

SUPPLEMENTARY FIGURES S1, S2

**Gene Flow Increases Phylogenetic Structure and Inflates Cryptic Species Estimations:
A Case Study on Widespread Philippine Puddle Frogs (*Occidozyga laevis*)**

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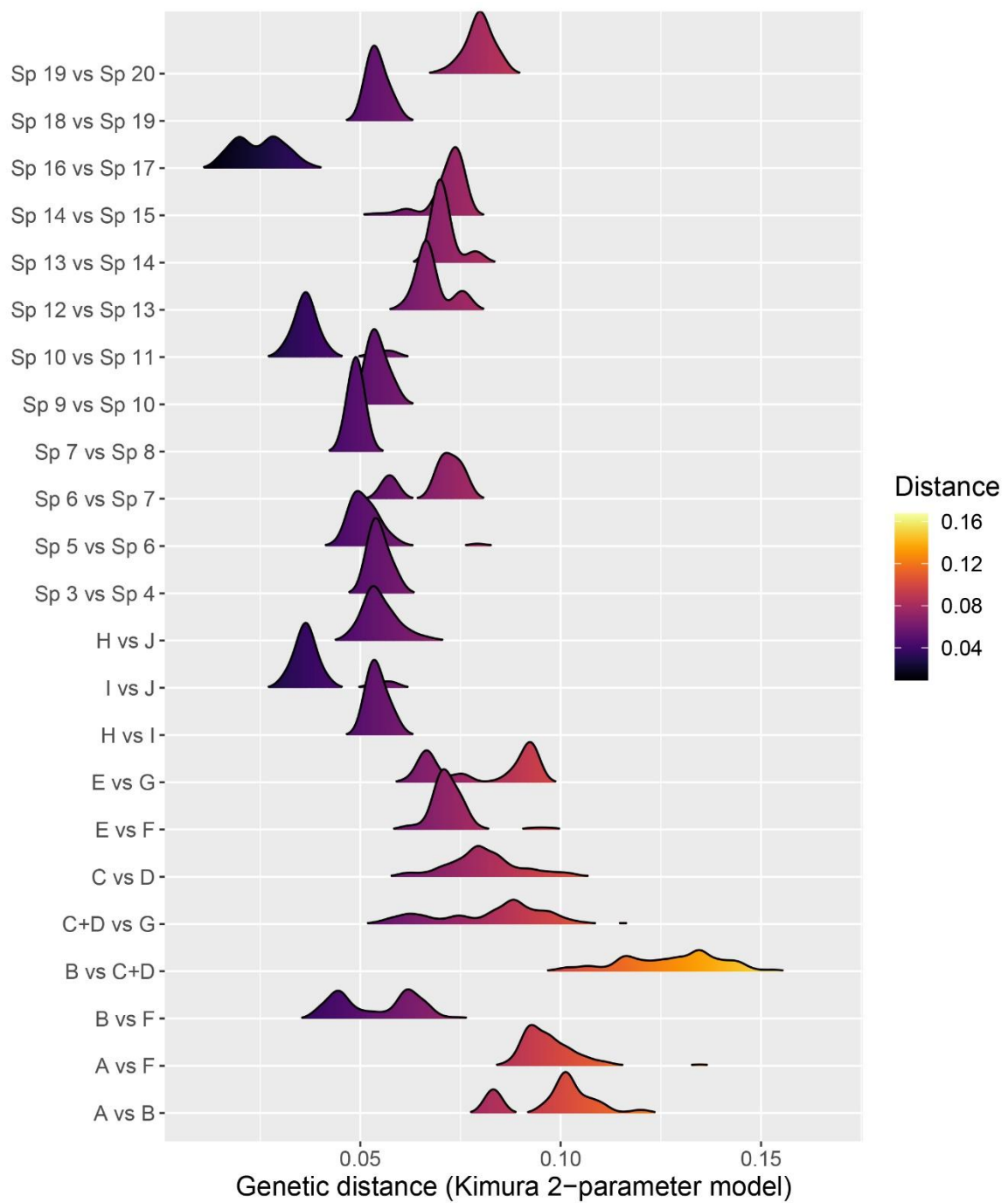


Fig. S1. Comparison of genetic distances among major clades and putative species calculated based on the Kimura 2-parameter model.

