

Figure S3. Phylogenetic reconstruction of tribes Arvicolini and Clethrionomyini using alignment with excluded third codon position

The trees were inferred from the concatenated dataset of 13 mitochondrial protein-coding genes. Node labels display BI/ML support, black circles show nodes with 0.95-1.0/95-100 support. Cytochrome *b* sequences are marked with asterisk

