



Supp. file 1. Maximum Likelihood (ML) phylogenetic tree based on the fragment of mitochondrial Cytochrome b (cytb) gene among the genera *Amphiesma*, *Amphiesmoides*, *Hebius*, *Herpetoreas*, and *Sahyadriophis*. The ultrafast bootstrap (UFB) support values are given at each branch. <https://doi.org/10.5852/ejt.2024.932.2519.11303>