



FIGURE S4 | Phylogenetic relationships among Dasyatinae, Neotrygoninae, and Urogyminae representatives based on a Bayesian Inference on an alignment of 11,471 base pairs with 135 sequences. For each node, posterior probabilities are given. Genus *Hypanus* in red, subfamily Dasyatinae in blue (except *Hypanus*), subfamily Neotrygoninae in green, and subfamily Urogyminae in orange. Download the image here: <https://www.ni.bio.br/content/v22n1/1982-0224-2023-0046/supplementary/1982-0224-ni-22-01-e230046-gfS4.zip>



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