

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. S6. MtSSU tree with *Japewia gyrophorica*. HKYG model with gamma distribution was used as a model of sequence evolution. Trees were constructed using Bayesian inference run for 53,000 generations.

