenetic analyses were conducted to determine the position of *P. vitream* sp. nov. based on the five abovementioned gene fragments. Available sequences of 22 representative species from the order Pectinida and Limidae (outgroup) were downloaded from GenBank (<https://www.ncbi.nlm.nih.gov/>) (Table 2). The longest fragment for each gene was selected if one species had two or more sequence records. The analyses were conducted using PhyloSuite v1.2.2 (Zhang et al., 2020) with several plug-in programs: (1) MAFFT v7.520 (Katoh & Standley, 2013) under the "auto" option was applied to align each gene fragment with the "Normal alignment" mode; (2) Gblocks v0.91b (Talavera & Castresana, 2007) was applied to remove ambiguously aligned fragments in batches, with missing genes or alignment gaps filled with "-"; (3) Then, the fragments were concatenated and ModelFinder v1.5.4 (Kalyaanamoorthy et al., 2017) was used to select the best-fit model according to the BIC criterion; (4) Bayesian inference (BI) and maximum-likelihood (ML) analyses were conducted using MrBayes v3.2.6 (Ronquist et al., 2012) and IQ-TREE2 v2.1.2 (Nguyen et al., 2015) with the GTR+R3+F model, under the partition model for 10 million generations and 100 thousand ultrafast bootstraps, respectively (Minh et al., 2013). The pairwise genetic distances between different species for each gene were estimated using the Kimura 2-parameter (K2P) model implemented in MEGA v7.0 (Kumar et al., 2016).

3 | RESULTS

3.1 | Systematics

Order: Pectinida Gray, 1854.

Superfamily: Anomioidea Rafinesque, 1815.

Family: Placunidae Rafinesque, 1815.

TABLE 2 Genbank accession numbers of the gene fragments used in the genetic distance calculations and phylogenetic analyses.

| Family | Genus | Species | cox1 | 16S rRNA | 18S rRNA | 28S rRNA | Histone H3 | |
|-----------------|---------------------|-----------------------------|----------------------|----------|----------|----------|------------|----------|
| Placunidae | <i>Placuna</i> | <i>vitreum</i> nov. sp. HT1 | PP711111 | PP599757 | PP599752 | PP599761 | PP663639 | |
| | | <i>vitreum</i> nov. sp. HT2 | PQ008987 | PQ032325 | PQ032329 | PQ008442 | PQ030828 | |
| | | <i>vitreum</i> nov. sp. HT3 | PQ008988 | PQ032326 | PQ032330 | PQ008443 | PQ030829 | |
| | | <i>vitreum</i> nov. sp. HT4 | PQ008989 | PQ032327 | PQ032331 | PQ008444 | PQ030830 | |
| | | <i>vitreum</i> nov. sp. HT5 | PQ008990 | PQ032328 | PQ032332 | PQ008445 | PQ030831 | |
| | | <i>vitreum</i> nov. sp. PT1 | PP711110 | PP599756 | PP599751 | PP599762 | PP663638 | |
| | | <i>vitreum</i> nov. sp. PT2 | PP711112 | PP599758 | PP599753 | PP599763 | PP663640 | |
| | | <i>ephippium</i> | PP711114 | PP599760 | PP599755 | PP599765 | PP663642 | |
| | | <i>quadrangula</i> | PP711113 | PP599759 | PP599754 | PP599764 | PP663641 | |
| | | <i>placenta</i> | KC429104 | HQ840731 | KC429343 | KC429442 | KC429180 | |
| | sp. HS0121 | MT896307 | / | / | / | / | | |
| Anomiidae | <i>Anomia</i> | <i>simplex</i> | KF850693 | JN133626 | / | / | / | |
| | | <i>ephippium</i> | KF369196 | KX713191 | AF120535 | KX713358 | KX713513 | |
| | | <i>chinensis</i> | MN608245 | / | / | AB105361 | / | |
| | | sp. FP2010 | GQ166573 | GQ166557 | / | / | / | |
| | | <i>Pododesmus</i> | <i>caelata</i> | / | / | AJ389650 | AJ307555 | / |
| | | | <i>patelliformis</i> | / | KC429261 | KC429342 | KC429441 | KC429179 |
| Dimyidae | <i>Dimya</i> | <i>lima</i> | / | KX713213 | KC429344 | KX713375 | KC429181 | |
| | | sp. DJC-2016 | / | / | KX713288 | KX713376 | KX713532 | |
| Entoliidae | <i>Pectinella</i> | <i>aequoris</i> | / | / | / | MH464049 | MH464038 | |
| Plicatuloidea | <i>Plicatula</i> | sp. DJC-2016 | / | / | KX713337 | KX713424 | KX713573 | |
| | | <i>australis</i> | / | / | AF229626 | AB102737 | KC429178 | |
| Spondylidae | <i>Spondylus</i> | <i>gaederopus</i> | JF496776 | KR676345 | KT757808 | KT757854 | KT757896 | |
| Propeamussiidae | <i>Parvamussium</i> | <i>torresi</i> | / | MH464019 | MH464099 | MH464043 | MH464032 | |
| | | sp. VLG-2013 | KC429103 | KC429259 | KC429340 | KC429437 | KC429176 | |
| Pectinidae | <i>Argopecten</i> | <i>purpuratus</i> | KP265825 | JN848518 | EU660809 | / | EU379526 | |
| | | <i>adamussium</i> | / | HM600752 | MH464058 | FJ263652 | EU379491 | |
| | | <i>chlamys</i> | / | FJ263648 | MH464068 | FJ263658 | FJ263667 | |
| | | <i>crassadoma</i> | / | EU379444 | L49050 | FJ263654 | EU379498 | |
| | | <i>flexopecten</i> | HQ197900 | MH490816 | AJ389662 | AJ307545 | JQ611569 | |
| | | <i>pecten</i> | KC429102 | X82501 | L49053 | KC429436 | KC429175 | |
| Outgroup | <i>Lima</i> | <i>lima</i> | AF120649 | KC429257 | KC429339 | AJ307558 | JQ611555 | |

Genus: *Placuna* Lightfoot, 1786.

Synonyms: *Ehippium* Röding, 1798; *Placenta* (Retzius, 1788);

Sellaria Link, 1807.

Type species: *Placuna placenta* Linnaeus, 1758.

Synonyms: *Anomia placenta* Linnaeus, 1758; *Ehippium transparens* Röding, 1798; *Placenta orbicularis* Retzius, 1788; *Placenta auriculata* Mörch, 1853; *Placenta communis* Megerle von Mühlfeld, 1811.

Diagnosis (modified from Matsukuma, 1987): Shell inequivalve, thin, very compressed, translucent to opaque, slightly fragile. Valves roughly subquadrate to subcircular, left valve more convex, ventral margin rounded. External surface lamellate, periostracum absent, internal surface smooth. Inverted V-shaped hinge ridges and ligaments equal or unequal. Adductor muscle scar subcircular, close to valve center.

3.2 | *Placuna vitream* sp. nov

<https://zoobank.org/NomenclaturalActs/cb26877e-51dd-4330-9de5-3bce8b151c1c>

3.2.1 | Type materials and locality

Holotype (TMBC031019), paratypes 1–4 (TMBC031020–031023), collected from the type locality Xincun Port (18°24.55' N, 109°58.49' E), Sanya City, Hainan Island, China by Mr. Yi-Tao Lin and Mr. Junhao Pan in May 2023. Paratypes 5–18 (TMBC031024–031036), collected from Dongmen Market (20°2'26", 110°20'45"), Haikou City, Hainan Island, China by Mr. Xiao Han, Dr. Yanjie Zhang, and Mr. Juhao Wang in May 2023.

