

Supplementary Fig. S2 Maximum clade credibility trees for mitochondrial COI haplotypes (on the left) and nuclear ITS (on the right) data sets. Vertical bars in the middle labelled Species I-X, CC ("*L. clapedianus-cervix*"), LM ("*L. maumeensis*") and LC ("*L. clapedianus*"). Dashed lines connecting the two trees connect shared specimens in the COI and ITS data sets. Nodal support of the tree corresponds to posterior supports from BEAST analysis, with a black dot signifying ≥ 0.99 value. The GAb, GA and Gb within the parentheses stands for Primary Hypothetical Species suggested by ABGD and bGMYC, or only by ABGD or bGMYC.

