

Supplementary Figure S2. Updated mtDNA-genealogy of the *Microhyla* – *Glyphoglossus* assemblage (collapsed tree).

BI genealogy of *Microhyla* and *Glyphoglossus* reconstructed from 2478 bp of mtDNA fragment. Values at nodes correspond to BI PP/ML BS, respectively; black and white circles correspond to well-supported ($\text{BI PP} \geq 0.95$; $\text{ML BS} \geq 90$) and moderately supported ($0.95 > \text{BI PP} \geq 0.90$; $90 > \text{ML BS} \geq 75$) nodes, respectively; no circles indicate unsupported nodes. Color marking of species groups in *Microhyla* species complex corresponds to Figs. 4–5, but not to Fig. 2. Photos by Nikolay A. Poyarkov, Indraneil Das, Vladislav A. Gorin, Parinya Pawangkhanant, Luan Thanh Nguyen, and Evgeniya N. Solovyeva.

