

Supplementary Information

¹ *Department of Integrative Zoology, University of Vienna, UZA1 Althanstrasse 14, 1090 Vienna, Austria*

² *Laboratory of Cell Differentiation, A.V. Zhirmunsky Institute for Marine Biology, Far East Branch of the Russian Academy of Sciences, Paltchevsky ulica 17, 690041 Vladivostok, Russian Federation*

A putative species complex in the Sea of Japan revealed by DNA sequence data: A study on *Lottia* cf. *kogamogai* (Gastropoda: Patellogastropoda)

ALEN KRISTOF^{1*}, ANDRÉ LUIZ DE OLIVEIRA¹, KONSTANTIN G. KOLBIN² & ANDREAS WANNINGER¹

**Corresponding author: alen.kristof@univie.ac.at*

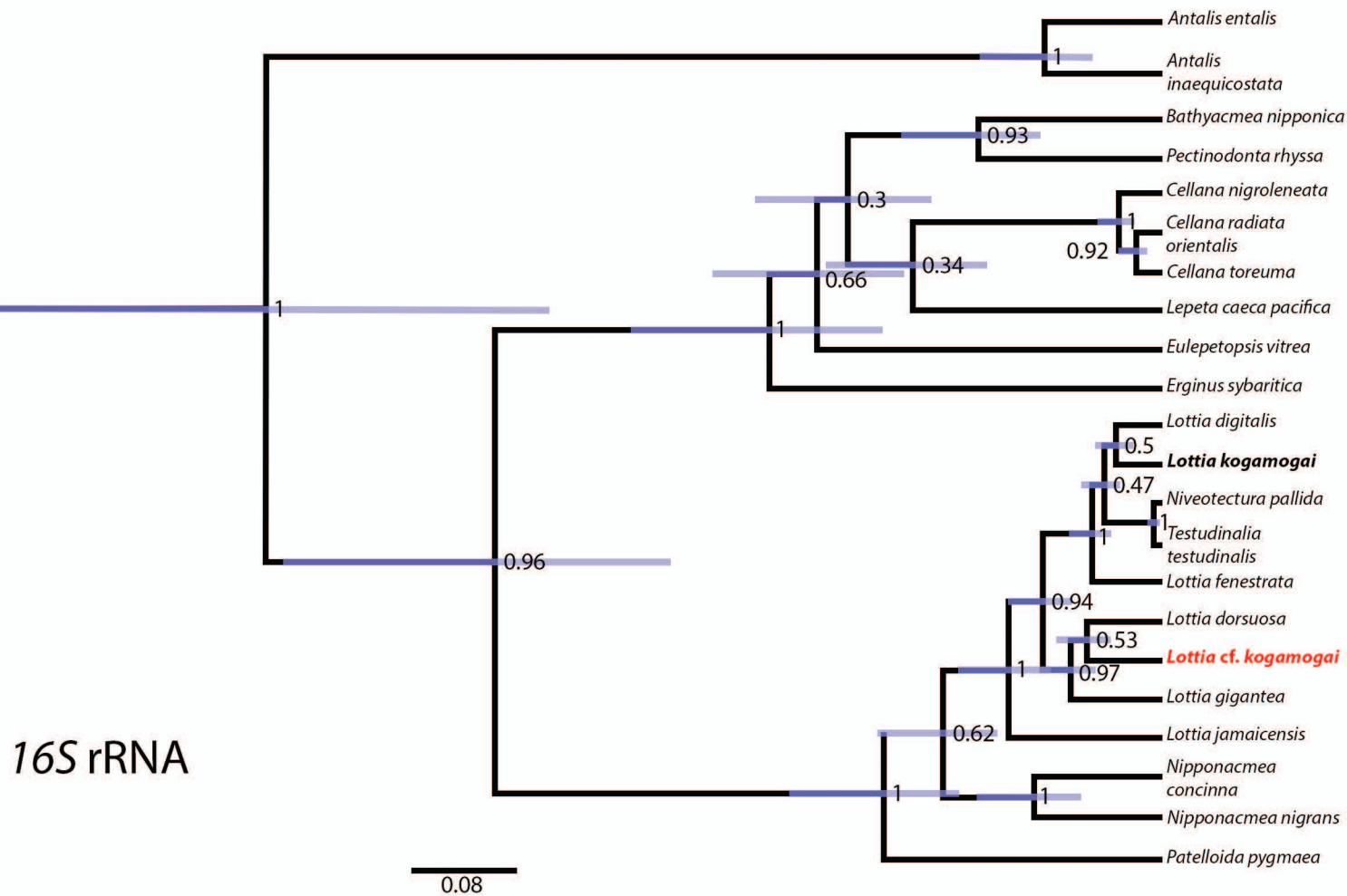
Contributing authors: andre.l Luiz.de.oliveira@univie.ac.at; konstantin.kolbin@gmail.com;

