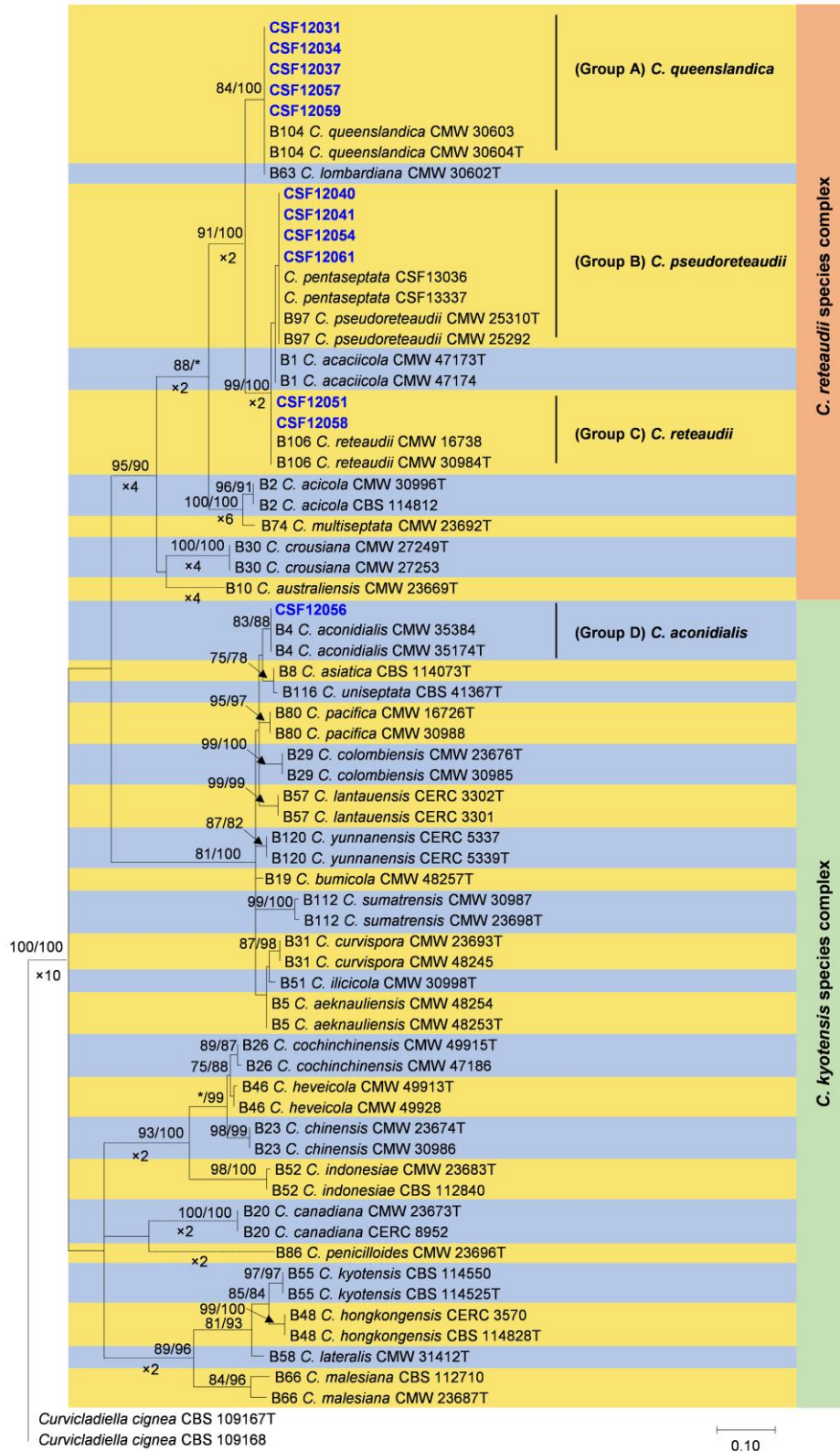Supplementary Fig. S4. Phylogenetic trees obtained from Maximum Likelihood (ML) analyses of *his3* region for species in the *C. reteaudii* and *C. kyotensis* species complexes. Bootstrap support values ≥70 % for ML and MP are indicated above the branches as ML/MP. Bootstrap support values < 70% are marked with "*". Isolates obtained in this study are highlighted in blue and in bold type; "T" represents ex-type isolates. "B" species codes are consistent with the published results in Liu et al. (2020). *Curvicladiella cignea* (CBS 109167 and CBS 109168) was used as the outgroup taxon.