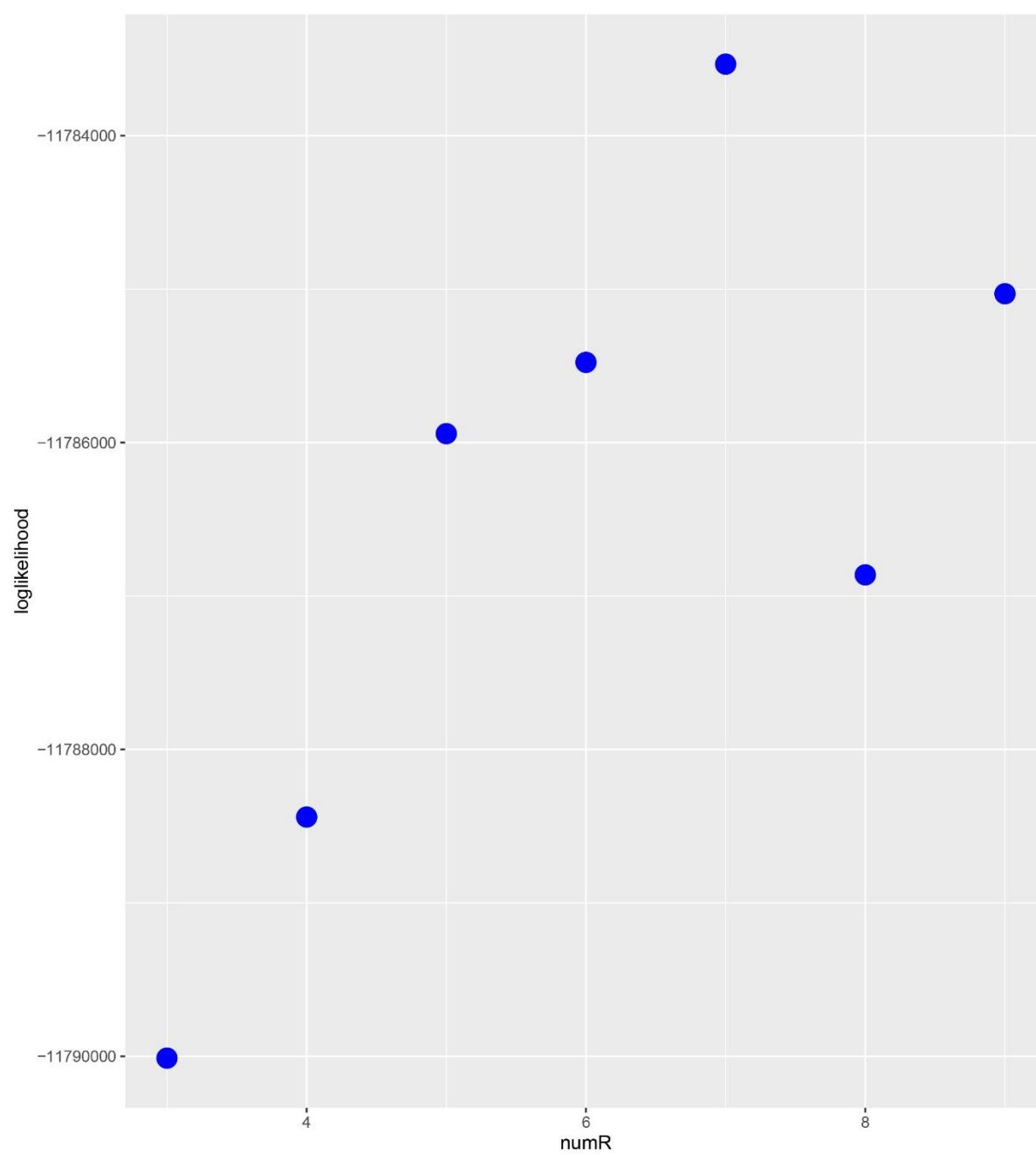


Fig. S2. Loglikelihoods vs. number of maximum reticulations (numR) of species networks inferred using the Maximum Pseudo-Likelihood approach.