andreas.wanninger@univie.ac.at

Table S1. List of species and genes with GenBank accession numbers used for phylogenetic analyses

Clade/Family	Species	16S	CO1	18S
Patellogastropoda				
Nacellidae	<i>Cellana nigrolineata</i>	DQ093467	AB548213	DQ013353
	<i>Cellana radiata</i>	AB106478	AB433647	AB282768
	<i>orientalis</i>			
	<i>Cellana toreuma</i>	GQ455937	JQ313557	AF308646
Lepetidae	<i>Lepeta caeca pacifica</i>	AB238347	AB543978	AB282759
Pectinodontidae	<i>Bathyacmaea</i>	AB238451	AB238588	AB282772
	<i>nipponica</i>			
	<i>Pectinodonta rhyssa</i>	AB238452	AB238589	AB282773
Lottiidae	<i>Erginus sybaritica</i>	AB238350	AB238461	AB282761
	<i>Lottia digitalis</i>	AB238352	KF643845	DQ248942
	<i>Lottia dorsuosa</i>	AB106502	KM221054	AF308645
	<i>Lottia fenestrata</i>	FJ977695	FJ977749	FJ977631
	<i>Lottia gigantea</i>	FJ977696	FJ977750	FJ977632
	<i>Lottia jamaicensis</i>	FJ977697	FJ977751	FJ977633
	<i>Lottia kogamogai</i>	AB106493	AB238467	-
	<i>Lottia cf. kogamogai</i>	KU053948	KU053950	KU053949
	<i>Nipponacmea</i>	AB106511	KM221077	DQ013354
	<i>concinna</i>			
	<i>Nipponacmea</i>	AB106516	AB238490	AB282763
	<i>nigrans</i>			
	<i>Niveotectura pallida</i>	AB106519	AB238494	AF308644
<i>Testudinalia</i>	FJ977694	FJ977748	FJ977630	
<i>testudinalis</i>				
Neolepetopsidae	<i>Eulepetopsis vitrea</i>	DQ093468	DQ093516	DQ093427
Patelloidinae	<i>Patelloida pygmaea</i>	AB161514	AB196505	AB282765
Outgroup				
Scaphopoda				
Dentaliidae	<i>Antalis entalis</i>	DQ280027	DQ280016	DQ279936
	<i>Antalis</i>	DQ280026	DQ280015	DQ279935
	<i>inaequicostata</i>			

