



**Fig. S1.** A maximum likelihood tree reconstructed using IQTREE with a concatenated dataset of five genes. The five mitochondrial gene tree also indicates the monophyly of *Ahaetulla flavescens* **comb. nov.** and the presence of two more lineages (Polillo & Luzon Island from Philippines, & Borneo) belonging to *Ahaetulla prasina* complex. However, the nodes for the Indonesian *A. prasina* complex are weakly supported as there is inadequate data for 12S or COI genes.  
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