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Torrenticolid water mites (Acari: Hydrachnidia: Torrenticolidae) from Malaysian Borneo

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Abstract

New records of water mites of the family Torrenticolidae (Acari: Hydrachnidia) from streams in two mountain ranges in northern Borneo are presented. Aims of this study were to reconstruct phylogenetic relationships of the newly collected torrenticolids using molecular methods, and describe all new species. A fragment of the mtCOI gene was successfully PCR-amplified from 18 torrenticolid specimens and 14 new species are described: *Torrenticola* (*Torrenticola*) *bornoensis* n. sp., *T.* (*T.*) *kinabaluensis* n. sp., *T.* (*T.*) *sabahensis* n. sp., *T.* (*T.*) *neoindica* n. sp., *T.* (*T.*) *schilthuizeni* n. sp., *Neoatractides* (*Allotorrenticola*) *sundaensis* n. sp., *N.* (*Heteratractides*) *uniscutatus* n. sp., *Pseudotorrenticola* *bornoensis* n. sp., *Monatractides* (*Monatractides*) *epiales* n. sp., *M.* (*M.*) *morpheus* n. sp., *M.* (*M.*) *phantastos* n. sp., *M.* (*M.*) *phobetor* n. sp., *M.* (*M.*) *hercules* n. sp. and *M.* (*M.*) *minuta* n. sp. Additionally, the first records for Borneo are given for *Torrenticola* (*Megapalpis*) cf. *pugionirostris* (K. Viets, 1939), *Monatractides* (*Monatractides*) *longiventris* (K. Viets, 1939), *M.* (*M.*) cf. *macroporus* (K. Viets, 1935) and *M.* (*M.*) *oxystomus* (K. Viets, 1935). *Monatractides tobaensis* (K. Viets, 1935) is transferred to the subgenus *Vietsclio* Pešić & Smit, 2014. A key to the species of *Monatractides* is presented.

Key words: Acari, water mites, running waters, taxonomy, new species

lowest divergences were found between *T. neoindica* n. sp. and *T. kinabaluensis* n. sp. and between *T. neoindica* n. sp. and *T. borneoensis* n. sp., respectively. The highest divergence (22 %) between *Torrenticola* species were found between *T. sabahensis* n. sp. and *T. tjiwalensis*. These high divergence values are generally indicative of distinct species. Average pairwise distances between *Monatractides longiventris* (K. Viets, 1939) and *M. morphicus* n. sp. were found to be high (21.8 %). These high divergence values suggest that these species are not closely related, what is confirmed by many morphological differences.

TABLE 7. Average pairwise maximum likelihood distances (TN model) among mtCOI sequences between each morpho-species (lower diagonal) and within morho-species (diagonal).

	1	2	3	4	5	6	7	8
1— <i>Hygrobates hamatus</i>	—							
2— <i>Torrenticola sabahensis</i>	0.431	0.002						
3— <i>Torrenticola neoindica</i>	0.415	0.191	0.051					
4— <i>Torrenticola kinabaluensis</i>	0.428	0.209	0.134	0.018				
5— <i>Torrenticola borneoensis</i>	0.403	0.209	0.149	0.130	—			
6— <i>Torrenticola tjiwalensis</i>	0.442	0.220	0.195	0.204	0.197	—		
7— <i>Monatractides morphicus</i>	0.455	0.251	0.245	0.238	0.241	0.251	—	
8— <i>Monatractides longiventris</i>	0.518	0.304	0.274	0.265	0.278	0.272	0.218	—

To use a threshold of genetic distance for example of 2%, has been proposed and used widely to identify samples or even to delimit species, especially in the DNA barcoding literature (e.g., Hebert *et al.* 2003, Kumar *et al.* 2007). As stated by Stålstedt *et al.* (2013), the genetic distances in the barcode region in water mites were larger than the distances among the majority of recognized, closely related species of other animal groups tested to date (see Stålstedt *et al.* 2013 for references). The differentiation in our study (4–6 %) however, is lower than in other studies on closely related species in water mites (Martin *et al.* 2010 [18–31 %], Stålstedt *et al.* 2013 [11–27 %]). Recently, Pešić *et al.* (2012b) showed that two torrenticolid species, *Torrenticola lukai* Pešić *et al.* 2011 (Montenegro) and *T. lundbladi* (K. Viets, 1930) (Spain) differed by 11.6–11.8 % in mtCOI. Both species belong to a clade with shoulder platelets fused with the dorsal plate, which include four species in our molecular analysis (*neoindica*, *kinabaluensis*, *borneoensis* and *sabahensis*).

The highest divergences within morpho-taxa were those between specimens of *Torrenticola neoindica* n. sp. (5.1 %), which came from two different sampling stations. Divergences between four specimens of *Torrenticola neoindica* n. sp. from Mahua stream were about 1.1 %. However, the single specimen examined from the first stream Minduk Sirung Trail was separated from the later specimens with a genetic distance of 11 %. The latter can be indicative of the presence of an undescribed sister species.

Divergences between three examined specimens of *T. kinabaluensis* n. sp. were about 1.8 %, and three examined specimens of *T. sabahensis* n. sp. about 0.2 % (Table 7). These are all indicative of intraspecific variability.

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