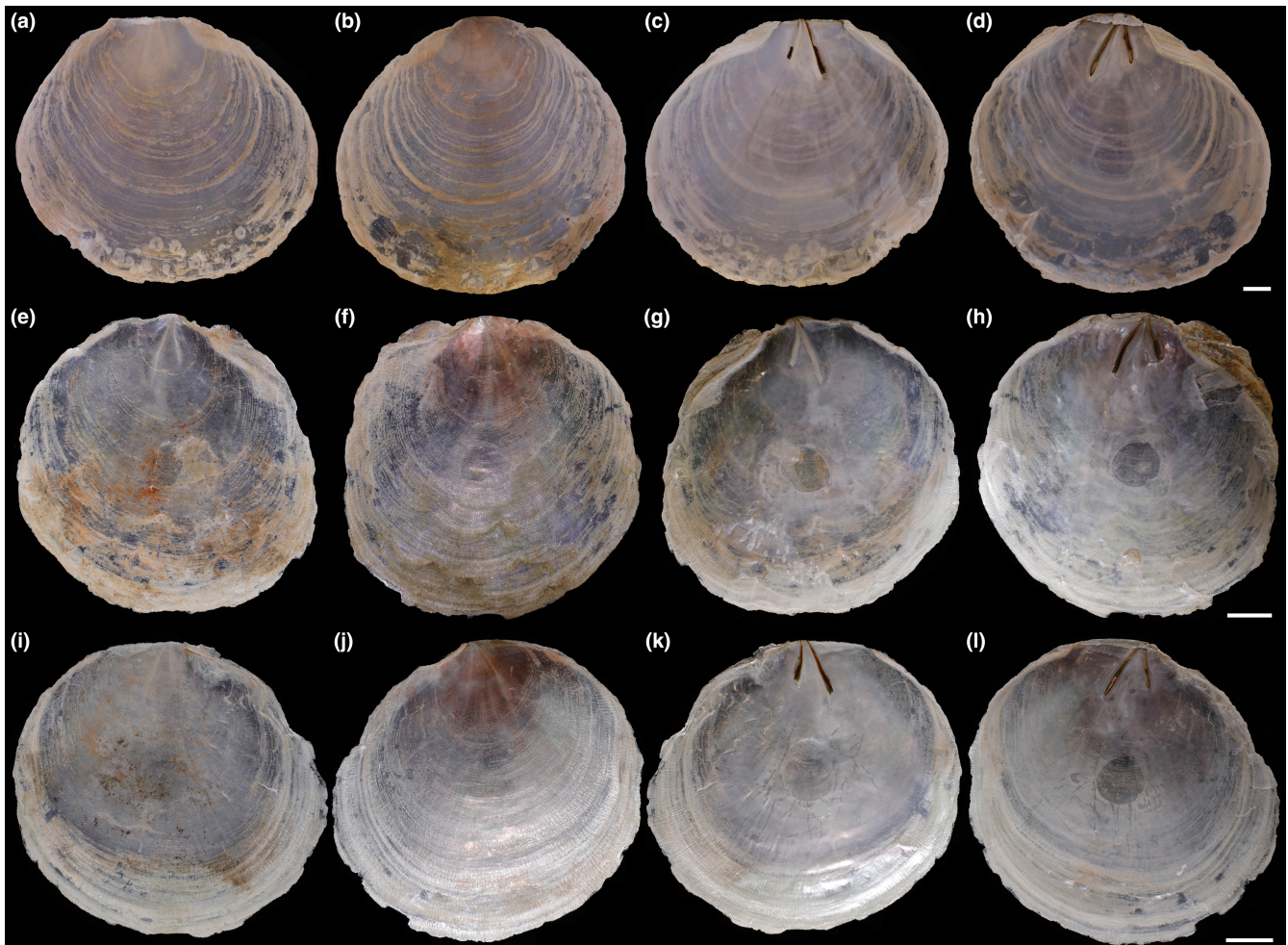


FIGURE 2 External and internal views of the left and right valves from three pairs of *Placuna vitream* sp. nov. specimens. (a–d) External view of left valve, external view of right valve, internal view of left valve, internal view of right valve of the holotype (TMBC031019), respectively; (e–h) External view of left valve, external view of right valve, internal view of left valve, internal view of right valve of the paratype 1 (TMBC031020), respectively; (i–l) External view of left valve, external view of right valve, internal view of left valve, internal view of right valve of the paratype 2 (TMBC031021), respectively. Scale bar: 10mm.

3.2.2 | Distribution

Currently known from Xincun Port, Sanya, and Xiajin Bay, Xiamen in China.

3.2.3 | Etymology

The epithet “*vitream*” refers to this species’ translucent and pearl-like glittery shells.

3.2.4 | Diagnosis

Shell up to 110mm, subcircular and translucent. Hinge slim with clear hinge teeth. Umbones slightly prominent, close to the anterior end of the hinge. Auricles obvious, anterior auricle larger than posterior auricle. Ridges angle moderate from 28° to 31°.

Hinge ridges and ligaments slightly curved; anterior ridge shorter than posterior ridge.

3.2.5 | Description

Shell (Figures 2 and 3a,b, Tables 3 and 4) subcircular, translucent, and very compressed. Left valve more convex than right valve. External surface with growth lines, mauve dorsal, and gray lamella, without radial lines. Length up to 110mm, nearly equal to height ($L/H=0.960-1.050$) and approximate equilateral ($AL/L=0.450-0.507$). Umbones slightly prominent, close to anterior end of hinge ($AHL/HL=0.205-0.325$). Auricles obvious, anterior auricle larger than posterior auricle. Hinge slim ($HH/HL=0.082-0.135$), slightly rounded, with obscure hinge teeth or without teeth. Hinge ridges and ligaments slightly curved, inverted V-shaped, with moderate ridge angle (RA) from 28° to 31°. Anterior hinge ridge and ligament shorter than posterior ridge ($ARL/PRL=0.594-0.761$). Anterior

