





Figure S1. Phylogenetic tree retrieved from RAxML analyses of a combined ITS, LSU and TEF1- α sequence data of *Apiospora*, and other related taxa in the family Apiosporaceae. Bootstrap support values for ML equal or greater than 60% and Bayesian posterior probabilities greater than 0.90 are indicated at the nodes as ML/PP. The ex-type strains are in bold. The new species are in red and new record and new combination species are in blue. The tree is rooted to *Sporocadus trimorphus* (CBS 114203).