Fig. 8.2: The Bayesian tree of the house shrew (Suncus murinus, x. variegatus species complex), based on the mitochondrial cytochrome b gene sequence. See Table S.1 for the operational taxonomic unit (OTU) code. Numbers near nodes denote the posterior probability of Bayesian method for the confidence value of the nodes. Posterior probabilities less than 50% are omitted.