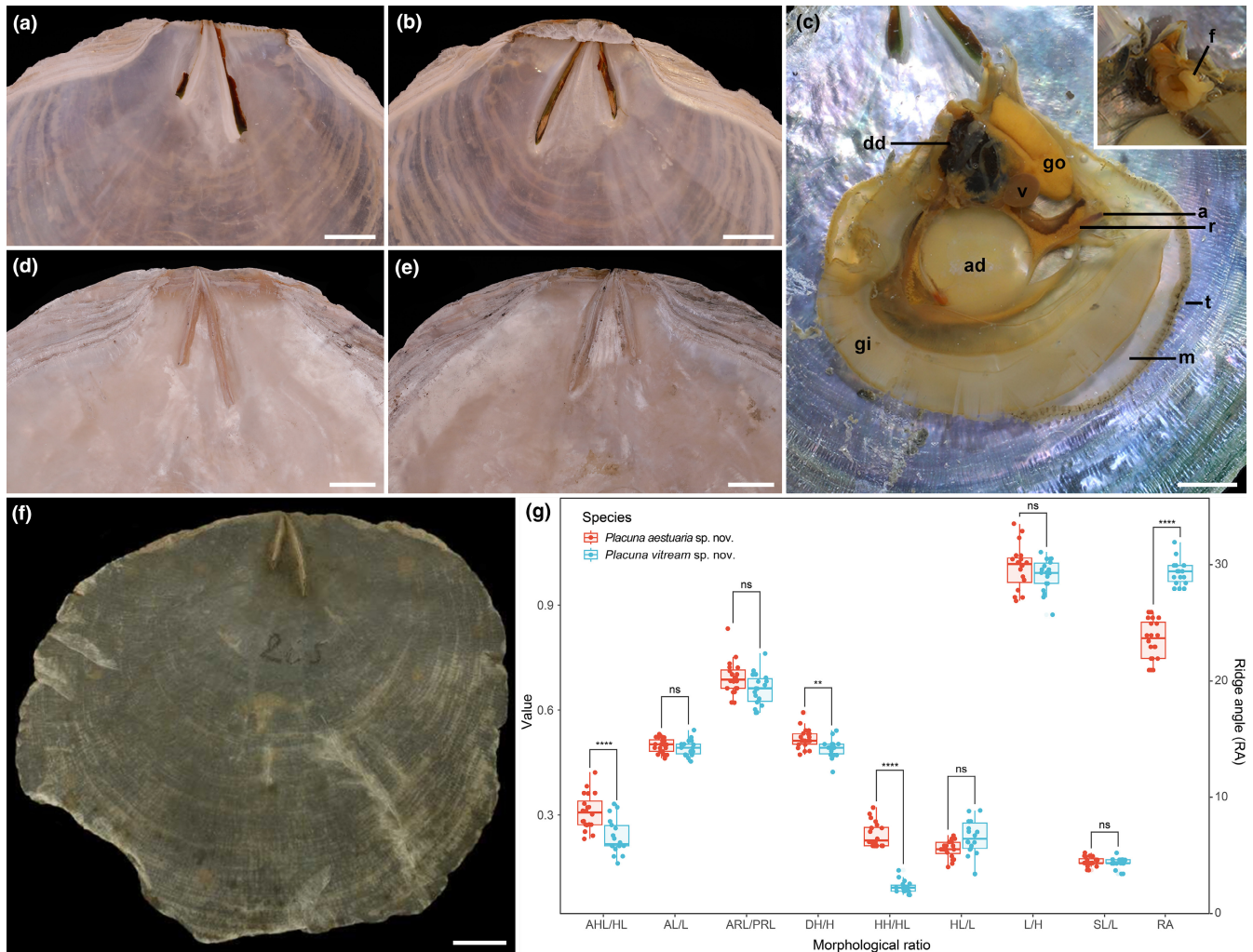


FIGURE 3 Internal views of three *Placuna* species and the statistical study of morphological features of the two new species. (a,b) The left and right valves of *P. vitream* sp. nov. holotype (TMBC031019); (c) Anatomy of *P. vitream* sp. nov. paratype 2 (TMBC031021); (d,e) The left and right valves of *P. aestuaria* sp. nov. holotype (TMBC031038). (f) The type specimen of *Placuna placenta* stored in the Linnean Society of London (<https://linnean-online.org/>) for morphological comparison. (g) Statistical study of the shell morphological indices and ridge angles between two new species. The details are shown in Tables 3 and 4. AL, anterior length; AHL, anterior hinge length; ARL, anterior ridges length; a, anus; ad, adductor muscle; dd, digestive diverticula; DH, dorsal height; f, foot; gi, gill; go, gonad; H, height; HH, hinge height; HL, hinge length; L, length; m, mantle; PRL, posterior ridges length; r, rectum; RA, ridge angle; SL, scar length; t, tentacle; v, ventricle. Scale bar: 10 mm.

pedal retractor scar oval, close to middle between ligament posterior ends. Adductor muscle scar subcircular, close to valve center.

Anatomy (Figure 3c): Mantle large, thin, and semitransparent, with distinct mantle edge and row of tentacles at the margin. Large and C-shaped gill at anterior side. Digestive diverticula subcircular. Adductor muscle large, circular, near shell center. Foot small, anteriorly located between digestive diverticula and adductor muscle. Ventricle circular, posteriorly located between digestive diverticula and adductor muscle. Gonad folded scrotiform, posterior to digestive diverticula.

3.2.6 | Remarks

Within the genus *Placuna*, *P. vitream* sp. nov. can be distinguished from *P. ephippium*, *P. quadrangula*, *P. lincolni*, and *P. lobata* by its

unequal length of the hinge ridges and ligaments (Das et al., 2019; Dunker, 1879; Matsukuma, 1987) (Figures 3a,b and 6, Tables 3 and 4). The AHL/HL ratio of *P. vitream* sp. nov. is much smaller than that of *P. ephippium* and *P. quadrangula*, which means the umbones of the former are located at the anterior end of the hinge, whereas those of the latter two are located near the hinge center. Although such data are unavailable for *P. lincolni* and *P. lobata*, previous studies showed that their umbones are in the middle of the hinge (Dunker, 1879; Gray, 1849; Matsukuma, 1987). *Placuna vitream* sp. nov. and two fossil species *P. pseudoplacenta* and *P. mandirantjanensis* exhibit similar shell shapes, outlines, and V-shaped hinges. However, the ridge angles of the two fossil species are substantially larger ($RA > 60^\circ$) (Martin, 1909) than *P. vitream* sp. nov. (28° – 31°), and the anterior hinge ridge of *P. vitream* sp. nov. is straighter. Notably, *P. vitream* sp. nov. is morphologically most

TABLE 3 Measurements of the left valves of the specimens (in mm where applicable).

| Species | Specimen ID | Length (L) | Height (H) | Hinge length (HL) | Hinge height (HH) | Anterior hinge length (AHL) | Anterior ridge length (ARL) | Posterior ridge length (PRL) | Scar length (SL) | Anterior length (AL) | Dorsal height (DH) | Ridge angle (RA) |
|---------------------------------|-------------|------------|------------|-------------------|-------------------|-----------------------------|-----------------------------|------------------------------|------------------|----------------------|--------------------|------------------|
| <i>Placuma vitream</i> sp. nov. | TMBC031019 | 108.3 | 103.1 | 23.7 | 3.2 | 6.1 | 18.6 | 26.3 | 17.4 | 53.7 | 55.6 | 29.5 |
| | TMBC031020 | 65.7 | 64.5 | 12.6 | 1.5 | 4.1 | 9.6 | 15.1 | 10.5 | 33.0 | 30.9 | 29.5 |
| | TMBC031021 | 64.4 | 65.2 | 18.2 | 1.6 | 4.0 | 9.3 | 15.0 | 10.1 | 30.8 | 32.4 | 29.0 |
| | TMBC031022 | 57.2 | 62.5 | 14.8 | 1.3 | 3.5 | 9.4 | 14.0 | 9.9 | 29.0 | 30.3 | 28.5 |
| | TMBC031023 | 59.3 | 61.6 | 18.2 | 1.5 | 3.9 | 8.2 | 13.8 | 9.2 | 27.9 | 30.4 | 30.0 |
| | TMBC031024 | 55.8 | 58.1 | 14.0 | 1.3 | 3.2 | 8.0 | 13.4 | 9.5 | 25.1 | 29.2 | 31.0 |
| | TMBC031025 | 50.3 | 50.7 | 12.2 | 1.0 | 2.5 | 6.7 | 10.1 | 8.0 | 24.5 | 24.8 | 30.0 |
| | TMBC031026 | 106.2 | 107.2 | 23.4 | 2.6 | 6.2 | 16.8 | 24.3 | 18.2 | 53.1 | 53.4 | 28.0 |
| | TMBC031027 | 104.0 | 100.8 | 21.2 | 2.0 | 6.8 | 16.6 | 21.8 | 19.5 | 48.8 | 52.9 | 28.0 |
| | TMBC031028 | 63.9 | 68.4 | 12.9 | 1.2 | 4.0 | 10.2 | 15.0 | 8.5 | 29.9 | 33.5 | 29.0 |
| | TMBC031029 | 58.2 | 57.9 | 12.8 | 1.1 | 2.3 | 8.0 | 12.1 | 10.0 | 30.4 | 27.0 | 29.5 |
| | TMBC031030 | 56.8 | 55.3 | 17.6 | 1.3 | 2.9 | 7.3 | 11.2 | 7.4 | 27.2 | 25.3 | 28.0 |
| | TMBC031031 | 57.2 | 58.3 | 14.1 | 1.1 | 2.5 | 7.8 | 11.2 | 9.1 | 26.5 | 27.5 | 30.0 |
| | TMBC031032 | 51.0 | 58.8 | 14.3 | 1.2 | 2.9 | 8.9 | 12.7 | 8.0 | 24.8 | 29.5 | 28.5 |
| | TMBC031033 | 55.1 | 55.0 | 15.5 | 1.2 | 3.2 | 7.6 | 12.5 | 9.3 | 27.8 | 26.9 | 32.0 |
| | TMBC031034 | 51.5 | 49.8 | 10.9 | 0.8 | 3.0 | 6.9 | 10.1 | 8.2 | 24.9 | 23.2 | 30.0 |
| TMBC031035 | 55.2 | 55.0 | 10.0 | 1.0 | 2.1 | 7.1 | 11.2 | 8.9 | 28.0 | 27.0 | 29.5 | |
| TMBC031036 | 58.2 | 62.2 | 7.3 | 0.7 | 1.5 | 7.0 | 11.8 | 8.3 | 31.4 | 26.3 | 29.0 | |
| <i>P. aestuaria</i> sp. nov. | TMBC031038 | 109.8 | 108.1 | 18.7 | 4.0 | 6.2 | 18.9 | 22.8 | 18.3 | 53.8 | 54.6 | 23.0 |
| | TMBC031039 | 107.2 | 116.0 | 22.0 | 5.7 | 7.0 | 18.5 | 28.0 | 17.0 | 54.0 | 60.4 | 26.0 |
| | TMBC031040 | 81.4 | 84.1 | 11.8 | 3.5 | 3.5 | 15.2 | 24.4 | 12.9 | 42.3 | 43.0 | 24.0 |
| | TMBC031041 | 121.1 | 133.2 | 24.6 | 7.2 | 10.3 | 30.0 | 41.6 | 21.1 | 62.5 | 78.0 | 25.0 |
| | TMBC031042 | 78.2 | 77.3 | 14.4 | 3.0 | 4.5 | 11.8 | 17.2 | 12.3 | 38.5 | 37.0 | 24.0 |
| | TMBC031043 | 95.7 | 95.0 | 18.8 | 4.2 | 5.0 | 16.7 | 24.5 | 15.0 | 45.2 | 48.0 | 25.5 |
| | TMBC031044 | 92.5 | 89.8 | 14.6 | 4.7 | 5.3 | 14.8 | 20.5 | 16.8 | 45.2 | 45.9 | 26.0 |
| | TMBC031045 | 92.2 | 81.8 | 21.0 | 4.5 | 5.0 | 13.5 | 19.2 | 14.8 | 44.6 | 43.9 | 22.0 |
| | TMBC031046 | 87.4 | 94.9 | 18.4 | 4.8 | 6.9 | 16.2 | 24.8 | 15.5 | 40.2 | 47.0 | 25.0 |
| | TMBC031047 | 86.5 | 78.0 | 15.1 | 4.3 | 4.1 | 13.5 | 21.9 | 12.0 | 44.0 | 41.7 | 23.0 |
| | TMBC031048 | 85.8 | 78.9 | 19.1 | 4.9 | 4.8 | 12.0 | 18.2 | 14.8 | 43.2 | 41.8 | 25.5 |
| | TMBC031049 | 82.9 | 81.0 | 18.0 | 3.8 | 6.5 | 13.0 | 18.2 | 12.6 | 41.5 | 40.9 | 21.0 |
| | TMBC031050 | 79.6 | 77.5 | 16.8 | 3.7 | 4.6 | 12.9 | 17.3 | 12.4 | 41.0 | 40.0 | 21.0 |