A



B

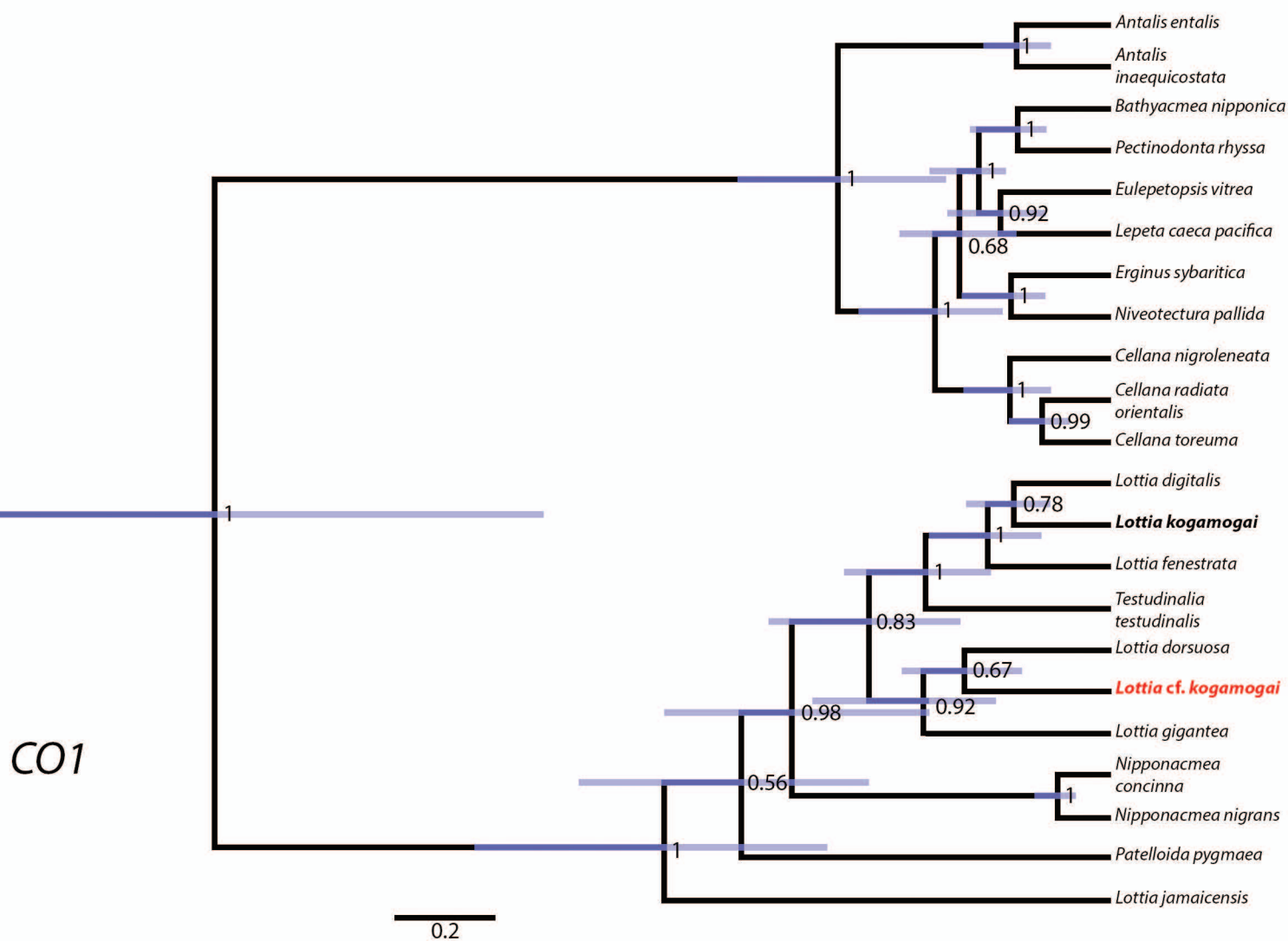


Figure S1. Maximum clade credibility (MCC) trees using individual *16S* rRNA and *cytochrome c oxidase* subunit I (*CO1*) mitochondrial markers resulting from BEAST2 Bayesian analysis. *Lottia cf. kogamogai* is indicated in red and *Lottia kogamogai* Sasaki and Okutani, 1994 in bold letters. All clades are shown with their associated 95% confidence intervals (blue bars). The support values are posterior probabilities. Outgroup: *Antalis entalis* (Linnaeus, 1758), *Antalis inaequicostata* (Dautzenberg, 1891). **A.** MCC tree based on the *16S* gene. **B.** MCC tree based on the *CO1* gene.

Table S2. Phenotypic distances (sequence divergence) between *16S* rRNA and *cytochrome c oxidase* subunit 1 (*CO1*). Proportion of base substitutions per site between *16S* (above the diagonal) and *CO1* (below the diagonal) sequences are shown. The analysis involved 22 nucleotide sequences. All positions containing gaps and missing data were eliminated. There are a total of 248 (*16S*) and 548 (*CO1*) positions in the final dataset. Calculation performed in MEGA 6 [1].

