

SUPPLEMENTARY INFORMATION

Discovery of *Novaculina myanmarensis* **sp. nov.** (Bivalvia: Pharidae: Pharellinae) closes the freshwater razor clams range disjunction in Southeast Asia

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Habitat of *Novaculina myanmarensis* **sp. nov.**, Ayeyarwady River, near Minbu, 20.1911° N, 94.8788° E (Photo: Nyein Chan).

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Supplementary Table 1. List of additional sequences from GenBank used in this study for phylogenetic reconstructions. Species names and environmental preference are on the basis of the World Register of Marine Species database¹.

Taxa	Environment	Locality	COI	16S rRNA	28S rRNA
<i>Sinonovacula constricta</i> (Lam., 1818)	Marine	China	JN859986	JN859901	AF131005
<i>S. mollis</i> (G. B. Sowerby II, 1874) (= <i>S. rivularis</i> Huang & Zhang, 2007)	Marine	China	JN859994	JN859908	n/a
<i>Pharella javanica</i> (Lam., 1818)	Marine	Thailand	n/a	n/a	AM779683
<i>Cultellus attenuatus</i> Dunker, 1862	Marine	China	JN859998	JN859913	n/a
<i>Ensiculus cultellus</i> (L., 1758)	Marine	Queensland, Australia	n/a	n/a	AM779682
<i>Pharus legumen</i> (L., 1758)	Marine	South Wales, UK	n/a	n/a	AM779684
<i>Phaxas pellucidus</i> (Pennant, 1777)	Marine	Sweden, Tjörnö	KC429145	KC429309	KC429508
<i>Ensis magnus</i> Schumacher, 1817*	Marine	Ireland & Spain**	EU523690	AJ586445	n/a
<i>E. leei</i> Huber, 2015*	Marine	Europe & USA**	EU523673	HF970450	JF909603
<i>E. macha</i> (Molina, 1782)*	Marine	Chile**	JF301780	HF970471	n/a
<i>Siliqua alta</i> (Broderip & Sowerby, 1829)*	Marine	Hokkaido, Japan	n/a	AB751362	AB746908
<i>S. radiata</i> (Linnaeus, 1758)*	Marine	China	JN860004	JN859917	n/a

*Outgroup taxa. **Chimeric sequence. n/a – not available.

Supplementary Table 2. List of available georeferenced records of *Novaculina* species

Species	Locality	Latitude	Longitude	Reference
<i>N. myanmarensis</i> sp. nov.	Myanmar: downstream of Donthami River	16.6935	97.5819	This study (RMBH)
<i>N. myanmarensis</i> sp. nov.	Myanmar: Ayeyarwady River, Pakokku Region, near Thin Baw Kone village*	21.3146	95.0591	This study (RMBH)
<i>N. myanmarensis</i> sp. nov.	Myanmar: Ayeyarwady River, large sandbar 0.5 mi SE of Nyaung-U	21.2066	94.9062	This study (CAS)
<i>N. myanmarensis</i> sp. nov.	Myanmar: Ayeyarwady River, near Minbu	20.1911	94.8788	This study (FFI)
<i>N. gangetica</i>	India: Ganges River at Vindhyachal upstream of Mirzapur	25.1666	82.5072	Ref. 5
<i>N. gangetica</i>	India: Hooghly River downstream of Howra	22.4278	88.1368	Ref. 5
<i>N. gangetica</i>	India: Ganges River at Varanasi	25.3279	83.0464	Ref. 6
<i>N. gangetica</i>	India: Ganges River at Patna	25.6589	85.1573	Ref. 7
<i>N. gangetica</i>	India: Calcutta*	22.6000	88.3000	MCZ 224805; UMZC I.102125
<i>N. gangetica</i>	India: Jumna [Yamuna] River at Hamirpur	25.9617	80.1525	Ref. 8
<i>N. gangetica</i>	India: Gomti River at Jaunpur	25.7474	82.6874	Ref. 8
<i>N. gangetica</i>	India: Burhi Gandak River at Khagaria	25.5009	86.4776	Ref. 9
<i>N. gangetica</i>	Bangladesh: Turag River at Tangi Bridge	23.8997	90.4081	Ref. 10
<i>N. gangetica</i>	Bangladesh: Moyur River	22.7719	89.5402	Ref. 11
<i>N. gangetica</i>	Myanmar: Kaladan River	21.0094	92.9813	Ref. 12
<i>N. gangetica</i>	Myanmar: Lemro River	20.6150	93.2481	Ref. 12
<i>N. siamensis</i>	Thailand: Chantakam [Bang Pakong River basin]*	14.0000	102.0000	Ref. 2
<i>N. siamensis</i>	Thailand: Pa Sak River near Saraburi	14.5304	100.9113	Ref. 3
<i>N. siamensis</i>	Vietnam: Ba Lai River, Mekong Delta	10.2647	106.4397	Ref. 4
<i>N. chinensis</i>	China: Lake Taihu, Wuxi*	31.4402	120.3143	Ref. 13, 14
<i>N. chinensis</i>	China: Lake Taihu, southwest	31.1843	120.4267	Ref. 15
<i>N. chinensis</i>	China: Lake Hongze	33.3075	118.7100	Ref. 16
<i>N. chinensis</i>	China: Lake Chaohu	31.5224	117.5616	Ref. 17
<i>N. chinensis</i>	China: Yintan	28.0333	117.0667	Ref. 18

*Type locality (see Taxonomic Account for details).

Supplementary Table 3. Models of sequence evolution for each partition based on corrected Akaike Information Criterion (AICc) of MEGA6¹⁹ that were applied within Bayesian models

Data set and partition	Model	Gamma	Invariant
1 st codon of <i>COI</i>	TN93+G	0.10	n/a
2 nd codon of <i>COI</i>	HKY+G	0.05	n/a
3 rd codon of <i>COI</i>	HKY+G	0.87	n/a
16S <i>rRNA</i>	GTR+G+I	0.40	0.51
28S <i>rRNA</i>	GTR+G+I	7.15	0.58

n/a – not available.

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