

Supplementary Fig. S4. A. Fragment of ITS alignment (positions 419–560 bp) with the single parsimony informative position at 513 bp (indicated by asterisk) based on which three ribotypes were defined: (i) ribotype ‘A’, (ii) ribotype ‘G’ and (iii) ribotype ‘R’. **B.** Fragment of trnH-psbA (cpDNA) alignment (positions 1-80 bp) with variable region at 43–52 bp (indicated by asterisks) based on which two haplotypes were defined (i) Haplotype ‘A’ (sequence ‘TTTTTTTGCT’ in positions 43–52 bp) and (ii) Haplotype ‘B’ (sequence ‘AGCA-AAAA’ in positions 43–52 bp). Sequences are labelled according to localities sampled: RKJ, Krivé jezero; P1, Plačkův les; S, Šúr; RS, Šúr (reference).