(Continues)

TABLE 3 (Continued)

| Species | Specimen ID | Length (L) | Height (H) | Hinge length (HL) | Hinge height (HH) | Anterior hinge length (AHL) | Anterior ridge length (ARL) | Posterior ridge length (PRL) | Scar length (SL) | Anterior length (AL) | Dorsal height (DH) | Ridge angle (RA) |
|-----------------------|-------------|------------|------------|-------------------|-------------------|-----------------------------|-----------------------------|------------------------------|------------------|----------------------|--------------------|------------------|
| | TMBC031051 | 75.0 | 79.8 | 15.3 | 3.2 | 4.3 | 13.0 | 18.9 | 10.7 | 39.5 | 38.0 | 23.0 |
| | TMBC031052 | 78.0 | 79.2 | 18.9 | 4.8 | 6.0 | 14.7 | 20.1 | 12.3 | 36.8 | 42.7 | 22.0 |
| | TMBC031053 | 72.4 | 69.7 | 14.8 | 3.1 | 5.4 | 10.3 | 15.5 | 11.2 | 34.8 | 33.0 | 25.5 |
| | TMBC031054 | 69.9 | 67.0 | 14.0 | 3.0 | 3.9 | 10.1 | 15.6 | 11.7 | 35.6 | 33.0 | 23.5 |
| | TMBC031055 | 104.1 | 104.5 | 23.5 | 5.3 | 7.3 | 18.7 | 27.6 | 17.5 | 52.5 | 58.1 | 24.0 |
| | TMBC031056 | 86.1 | 91.3 | 20.2 | 5.4 | 5.5 | 17.7 | 25.2 | 16.0 | 45.2 | 48.4 | 22.0 |
| | TMBC031057 | 69.8 | 68.1 | 13.0 | 2.8 | 3.0 | 10.2 | 15.1 | 12.6 | 33.0 | 34.4 | 21.0 |
| <i>P. ephippium</i> | TMBC031058 | 156.8 | 105.9 | 20.7 | 1.7 | 10.0 | 20.0 | 20.7 | 17.7 | 78.5 | 53.0 | 68.0 |
| | TMBC031059 | 122.2 | 101.7 | 27.0 | 2.3 | 13.8 | 17.7 | 18.7 | 14.9 | 67.0 | 48.2 | 70.0 |
| | TMBC031060 | 101.2 | 87.8 | 24.4 | 2.3 | 12.2 | 13.6 | 15.2 | 14.2 | 53.0 | 43.3 | 69.0 |
| | TMBC031061 | 89.8 | 78.9 | 23.2 | 1.9 | 8.2 | 12.9 | 13.2 | 12.9 | 47.0 | 36.3 | 66.0 |
| | TMBC031062 | 91.2 | 87.4 | 25.8 | 2.8 | 13.5 | 14.5 | 17.0 | 11.3 | 47.2 | 38.7 | 66.0 |
| | TMBC031063 | 119.8 | 107.7 | 17.4 | 1.4 | 8.1 | 14.5 | 15.6 | 13.4 | 60.3 | 42.6 | 70.0 |
| <i>P. quadrangula</i> | TMBC031064 | 80.9 | 69.7 | 19.0 | 2.1 | 9.5 | 10.5 | 10.4 | 10.7 | 39.7 | 28.0 | 75.0 |
| | TMBC031065 | 81.9 | 71.2 | 19.7 | 2.0 | 10.1 | 11.2 | 12.0 | 10.4 | 40.2 | 31.5 | 75.0 |
| | TMBC031066 | 83.8 | 72.7 | 20.8 | 2.2 | 10.3 | 10.7 | 11.5 | 10.3 | 44.0 | 28.0 | 80.0 |

TABLE 4 Shell morphological ratios of the specimens used in this study.

