



Fig 2 Maximum likelihood phylogenetic trees for the ITS, *RPB2*, *MCM7*, *TEF1*, LSU and SSU gene sequences. *Caliciopsis pinea* (*Coryneliaceae*, *Coryneliales*) was chosen as the outgroup. Isolates included in this study representing new species are shown in colour. Numbers on branches correspond to bootstrap values (n = 1,000).