



Fig. S.2. The Bayesian tree of the house shrew (*Suncus murinus*-*S. montanus* species complex), based on the mitochondrial cytochrome *b* gene sequences. See Table S.1 for the operational taxonomic unit (OTU) code. Numbers near nodes denote the posterior probabilities of Bayesian method for the confidence value of the nodes. Posterior probabilities less than 50 % are omitted.