| Species | Specimen ID | L/H | HL/L | HH/HL | AHL/HL | ARL/PRL | SL/L | AL/L | DH/H |
|------------------------------|-------------|-------|-------|-------|--------|---------|-------|-------|-------|
| <i>P. vitream</i> sp. nov. | TMBC031019 | 1.050 | 0.219 | 0.135 | 0.257 | 0.707 | 0.161 | 0.496 | 0.539 |
| | TMBC031020 | 1.019 | 0.192 | 0.119 | 0.325 | 0.636 | 0.160 | 0.502 | 0.479 |
| | TMBC031021 | 1.032 | 0.204 | 0.094 | 0.321 | 0.761 | 0.188 | 0.469 | 0.525 |
| | TMBC031022 | 0.915 | 0.259 | 0.088 | 0.236 | 0.671 | 0.173 | 0.507 | 0.485 |
| | TMBC031023 | 0.963 | 0.307 | 0.082 | 0.214 | 0.594 | 0.155 | 0.470 | 0.494 |
| | TMBC031024 | 0.960 | 0.251 | 0.093 | 0.229 | 0.597 | 0.170 | 0.450 | 0.503 |
| | TMBC031025 | 0.992 | 0.243 | 0.082 | 0.205 | 0.663 | 0.159 | 0.487 | 0.489 |
| | TMBC031026 | 0.991 | 0.220 | 0.111 | 0.265 | 0.691 | 0.171 | 0.500 | 0.498 |
| | TMBC031027 | 0.988 | 0.283 | 0.088 | 0.220 | 0.620 | 0.157 | 0.478 | 0.497 |
| | TMBC031028 | 0.934 | 0.202 | 0.093 | 0.310 | 0.680 | 0.133 | 0.468 | 0.490 |
| | TMBC031029 | 1.005 | 0.220 | 0.086 | 0.180 | 0.661 | 0.172 | 0.522 | 0.466 |
| | TMBC031030 | 1.027 | 0.310 | 0.074 | 0.165 | 0.652 | 0.130 | 0.479 | 0.458 |
| | TMBC031031 | 0.981 | 0.247 | 0.078 | 0.177 | 0.696 | 0.159 | 0.463 | 0.472 |
| | TMBC031032 | 0.867 | 0.280 | 0.084 | 0.203 | 0.701 | 0.157 | 0.486 | 0.502 |
| | TMBC031033 | 1.002 | 0.281 | 0.077 | 0.206 | 0.608 | 0.169 | 0.505 | 0.489 |
| | TMBC031034 | 1.034 | 0.212 | 0.073 | 0.275 | 0.683 | 0.159 | 0.483 | 0.466 |
| TMBC031035 | 1.004 | 0.181 | 0.100 | 0.210 | 0.634 | 0.161 | 0.507 | 0.491 | |
| TMBC031036 | 0.936 | 0.125 | 0.096 | 0.205 | 0.593 | 0.143 | 0.540 | 0.423 | |
| <i>P. aestuaria</i> sp. nov. | TMBC031038 | 1.016 | 0.170 | 0.214 | 0.332 | 0.829 | 0.167 | 0.490 | 0.505 |
| | TMBC031039 | 0.924 | 0.205 | 0.259 | 0.318 | 0.661 | 0.159 | 0.504 | 0.521 |
| | TMBC031040 | 0.968 | 0.145 | 0.297 | 0.297 | 0.623 | 0.158 | 0.520 | 0.511 |
| | TMBC031041 | 0.909 | 0.203 | 0.293 | 0.419 | 0.721 | 0.174 | 0.516 | 0.586 |
| | TMBC031042 | 1.012 | 0.184 | 0.208 | 0.313 | 0.686 | 0.157 | 0.492 | 0.479 |
| | TMBC031043 | 1.007 | 0.196 | 0.223 | 0.266 | 0.682 | 0.157 | 0.472 | 0.505 |
| | TMBC031044 | 1.030 | 0.158 | 0.322 | 0.363 | 0.722 | 0.182 | 0.489 | 0.511 |
| | TMBC031045 | 1.127 | 0.228 | 0.214 | 0.238 | 0.703 | 0.161 | 0.484 | 0.537 |
| | TMBC031046 | 0.921 | 0.211 | 0.261 | 0.375 | 0.653 | 0.177 | 0.460 | 0.495 |
| | TMBC031047 | 1.109 | 0.175 | 0.285 | 0.272 | 0.616 | 0.139 | 0.509 | 0.535 |
| | TMBC031048 | 1.087 | 0.223 | 0.257 | 0.251 | 0.659 | 0.172 | 0.503 | 0.530 |
| | TMBC031049 | 1.023 | 0.217 | 0.211 | 0.361 | 0.714 | 0.152 | 0.501 | 0.505 |
| | TMBC031050 | 1.027 | 0.211 | 0.220 | 0.274 | 0.746 | 0.156 | 0.515 | 0.516 |
| | TMBC031051 | 0.940 | 0.204 | 0.209 | 0.281 | 0.688 | 0.143 | 0.527 | 0.476 |
| | TMBC031052 | 0.985 | 0.242 | 0.254 | 0.317 | 0.731 | 0.158 | 0.472 | 0.539 |
| | TMBC031053 | 1.039 | 0.204 | 0.209 | 0.365 | 0.665 | 0.155 | 0.481 | 0.473 |
| TMBC031054 | 1.043 | 0.200 | 0.214 | 0.279 | 0.647 | 0.167 | 0.509 | 0.493 | |
| TMBC031055 | 0.996 | 0.226 | 0.226 | 0.311 | 0.678 | 0.168 | 0.504 | 0.556 | |
| TMBC031056 | 0.943 | 0.235 | 0.267 | 0.272 | 0.702 | 0.186 | 0.525 | 0.530 | |
| TMBC031057 | 1.025 | 0.186 | 0.215 | 0.231 | 0.675 | 0.181 | 0.473 | 0.505 | |
| <i>P. ehippium</i> | TMBC031058 | 1.481 | 0.132 | 0.082 | 0.483 | 0.966 | 0.113 | 0.501 | 0.500 |
| | TMBC031059 | 1.202 | 0.221 | 0.085 | 0.511 | 0.947 | 0.122 | 0.548 | 0.474 |
| | TMBC031060 | 1.153 | 0.241 | 0.094 | 0.500 | 0.895 | 0.140 | 0.524 | 0.493 |
| | TMBC031061 | 1.138 | 0.258 | 0.082 | 0.353 | 0.977 | 0.144 | 0.523 | 0.460 |
| | TMBC031062 | 1.043 | 0.283 | 0.109 | 0.523 | 0.853 | 0.124 | 0.518 | 0.443 |
| | TMBC031063 | 1.112 | 0.145 | 0.080 | 0.466 | 0.929 | 0.112 | 0.503 | 0.396 |
| <i>P. quadrangula</i> | TMBC031064 | 1.161 | 0.235 | 0.111 | 0.500 | 1.010 | 0.132 | 0.491 | 0.402 |
| | TMBC031065 | 1.150 | 0.241 | 0.102 | 0.513 | 0.933 | 0.127 | 0.491 | 0.442 |
| | TMBC031066 | 1.153 | 0.248 | 0.106 | 0.495 | 0.930 | 0.123 | 0.525 | 0.385 |

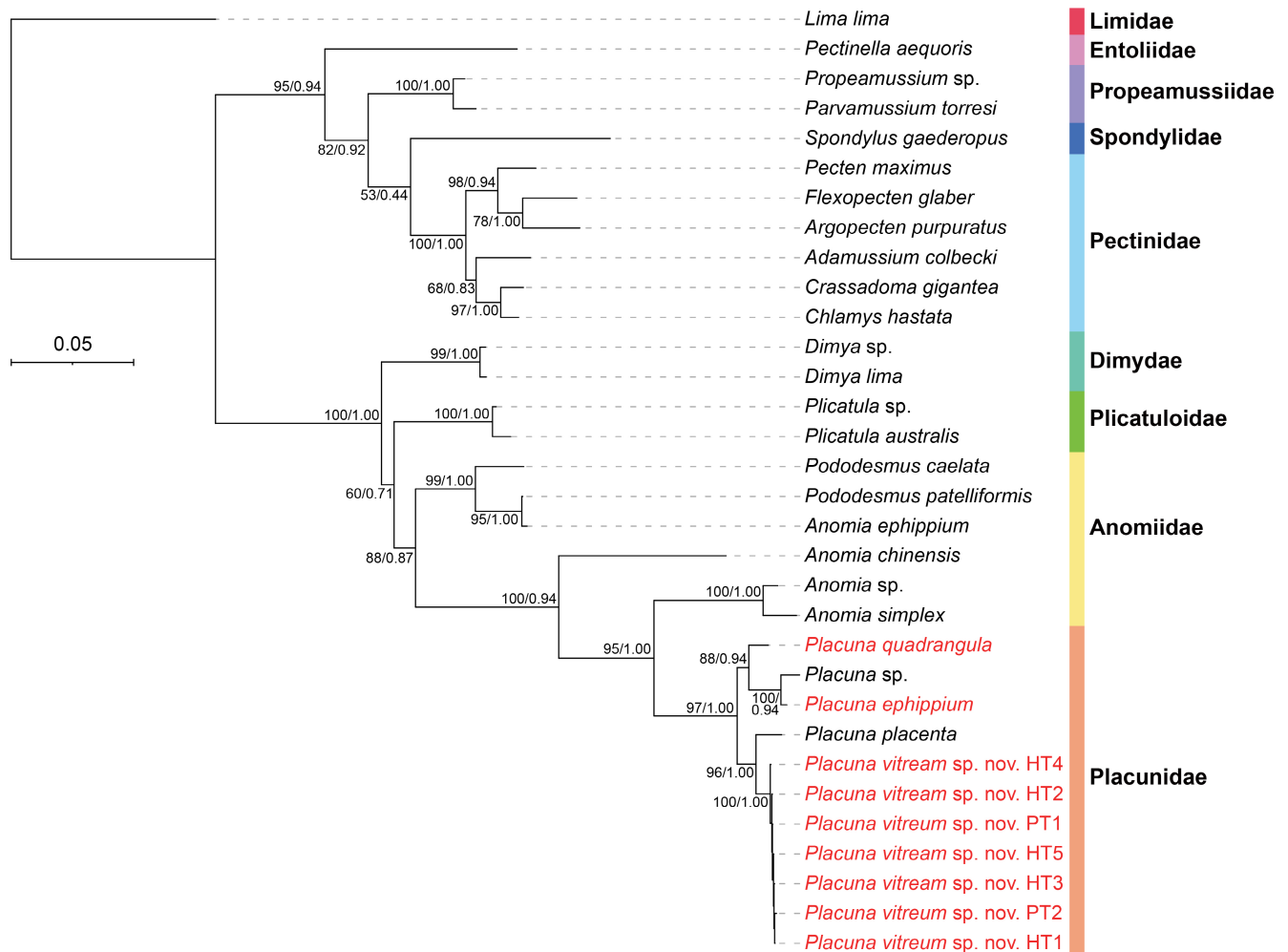


FIGURE 4 Phylogenetic relationships of Pectinida. (a) The placement of *Placuna vitream* sp. nov. in Pectinida revealed by a 4745-bp concatenated alignment (*cox1-16S rRNA-18S rRNA-28S rRNA-histone H3*) using Bayesian inference (BI) analysis. Bootstrap values from maximum-likelihood analysis and posterior probabilities values from BI analysis are given at nodes. Sequences generated from this study are highlighted in red color. The sequences *P. vitream* sp. nov. HT1-5 represent the specimens TMBC031019-TMBC031023 collected from the type locality, while the sequences *P. vitream* sp. nov. PT1-2 represent the specimens collected from Haikou and Xiamen, respectively.

