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Title	Re-evaluation of the phylogeny based on mitochondrial cytochrome b gene in the house shrew, Suncus murinus-S. montanus species complex, with special reference to Yemen and Myanmar populations
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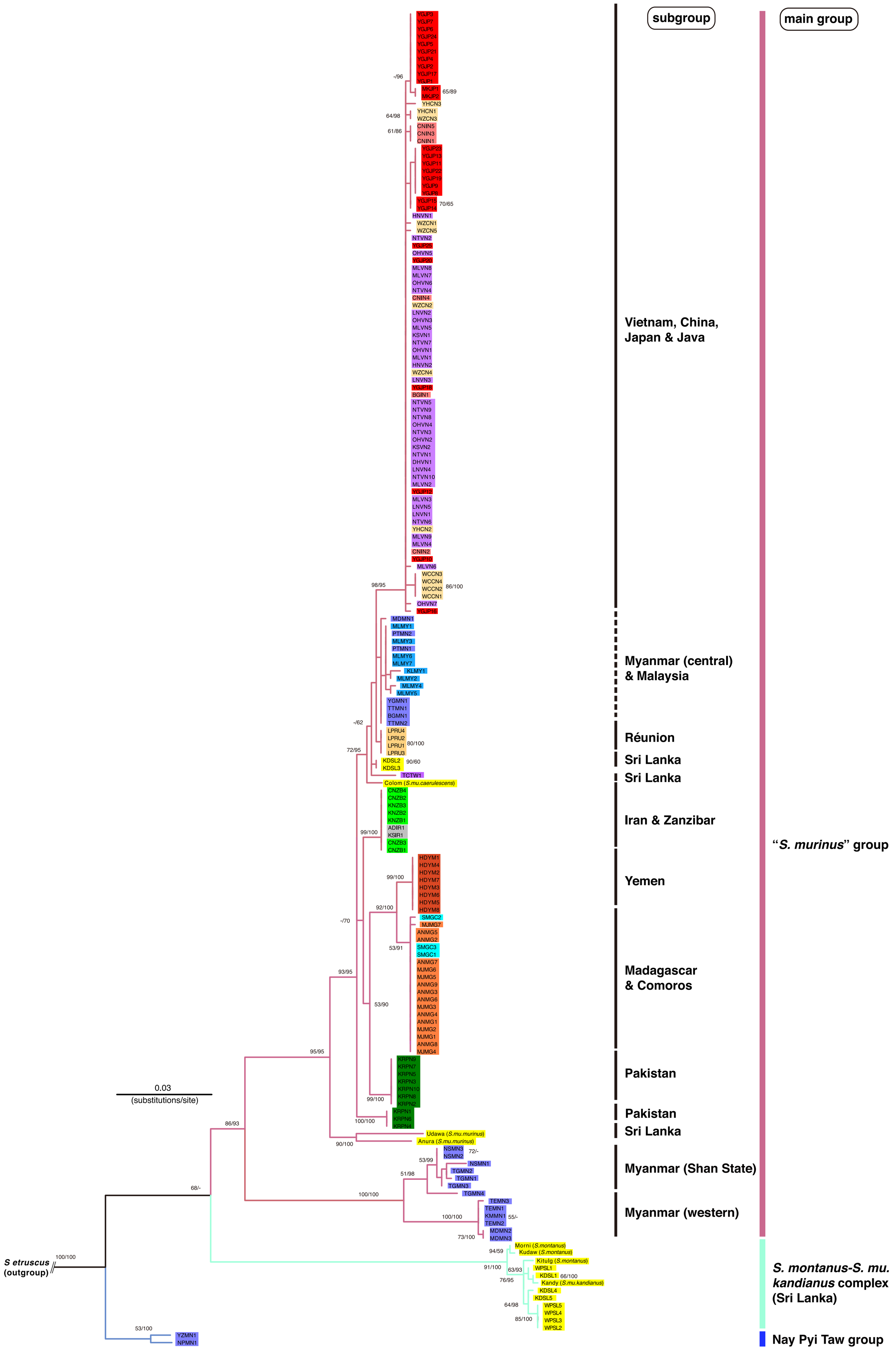

Fig. S.1. The maximum likelihood tree of the house shrew (*Suncus murinus*-*S. montanus* species complex), based on the mitochondrial cytochrome *b* gene sequences. See Table S.1 for the operational taxonomic unit (OTU) code. The fill colors of the OTUs indicates the locality of the shrew and is similar to that in Table S.1. Numbers near nodes denote the bootstrap values of the maximum likelihood method and the posterior probabilities of Bayesian method, respectively, for the confidence value of the nodes. Bootstrap values and poste-