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**Wildlife Conservation Society Malaysia Program, 7
Jalan Ridgeway, 93200 Kuching, Sarawak, Malaysia
Emails: greensmoke72@yahoo.com (John Mathai),
ngumbangjuat@yahoo.com (Ngumbang Juat)**

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.