similar to the windowpane shell *P. placenta*, which may explain its records as a common species in Chinese coastal waters (Li et al., 2019; Liu, 2008). They are extremely similar in shell shape, outline, hinge ridge and ligament form, and autonomy (Yonge, 1977). Considering the widely distributed so-called “*P. placenta*” and the taxonomic uncertainty of the specimens from other locations, we only compared *P. vitream* sp. nov. with the holotype of *P. placenta* (Figure 3f) stored in the Linnean Society of London (<https://linnean-online.org/>). The *P. placenta* holotype, whose sampling locality is unknown as Carl Linnaeus only wrote “Pelago”=the Ocean for its habitat, processes a gentle dorsal outline and inconspicuous umbones (Linnaeus, 1758). Our statistical study shows that ridge angle is a significant feature for species identification within *Placuna* (Figure 3g). The ridge angle of *P. placenta* is about 21° (Linnaeus, 1758), which is slightly smaller than 28° to 31° in *P. vitream* sp. nov. (Figures 2 and 3a,b, Table 2). In conclusion, *P. vitream* sp. nov. can be distinguished from its congeneric species and regarded as a new species.

3.2.7 | Phylogenetic and genetic distance analyses

Sequencing the target gene fragments of *P. vitream* sp. nov., *P. ephippium*, and *P. quadrangula* produced 657bp *cox1*, 384–388bp 16S *rRNA*, 1765bp 18S *rRNA*, 2098–2102bp 28S *rRNA*, and 328bp *histone H3*. After alignment, trimming, and concatenation of these five fragments with 22 other Pectinida species and an outgroup Limida, a 4745bp matrix was generated. The BI and ML trees are identical in topology (Figure 4). The *Placuna* species form a single clade of Placunidae with new data from three species: *P. quadrangula*, *P. ephippium*, and the new species. Among them, the *P. vitream* sp. nov. specimens are fully supported to be sister to the specimen from Singapore identified as *P. placenta* (posterior probability=1.0, bootstrap value=100). Among the eight families of Pectinida included in the analyses, only Anomiidae is paraphyletic, with *Anomia simplex* and *Anomia* sp. being sister to Placunidae, and together they are sister to *A. chinensis* and a clade comprising *A. ephippium*, *Pododesmus caelata*, and *P. patelliformis*.

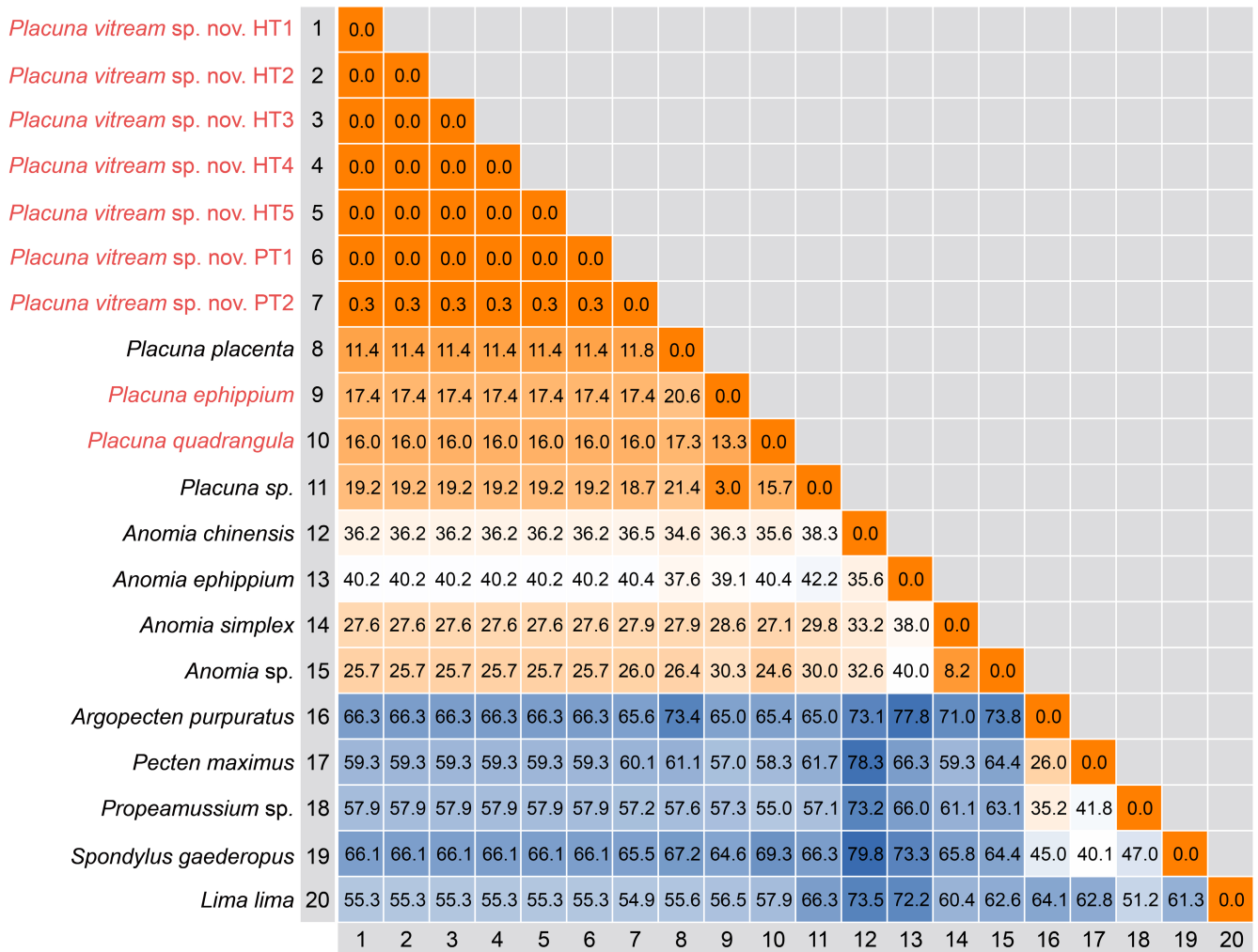


FIGURE 5 The Kimura 2-parameter (K2P) genetic distances (%) based on the *cox1* fragments among Pectinida. The analysis was performed based on a 615-bp matrix, with the species *Lima lima* as the outgroup. The sequences *P. vitream* sp. nov. HT1-5 represent the specimens TMBC031019-TMBC031023 collected from the type locality, while the sequences *P. vitream* sp. nov. PT1-2 represent the specimens collected from Haikou and Xiamen, respectively.

The intraspecific K2P genetic distances within *P. vitream* sp. nov. are 0%–0.31% for *cox1* (Figure 5) and 0% for the other four gene fragments. By contrast, the K2P distances between *P. vitream* sp. nov. and other congeneric species are much larger, ranging from 11.43% to 19.17%. Even for the sister species *P. placenta*, their divergences were 11.43%–11.82% for *cox1*, 0.38% for 16S *rRNA*, and 0.91% for 18S *rRNA* (Figure 5). However, 28S *rRNA* and *histone H3* were highly conserved with genetic distances of 0.17% and 0% between *P. vitream* sp. nov. and *P. placenta*, respectively. The large interspecific genetic distances of *cox1* between *P. vitream* sp. nov. and other *Placuna* species support our recognition of the new species.

