



**Figure S6.** ITS1 phylogenetic analysis under parsimony as the optimality criterion. Numbers on nodes indicate Goodman-Bremer support values and colored circles represent bootstrap resampling frequencies (black =  $\geq 95$ , grey =  $\geq 75$ , absent =  $< 75$ ). Scale bar represents the number of nucleotide transformations. Table S4 contains further details on sequences used to reconstruct this phylogeny.