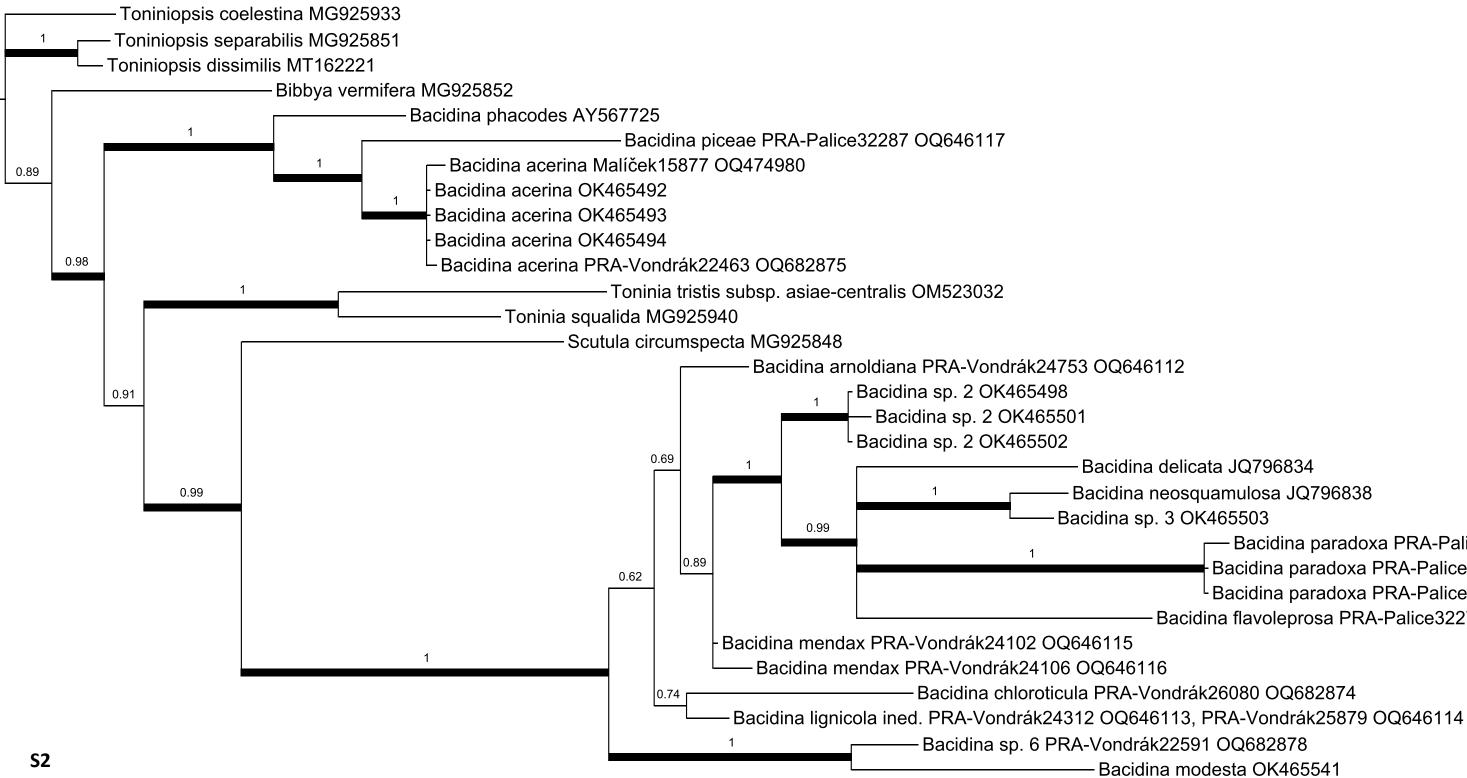
Vondrák J., Svoboda S., Košnar J., Malíček J., Šoun J., Frolov I., Svensson M., Novotný P. & Palice Z. (2023) Martin7: a reference database of DNA barcodes for European epiphytic lichens and its taxonomic implications. – Preslia 95: 311–345.

**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.



Bacidina neosquamulosa JQ796838 - Bacidina paradoxa PRA-Palice26142 OQ682876 Bacidina paradoxa PRA-Palice32056 OQ646118 <sup>L</sup>Bacidina paradoxa PRA-Palice33646 OQ682877 Bacidina flavoleprosa PRA-Palice32275 OQ646119

Bacidina modesta OK465541