



Supplementary 6. ML tree showing the internal phylogeny of *Astrothelium* based on ITS with an alignment length of 455 bp. Maximum likelihood bootstrap support above 70% (left) and Bayesian inference posterior probabilities above 0.95 (right) are shown at nodes (ML-BP/B-PP). Terminals in boldface indicate newly generated sequences for this study. The type specimens in the phylogenetic tree were labeled by letter T.