

report 1997. SFD, Kuching, Sarawak, Malaysia.

Schreiber, A., Wirth, R., Riffel, M., & Van Rompaey, H. 1989. *Weasels, civets, mongooses, and their relatives: an action plan for the conservation of mustelids and viverrids*. IUCN, Gland, Switzerland.

Than Zaw, Saw Htun, Saw Htoo Tha Po, Myint Maung, Lynam, A. J., Kyaw Thinn Latt & Duckworth, J. W. 2008. Status and distribution of small carnivores in Myanmar. *Small Carnivore Conservation* 38: 2–28.

Van Rompaey, H. & Azlan J., M. 2004. Hose's Civet, *Diplogale hosei*. *Small Carnivore Conservation* 30: 18–19.

Wells, K., Bium, A. & Gabin, M. 2005. Viverrid and herpestid observations by camera and small cage trapping in the lowland rainforests on Borneo including a record of the Hose's Civet, *Diplogale hosei*. *Small Carnivore Conservation* 32: 12–14.

Wild Life Protection Ordinance 1998. *Sarawak Wild Life Protectirinance 1998*. SFD, Kuching, Sarawak, Malaysia.

Wilting, A., Samejima, H. & Mohamed, A. 2010. Diversity of Bornean viverrids and other small carnivores in Deramakot Forest Reserve, Sabah, Malaysia. *Small Carnivore Conservation* 42: **[[Divya: insert pg nos!]]**.

Yasuma, S. 2004. Observations of a live Hose's Civet *Diplogale hosei*. *Small Carnivore Conservation* 31: 3–5.

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ERRATUM

Helgen, K. M., Kays, R., Helgen, L. E., Tsuchiya-Jerep, M. T. N., Pinto, C. M., Koepfli, K. P., Eizirik, E. & Maldonado, J. E. 2009. Taxonomic boundaries and geographic distributions revealed by an integrative systematic overview of the mountain coatis, *Nasuella* (Carnivora: Procyonidae). *Small Carnivore Conservation* 41: 65–74.

*Fig. 3 (page 70) revised. Molecular relationships of coatis based on partial cytochrome b sequences. One of three most parsimonious trees (length = 167, retention index = 0.763, consistency index = 0.760) from the partial sequence of the cyt b gene (366 bp). This comparison allows for the inclusion of the short sequence generated from DNA extracted from the turbinates of a specimen of *N. meridensis*. Branch support values represent maximum parsimony and maximum likelihood bootstrap support, followed by Bayesian posterior probabilities values, respectively.*

The bootstrap support values for the maximum likelihood estimate were incorrect in the original article. Our overall taxonomic conclusions remain unchanged.

