Phylogenetic tree of *Rattus sensu lato* estimated using Cyt B sequences from Genbank and this study (5_Rodent_cytb_full.nex). Groups labelled in boldface are clade names taken from [4], groups labelled with a binomial share the same species label but were not included in [4], and other sequences are labelled as the Genbank accession followed by the species name associated with the sample. The tree was estimated using Bayesian inference and clades with posterior probability > 0.8 are marked with *. The full tree is given as Additional file 7_Rodent_cytb_full.tre.