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22
<i>Antalis entalis</i> (1)	-	0.150	0.315	0.335	0.339	0.347	0.351	0.355	0.331	0.399	0.403	0.399	0.379	0.411	0.403	0.403	0.379	0.379	0.403	0.367	0.335	0.395
<i>Antalis inaequicostata</i> (2)	0.171	-	0.302	0.306	0.290	0.294	0.327	0.355	0.306	0.395	0.411	0.403	0.387	0.419	0.391	0.399	0.383	0.379	0.403	0.351	0.310	0.395
<i>Bathycypraea nipponica</i> (3)	0.262	0.297	-	0.262	0.25	0.254	0.254	0.234	0.246	0.355	0.347	0.355	0.343	0.391	0.351	0.347	0.355	0.331	0.347	0.323	0.218	0.347
<i>Cellana nigrolineata</i> (4)	0.297	0.301	0.234	-	0.0726	0.077	0.222	0.234	0.214	0.319	0.315	0.310	0.315	0.339	0.331	0.323	0.335	0.323	0.327	0.315	0.198	0.323
<i>Cellana radiata orientalis</i> (5)	0.294	0.301	0.234	0.126	-	0.052	0.214	0.230	0.206	0.302	0.306	0.302	0.290	0.319	0.319	0.302	0.315	0.306	0.315	0.302	0.169	0.306
<i>Cellana toreuma</i> (6)	0.308	0.332	0.28	0.178	0.157	-	0.222	0.226	0.222	0.302	0.306	0.306	0.302	0.335	0.331	0.306	0.319	0.310	0.315	0.290	0.198	0.302
<i>Erginus sybaritica</i> (7)	0.262	0.276	0.185	0.161	0.175	0.224	-	0.242	0.226	0.290	0.282	0.282	0.298	0.319	0.302	0.286	0.298	0.286	0.302	0.310	0.242	0.302
<i>Eulepetopsis vitrea</i> (8)	0.287	0.266	0.206	0.22	0.238	0.231	0.192	-	0.262	0.282	0.274	0.298	0.258	0.282	0.278	0.274	0.286	0.266	0.302	0.290	0.266	0.298
<i>Lepeta caeca pacifica</i> (9)	0.273	0.294	0.192	0.21	0.224	0.224	0.185	0.15	-	0.343	0.331	0.335	0.343	0.363	0.339	0.339	0.310	0.327	0.359	0.315	0.206	0.351
<i>Lottia digitalis</i> (10)	0.413	0.381	0.43	0.381	0.413	0.427	0.406	0.381	0.395	-	0.097	0.065	0.109	0.149	0.093	0.028	0.173	0.153	0.073	0.198	0.347	0.069
<i>Lottia dorsuosa</i> (11)	0.381	0.385	0.413	0.374	0.402	0.392	0.374	0.385	0.371	0.224	-	0.121	0.101	0.153	0.089	0.081	0.173	0.157	0.121	0.206	0.335	0.113
<i>Lottia fenestrata</i> (12)	0.406	0.392	0.434	0.406	0.399	0.413	0.406	0.381	0.392	0.213	0.287	-	0.121	0.149	0.109	0.056	0.165	0.145	0.089	0.206	0.335	0.089
<i>Lottia gigantea</i> (13)	0.402	0.399	0.406	0.374	0.406	0.406	0.381	0.406	0.395	0.238	0.203	0.28	-	0.145	0.105	0.101	0.157	0.157	0.117	0.198	0.323	0.105
<i>Lottia jamaicensis</i> (14)	0.374	0.402	0.451	0.399	0.402	0.402	0.423	0.409	0.413	0.332	0.325	0.315	0.315	-	0.133	0.145	0.181	0.198	0.161	0.218	0.371	0.153
<i>Lottia cf. kogamogai</i> (15)	0.364	0.378	0.395	0.378	0.409	0.399	0.388	0.364	0.388	0.276	0.217	0.301	0.213	0.329	-	0.085	0.173	0.161	0.109	0.198	0.339	0.117
<i>Lottia kogamogai</i> (16)	0.43	0.441	0.451	0.434	0.434	0.444	0.434	0.441	0.43	0.199	0.276	0.227	0.28	0.322	0.311	-	0.169	0.149	0.077	0.202	0.343	0.073
<i>Nipponacmea concinna</i> (17)	0.381	0.399	0.385	0.381	0.409	0.43	0.392	0.399	0.402	0.283	0.245	0.318	0.22	0.357	0.22	0.301	-	0.141	0.173	0.222	0.323	0.165
<i>Nipponacmea nigrans</i> (18)	0.399	0.406	0.409	0.388	0.399	0.434	0.402	0.416	0.392	0.315	0.259	0.294	0.227	0.343	0.245	0.308	0.112	-	0.169	0.210	0.323	0.165
<i>Niveotectura pallida</i> (19)	0.294	0.311	0.227	0.203	0.224	0.203	0.154	0.21	0.22	0.409	0.364	0.434	0.374	0.399	0.367	0.437	0.413	0.409	-	0.202	0.339	0.02
<i>Patelloida pygmaea</i> (20)	0.395	0.402	0.399	0.406	0.409	0.416	0.395	0.406	0.402	0.332	0.308	0.336	0.28	0.325	0.304	0.35	0.325	0.318	0.406	-	0.319	0.19
<i>Pectinodonta rhyssa</i> (21)	0.276	0.283	0.168	0.231	0.217	0.255	0.196	0.206	0.175	0.406	0.378	0.399	0.392	0.423	0.392	0.434	0.378	0.381	0.213	0.392	-	0.335
<i>Testudinalia testudinalis</i> (22)	0.395	0.437	0.385	0.395	0.402	0.378	0.42	0.395	0.371	0.21	0.262	0.252	0.241	0.318	0.245	0.241	0.28	0.297	0.42	0.308	0.395	-

