

SUPPLEMENTARY DATA 4

SUPPLEMENTARY DATA 4.1

Phylogenetic relationships of *Rhinella* and outgroups recovered in the maximum likelihood analysis with IQ-TREE (log-likelihood of consensus tree: -1202690.336). Numbers around nodes are the ultrafast bootstrap support values. An asterisk (*) indicates 100% ultrafast bootstrap support. A skeletal topology of the complete tree is shown for reference, with the magnified section marked in white.

SUPPLEMENTARY DATA 4.2

Phylogenetic relationships of *Rhinella* recovered in the maximum likelihood analysis with IQ-TREE (log-likelihood of consensus tree: -1202690.336). The clades and species groups shown are those recognized in this study. **Part 1 of 4. The *R. marina* Clade (1): the *R. arunco*, *R. spinulosa*, and *R. granulosa* Groups.** Numbers around nodes are the ultrafast bootstrap support values. An asterisk (*) indicates 100% ultrafast bootstrap support. A skeletal topology of the complete tree is shown for reference, with the magnified section marked in white.

SUPPLEMENTARY DATA 4.3

Phylogenetic relationships of *Rhinella* recovered in the maximum likelihood analysis with IQ-TREE (log-likelihood of consensus tree: -1202690.336). The clades and species groups shown are those recognized in this study. **Part 2 of 4. The *R. marina* Clade (2): the GIM and the *R. crucifer* and *R. marina* Groups.** Numbers around nodes are the ultrafast bootstrap support values. An asterisk (*) indicates 100% ultrafast bootstrap support. A skeletal topology of the complete tree is shown for reference, with the magnified section marked in white.

SUPPLEMENTARY DATA 4.4

Phylogenetic relationships of *Rhinella* recovered in the maximum likelihood analysis with IQ-TREE (log-likelihood of consensus tree: -1202690.336). The clades and species groups shown are those recognized in this study. Part 3 of 4. **The *R. margaritifera* Clade (1): *R. sternosignata* and the *R. veraguensis* and *R. festae* Groups.** Numbers around nodes are the ultrafast bootstrap support values. An asterisk (*) indicates 100% ultrafast bootstrap support. A skeletal topology of the complete tree is shown for reference, with the magnified section marked in white.

SUPPLEMENTARY DATA 4.5

Phylogenetic relationships of *Rhinella* recovered in the maximum likelihood analysis with IQ-TREE (log-likelihood of consensus tree: -1202690.336). The clades and species groups shown are those recognized in this study. **Part 4 of 4. The *R. margaritifera* Clade (2): the *R. margaritifera* Group.** Numbers around nodes are the ultrafast bootstrap support values. An asterisk (*) indicates 100% ultrafast bootstrap support. A skeletal topology of the complete tree is shown for reference, with the magnified section marked in white.