3.3 | *Placuna aestuaria* sp. nov

<https://zoobank.org/NomenclaturalActs/4FBABB26-BFE7-41DF-8317-809F8E621BBA>

3.3.1 | Type materials and locality

Holotype (TMBC031038), paratypes 1–19 (TMBC031039–031057), collected in the type locality Mai Po Nature Reserve (22°29.03' N, 114°01.56' E), Hong Kong SAR, China by Mr. Yi-Tao Lin and Dr. Carmen K. M. Or, in July 2023.

3.3.2 | Distribution

Currently known only from the Mai Po Nature Reserve, Hong Kong SAR, China.

3.3.3 | Etymology

The species epithet “*aestuaria*” comes from “estuarial” in Latin, which refers to the estuarine waters of the type locality of this species.

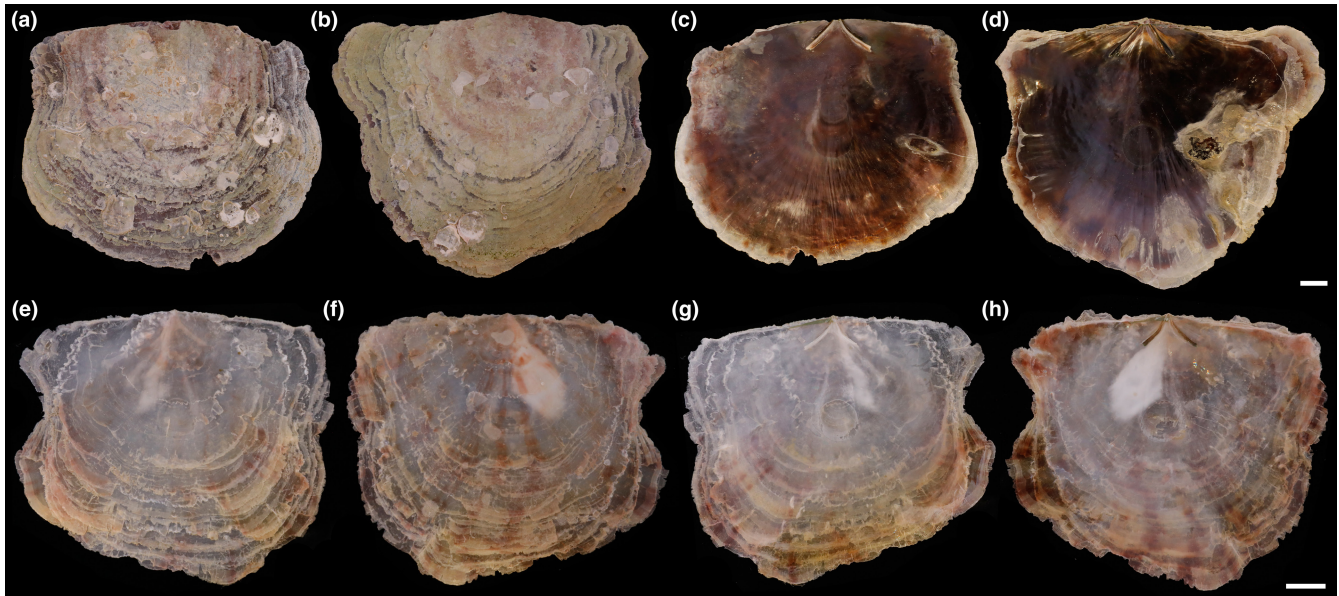


FIGURE 6 External and internal views of the left and right valves from two species of *Placuna*. (a–d) External view of left valve, external view of right valve, internal view of left valve, internal view of right valve of *P. ephippium* (TMBC031059), respectively; (e–h) External view of left valve, external view of right valve, internal view of left valve, internal view of right valve of *P. quadrangula* (TMBC031064), respectively. Scale bar: 10mm.

3.3.4 | Diagnosis

Placuna with a large shell up to 122 mm, oval to subcircular, and semi-transparent to opaque. Hinge broad with clear hinge teeth. Umbones anteriorly close to the hinge center. Ridges angle 23° to 26°. Anterior hinge ridge and ligament shorter than posterior. Auricle broad and large, and anterior auricle approximately equal to or slightly larger than posterior auricle.

3.3.5 | Description

Shell (Figures 3d,e and 7, Tables 3 and 4) oval to subcircular, opaque, and very compressed. Left valve more convex than right valve. External surface with growth lines and gray lamella, without radial lines. Length up to 122mm, nearly equal to the height ($L/H=0.909-1.016$) and approximate equilateral ($AL/L=0.490-0.520$). Hinge broad ($HH/HL=0.208-0.297$), slightly rounded, with clear hinge teeth. Umbones anteriorly close to middle of hinge ($AHL/HL=0.297-0.419$). Hinge ridges and ligaments nearly straight, inverted V-shaped with moderate ridge angle (RL) from 23° to 26°. Anterior hinge ridge and ligament shorter than posterior ridge ($ARL/PRL=0.623-0.829$). Auricles broad and large, anterior auricle approximately equal or slightly larger than posterior auricle. Anterior pedal retractor scar oval, located between the distal ends of two ridges, slightly close to anterior ridge. Adductor muscle scar subcircular, close to valve center.

3.3.6 | Remarks

The unequal hinge ridges and ligaments allow *P. aestuaria* sp. nov. to be distinguished from *P. ephippium*, *P. quadrangula*, *P. lincolni*, and

P. lobata with equal hinge ridges and ligaments, and the ratio AHL/HL of *P. aestuaria* sp. nov. is much smaller than that of *P. ephippium* and *P. quadrangula* (Das et al., 2019; Dunker, 1879; Matsukuma, 1987) (Figures 3d,e, 6, and 7, Tables 3 and 4). The ridge angle of *P. aestuaria* sp. nov. (23°–26°) is slightly larger than that of *P. placenta* (21°), and much smaller than that (60°) of the fossil species *P. pseudoplacenta* and *P. mandirantjanensis* (Martin, 1909). Besides, our statistical study between *P. aestuaria* sp. nov. and *P. vitream* sp. nov. clearly shows that the morphological indices, including HH/HL and AHL/HL , and ridge angle are significantly different (Figure 3g). Furthermore, the hinge and auricles of *P. aestuaria* sp. nov. are substantially distinct and broad with nearly straight hinge ridges (both anterior and posterior), which could not be observed in the congeneric species (Figures 6d,e and 7). These distinct characteristics support our description of *P. aestuaria* sp. nov.

4 | DISCUSSION

The identification of *Placuna* species is challenging due to the variability in shell size, shape, outline, and even the inner surface color, which varies from light purple to dark brown. *Placuna* contains many synonyms, such as *Sellaria* Link, 1807 and *P. planicostata* Dunker, 1879. A previous morphological study posited the hinge as a more appropriate shell section for species identification within this genus, whereas ridge length was also used to separate two groups (or subgenera) of *Placuna* (i.e., *Ephippium* and *Placenta*, currently unaccepted) (Gray, 1849). Besides, the wide geographical ranges of *P. placenta*, *P. ephippium*, and *P. quadrangula* make it difficult to confirm their identification by their geographic locality (Hung & Carson, 2014; Retzius, 1788; Rustia et al., 2023). In this study, we successfully applied a combination of hinge morphology (including