16S

CO1

[1] Tamura K, Stecher G, Peterson D, Filipski A, Kumar S (2013) MEGA6: Molecular evolutionary genetics analysis version 6.0. *Mol Biol Evol* **30**: 2725-2729.

Table S3. Estimates of evolutionary divergence using the Jukes-Cantor model [1] between *16S* rRNA and *cytochrome c oxidase* subunit 1 (*CO1*). The number of base substitutions per site between *16S* (above the diagonal) and *CO1* (below the diagonal) sequences are shown. The analysis involved 22 nucleotide sequences. All positions containing gaps and missing data were eliminated. There are a total of 248 (*16S*) and 548 (*CO1*) positions in the final dataset. Evolutionary analyses were conducted in MEGA6 [2].

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22
<i>Antalis entalis</i> (1)	-	0.166	0.408	0.443	0.451	0.465	0.473	0.481	0.436	0.57	0.579	0.57	0.528	0.596	0.579	0.579	0.528	0.528	0.579	0.504	0.443	0.561
<i>Antalis inaequicostata</i> (2)	0.195	-	0.387	0.394	0.367	0.374	0.429	0.481	0.394	0.561	0.596	0.579	0.544	0.614	0.553	0.57	0.536	0.528	0.579	0.473	0.401	0.561
<i>Bathyacmaea nipponica</i> (3)	0.323	0.378	-	0.322	0.304	0.31	0.31	0.28	0.298	0.481	0.465	0.481	0.458	0.553	0.473	0.465	0.481	0.436	0.465	0.422	0.257	0.465
<i>Cellana nigrolineata</i> (4)	0.378	0.384	0.281	-	0.076	0.081	0.263	0.28	0.252	0.415	0.408	0.401	0.408	0.451	0.436	0.422	0.443	0.422	0.429	0.408	0.229	0.422
<i>Cellana radiata orientalis</i> (5)	0.373	0.384	0.281	0.138	-	0.054	0.252	0.274	0.24	0.387	0.394	0.387	0.367	0.415	0.415	0.387	0.408	0.394	0.408	0.387	0.192	0.394
<i>Cellana toreuma</i> (6)	0.396	0.439	0.35	0.204	0.177	-	0.263	0.269	0.263	0.387	0.394	0.394	0.387	0.443	0.436	0.394	0.415	0.401	0.408	0.367	0.229	0.387
<i>Erginus sybaritica</i> (7)	0.323	0.345	0.213	0.181	0.199	0.266	-	0.292	0.269	0.367	0.354	0.354	0.38	0.415	0.387	0.361	0.38	0.361	0.387	0.401	0.292	0.387
<i>Eulepetopsis vitrea</i> (8)	0.361	0.328	0.241	0.261	0.286	0.276	0.222	-	0.322	0.354	0.341	0.38	0.316	0.354	0.348	0.341	0.361	0.329	0.387	0.367	0.329	0.38
<i>Lepeta caeca pacifica</i> (9)	0.339	0.373	0.222	0.246	0.266	0.266	0.213	0.168	-	0.458	0.436	0.443	0.458	0.496	0.451	0.451	0.401	0.429	0.488	0.408	0.24	0.473
<i>Lottia digitalis</i> (10)	0.599	0.532	0.639	0.532	0.599	0.631	0.584	0.532	0.561	-	0.104	0.067	0.118	0.166	0.099	0.029	0.197	0.171	0.076	0.229	0.465	0.072
