

SUPPLEMENTARY FIGURES S1, S2

Gene Flow Increases Phylogenetic Structure and Inflates Cryptic Species Estimations:

A Case Study on Widespread Philippine Puddle Frogs (*Occidozyga laevis*)

Kin Onn Chan, Carl R. Hutter, Perry L. Wood, Jr., Yong-Chao Su, Rafe M. Brown

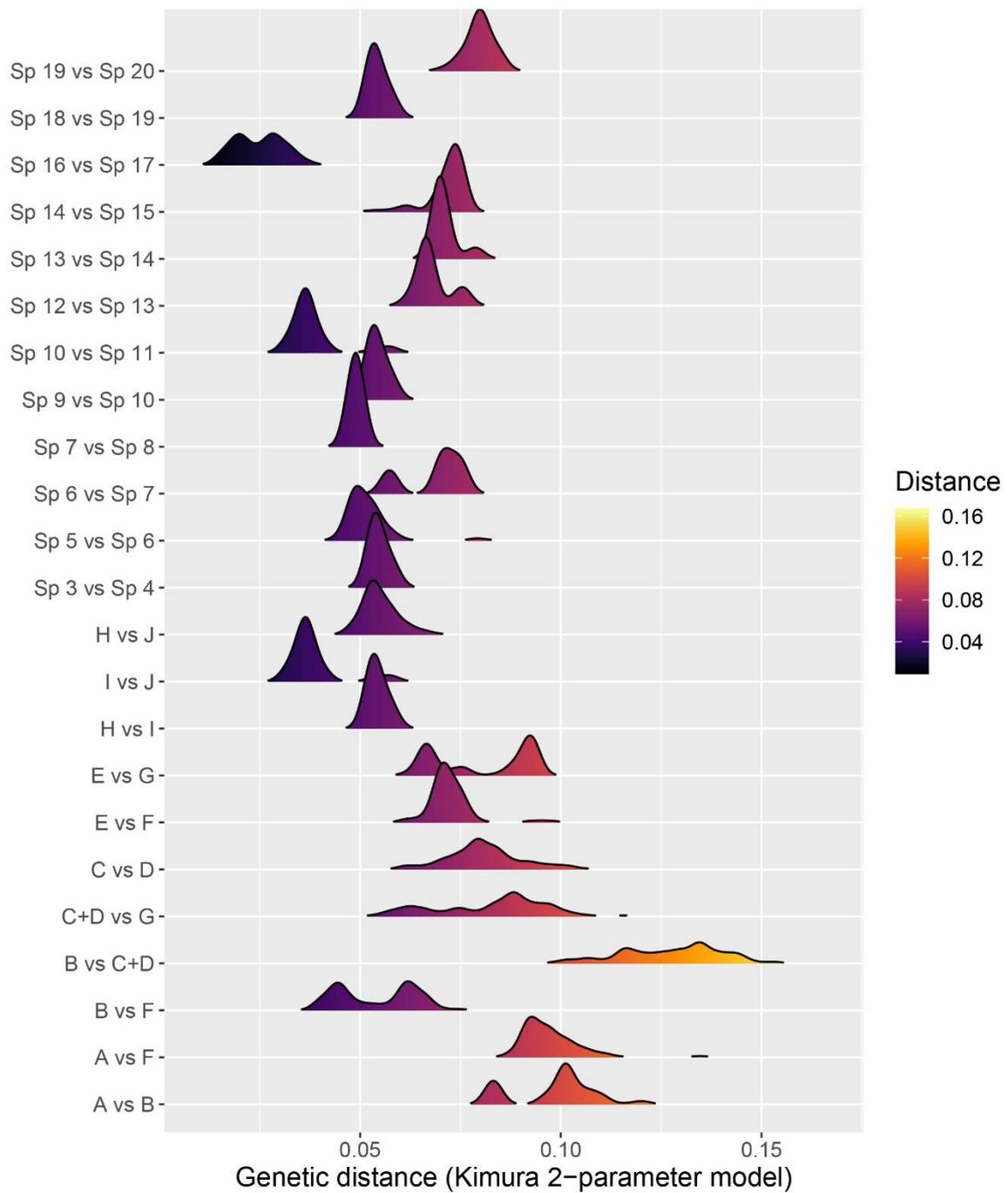


Fig. S1. Comparison of genetic distances among major clades and putative species calculated based on the Kimura 2-parameter model.