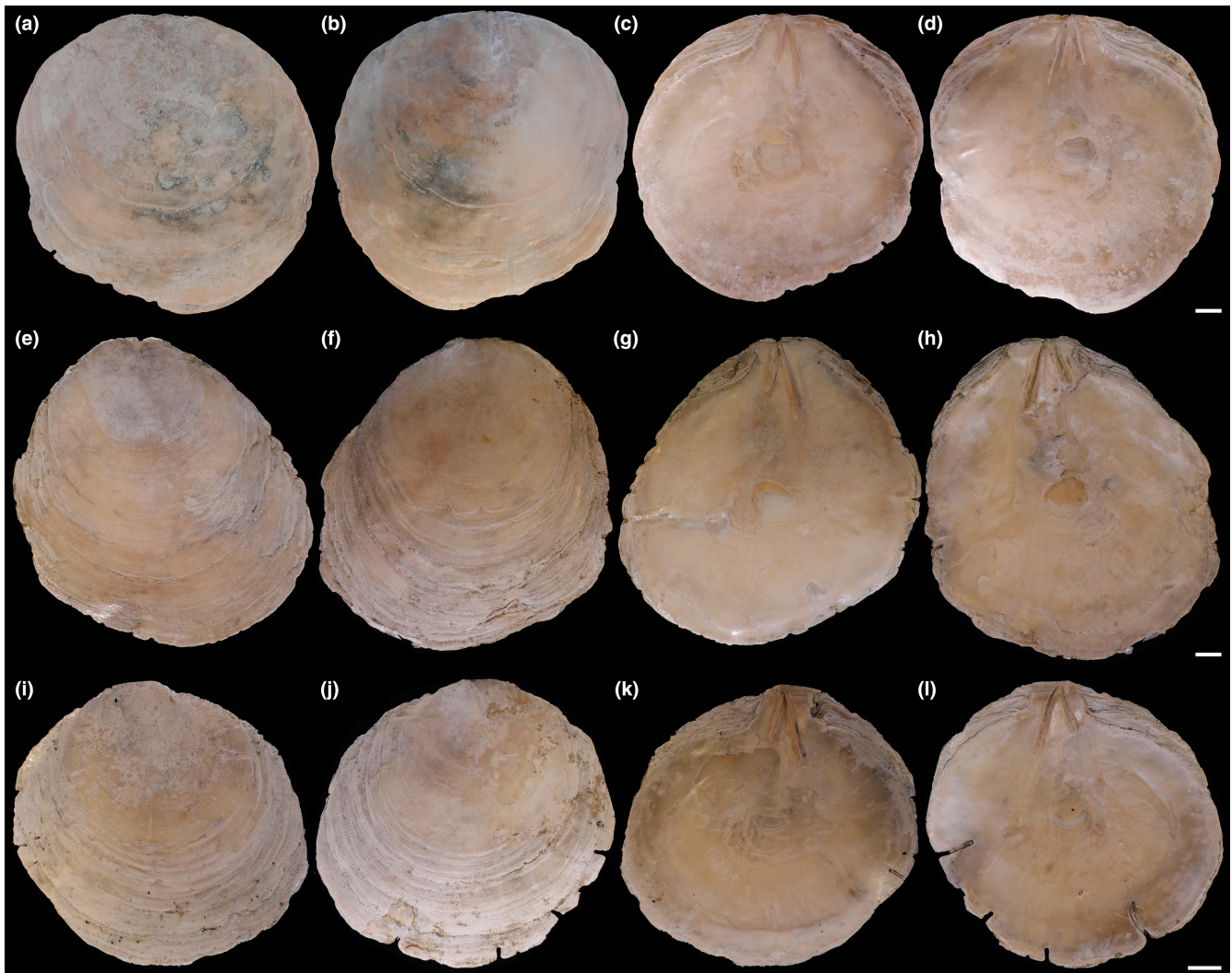


FIGURE 7 External and internal views of the left and right valves from three pairs of *Placuna aestuaria* sp. nov. specimens. (a–d) External view of left valve, external view of right valve, internal view of left valve, internal view of right valve of the holotype (TMBC031038), respectively; (e–h) External view of left valve, external view of right valve, internal view of left valve, internal view of right valve of the paratype 1 (TMBC031039), respectively; (i–l) External view of left valve, external view of right valve, internal view of left valve, internal view of right valve of the paratype 2 (TMBC031040), respectively. Scale bar: 10mm.

the AHL/HL and HH/HL ratios), ridge angle, and auricle features to distinguish the species of *Placuna* (Figure 3g).

Molecular data are also very limited in Placunidae. Prior to this study, only five gene fragments from *P. placenta* and a *cox1* sequence of an undetermined *Placuna* species have been published (Bieler et al., 2014; Chang et al., 2020; Sharma et al., 2013). The lack of molecular data makes it difficult to determine the phylogenetic relationships among *Placuna* spp. and the position of *Placuna* in Pectinida. The two mitochondrial and three nuclear gene fragments for *P. ephippium*, *P. quadrangula*, and *P. vitream* sp. nov. generated in this study will be useful for future phylogenetic studies of *Placuna* and even Bivalvia. Furthermore, given that DNA-based analyses have been widely used to distinguish morphologically similar species, such as species in Mytilida, Pteriida, and Venerida (Lemer et al., 2014; Ni et al., 2012; Shen et al., 2014), our discovery of *P. vitream* sp. nov. suggests that *Placuna* might be more diverse than previously

thought, and “*P. placenta*” specimens from other locations should be examined to determine their real identities (Gallardo et al., 1995; Li et al., 2019; Rustia et al., 2023; Song et al., 2022).

Key to *Placuna* Lightfoot, 1786:

- 1a. Hinge ridges equal in length.....2.
- 1b. Anterior ridge shorter than posterior ridge.....4.
- 2a. Shell saddle or subquadrate shaped.....3.
- 2b. Shell oval to subcircular.....*P. lincolni*.
- 3a. Shell flexuous with purple or brown color inter surface.....*P. ephippium*.
- 3b. Outer surface ornamented with ribs.....*P. lobata*.
- 3c. Shell semitransparent with radial reddish-brown stripes.....*P. quadrangula*.
- 4a. Ridges gently diverged with an angle less than 40°.....5.
- 4b. Anterior ridge strongly curved and ridge angle 60° or larger.....7.

5a. Hinge and auricles obvious.....6.

5b. The ridge angle smaller than 25°.....*P. placenta*.

6a. The ridge angle larger than 25°.....*P. vitream* sp. nov.

6b. Hinge broad with nearly straight ridges.....*P. aestuaria*.

7a. Shell outer surface with feather-like, sharply prominent radial sculpture.....*P. mandirantjanensis*.

7b. Shell outer surface dashed by fine and radial ribs.....*P. pseudoplacenta*.

5 | CONCLUSION

We reported two new species of *Placuna*, *P. vitream* sp. nov., and *P. aestuaria*, and sequenced two mitochondrial and three nuclear genes for *P. ephippium*, *P. quadrangula*, and *P. vitream* sp. nov. Our K2P genetic distance analyses showed that the three specimens from different locations in China are *P. vitream* sp. nov. The genetic distance and phylogenetic analyses confirmed the distinction between *P. vitream* sp. nov. and its sister species from Singapore, which was identified as *P. placenta*, as well as other congeneric species with corresponding DNA sequences. Besides, we described the shell morphological features of *P. vitream* sp. nov. and *P. aestuaria*, and the internal anatomy of *P. vitream* sp. nov. and *P. aestuaria* can be distinguished from congeneric species by their hinge structures, ridge angles, and auricle features. Our discovery of *P. vitream* sp. nov. and *P. aestuaria* enhances our understanding of the diversity of *Placuna*. This case study also highlights the urgency of reevaluation of the identity and distribution of marine species in Asian waters, especially those described in the 1970s to early 1900s and considered to have wide distribution ranges.

AUTHOR CONTRIBUTIONS

Yi-Tao Lin: Data curation (lead); methodology (lead); visualization (lead); writing – original draft (lead). **Yi-Xuan Li:** Methodology (equal); visualization (equal). **Hai-Xin Loke:** Data curation (equal). **Xiao Han:** Data curation (equal). **Jian-Wen Qiu:** Conceptualization (lead); funding acquisition (lead); project administration (lead); supervision (lead).

ACKNOWLEDGMENTS

This project was supported by the Lantau Conservation Fund (RE-2020-22) and the Hong Kong Offshore LNG Terminal Project (MCEF22003). We thank Dr. Carmen K. M. Or (WWF Hong Kong), Dr. Yanjie Zhang (Hainan University), Mr. Juhao Wang, and Mr. Junhao Pan for their assistance in sample collection.

CONFLICT OF INTEREST STATEMENT

The authors declare that they have no competing interests.

DATA AVAILABILITY STATEMENT

The genetic sequences generated in this study are available in GenBank (<https://www.ncbi.nlm.nih.gov/genbank/>) under the accession numbers shown in Table 2.

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How to cite this article: Lin, Y.-T., Li, Y.-X., Loke, H.-X., Han, X., & Qiu, J.-W. (2024). One becomes three: An integrative morphological and molecular analysis of the windowpane oyster *Placuna* (Bivalvia: Pectinida) reveals new species. *Ecology and Evolution*, 14, e70260. <https://doi.org/10.1002/ece3.70260>