<i>Lottia dorsuosa</i> (11)	0.532	0.539	0.599	0.518	0.576	0.554	0.518	0.539	0.511	0.266	-	0.132	0.108	0.171	0.094	0.085	0.197	0.176	0.132	0.24	0.443	0.122
<i>Lottia fenestrata</i> (12)	0.584	0.554	0.647	0.567	0.569	0.599	0.584	0.532	0.554	0.251	0.361	-	0.132	0.166	0.118	0.059	0.187	0.161	0.094	0.24	0.443	0.094
<i>Lottia gigantea</i> (13)	0.576	0.569	0.584	0.518	0.584	0.584	0.532	0.584	0.561	0.286	0.236	0.35	-	0.161	0.113	0.108	0.176	0.176	0.127	0.229	0.422	0.113
<i>Lottia jamaicensis</i> (14)	0.518	0.576	0.69	0.569	0.576	0.576	0.623	0.591	0.599	0.439	0.426	0.408	0.408	-	0.146	0.161	0.208	0.229	0.182	0.257	0.512	0.171
<i>Lottia cf. kogamogai</i> (15)	0.497	0.525	0.561	0.525	0.591	0.569	0.547	0.497	0.547	0.345	0.256	0.384	0.251	0.432	-	0.09	0.197	0.182	0.118	0.229	0.451	0.127
<i>Lottia kogamogai</i> (16)	0.639	0.664	0.69	0.647	0.647	0.673	0.647	0.664	0.639	0.232	0.345	0.271	0.35	0.42	0.402	-	0.192	0.166	0.081	0.235	0.458	0.076
<i>Nipponacmea concinna</i> (17)	0.532	0.569	0.539	0.532	0.591	0.639	0.554	0.569	0.576	0.356	0.296	0.414	0.261	0.484	0.261	0.384	-	0.156	0.197	0.263	0.422	0.187
<i>Nipponacmea nigrans</i> (18)	0.569	0.584	0.591	0.547	0.569	0.647	0.576	0.607	0.554	0.408	0.317	0.373	0.271	0.458	0.296	0.396	0.121	-	0.192	0.246	0.422	0.187
<i>Niveotectura pallida</i> (19)	0.373	0.402	0.271	0.236	0.266	0.236	0.172	0.246	0.261	0.591	0.497	0.647	0.518	0.569	0.504	0.656	0.599	0.591	-	0.235	0.451	0.02
<i>Patelloida pygmaea</i> (20)	0.561	0.576	0.569	0.584	0.591	0.607	0.561	0.584	0.576	0.439	0.396	0.445	0.35	0.426	0.39	0.471	0.426	0.414	0.584	-	0.415	0.218
<i>Pectinodonta rhyssa</i> (21)	0.345	0.356	0.19	0.276	0.256	0.312	0.227	0.241	0.199	0.584	0.525	0.569	0.554	0.623	0.554	0.647	0.525	0.532	0.251	0.554	-	0.443
<i>Testudinalia testudinalis</i> (22)	0.561	0.656	0.539	0.561	0.576	0.525	0.615	0.561	0.511	0.246	0.323	0.307	0.291	0.414	0.296	0.291	0.35	0.378	0.615	0.396	0.561	-

16S

CO1

[1] Jukes TH, Cantor CR (1969) Evolution of protein molecules. In: Munro HN (ed), Mammalian Protein Metabolism. Academic Press, New York, pp 21-132.

[2] Tamura K, Stecher G, Peterson D, Filipski A, Kumar S (2013) MEGA6: Molecular evolutionary genetics analysis version 6.0. Mol Biol Evol **30**: 2725-2729.