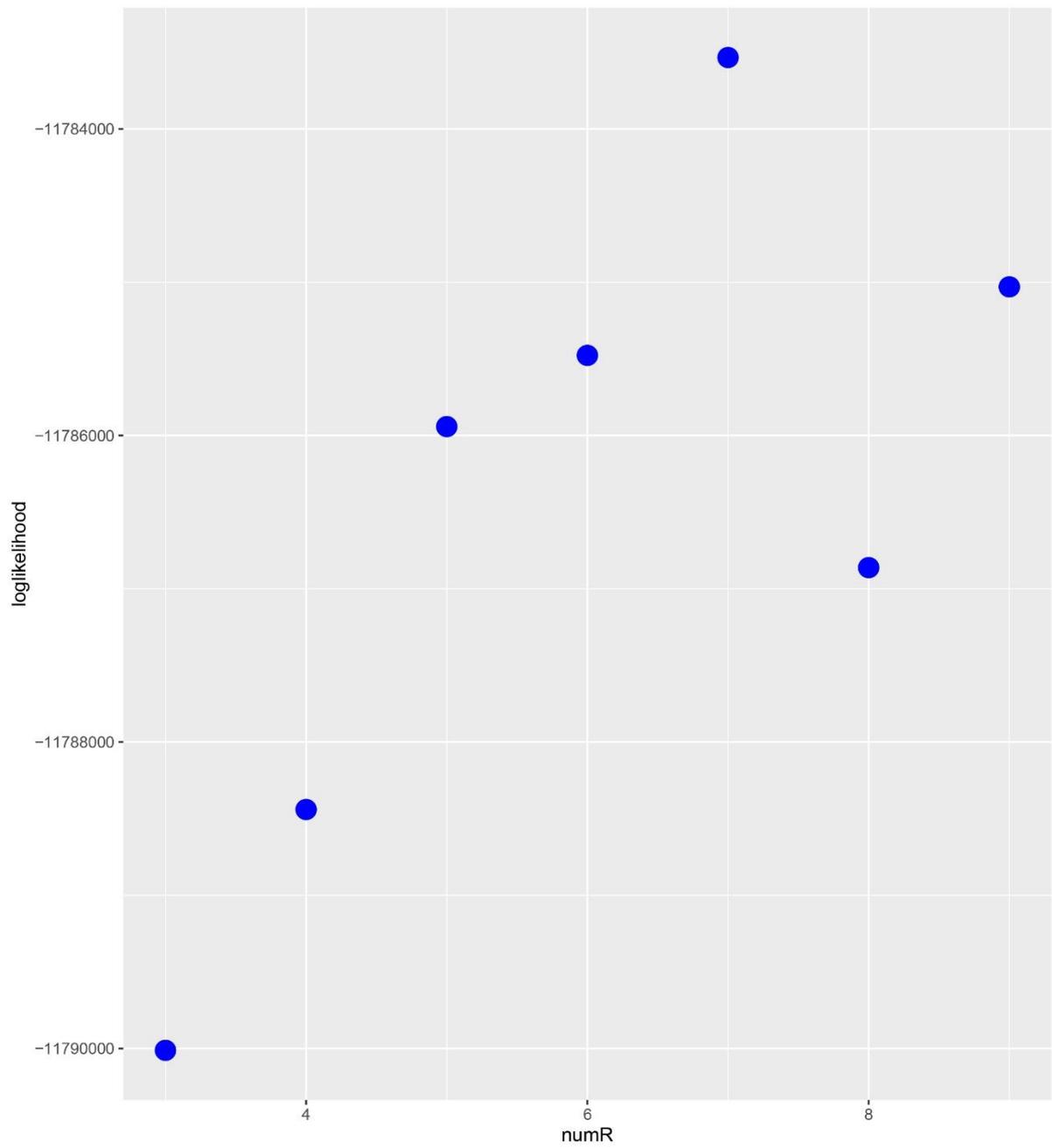


Fig. S2. Loglikelihoods vs. number of maximum reticulations (numR) of species networks inferred using the Maximum Pseudo-Likelihood approach.