



FIGURE S2 | Maximum likelihood tree of *Rineloricaria* species using 223 COI sequences from Costa-Silva *et al.* (2015) and 10 sequences of *Rineloricaria nudipectoris* (657 bp) depicting the *Rineloricaria nudipectoris* clade. Sequence name and associated data are listed in Tab. 1 and Tab. S1. Numbers above branches represent bootstrap support, except for those below 0.5% which were suppressed. (Click here to see the full image)



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REFERENCE

- Costa-Silva GJ, Rodriguez MS, Roxo FF, Foresti F, Oliveira C. Using different methods to access the difficult task of delimiting species in a complex Neotropical hyperdiverse group. PLoS ONE. 2015; 10:1–12. <https://doi.org/10.1371/journal.pone.0135075>

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