



Figure S1. UCE phylogeny. The maximum-likelihood phylogeny of the 90% complete concatenated matrix containing 799 UCE loci (702,574 base pairs), adapted from Ješovnik et al., 2017. Black circles at nodes indicate 98–100 bootstrap frequencies (N=500). Taxon names include DNA extraction codes (numbers following species names) and country codes as follows: BR–Brazil, CO–Colombia, CR–Costa Rica, HN–Honduras, EC–Ecuador, GY–Guyana, GF– French Guiana, GT–Guatemala, MX–Mexico, NI–Nicaragua, PA–Panama, PE–Peru, SR–Suriname, TT– Trinidad and Tobago, VE– Venezuela.