



Fig. S6. A maximum likelihood phylogenetic tree derived from the combined *ITS*, *LSU*, *SSU*, *MCM7*, *RPB1*, and *TEF1* DNA sequence data sets. Nodes having Bayesian inference posterior probabilities ≥ 0.95 and maximum likelihood bootstrap values $\geq 75\%$ are shown. The tree is rooted with *Pyricularia grisea* M83.