



0.07

Supplementary figure 11. Single-locus tree for Lecanoraceae *MCM7*. Maximum-likelihood tree inferred with the TIM+F+I+G4 substitution model. Support values are UFboot2 with 5000 replicates. Bipartitions with bootstrap support ≥ 95 are indicated with bold branches. Bold tip names indicate specimens sequenced in this study. Scale indicates substitutions per site.