

Vondrák J., Svoboda S., Košnar J., Malíček J., Šoun J., Frolov I., Svensson M., Novotný P. & Palice Z. (2023) Martin7: a reference database of DNA barcodes for European epiphytic lichens and its taxonomic implications. – *Preslia* 95: 311–345.

Supplementary Fig. S2. MtSSU tree with *Bacidina acerina* and *B. paradoxa*. GTRG model with gamma distribution was used as a model of sequence evolution. Trees were constructed using Bayesian inference run for 158,000 generations.

