

Supplementary Materials

Supplementary Figure S1. Phylogenetic tree of *Calonectria* species based on maximum likelihood (ML) analyses of the *tef1* gene sequences in this study. Bootstrap support values $\geq 70\%$ are presented above the branches as follows: ML/MP. Bootstrap values $< 70\%$ and absent are marked with “*”. Isolates highlighted in five different colours and bold were obtained in this study. Ex-type isolates are marked with “T”. The “B” species codes are consistent with the recently published results in Liu and co-authors [1]. *Curviciadiella cignea* (CBS 109167 and CBS 109168) was used as an outgroup taxon.

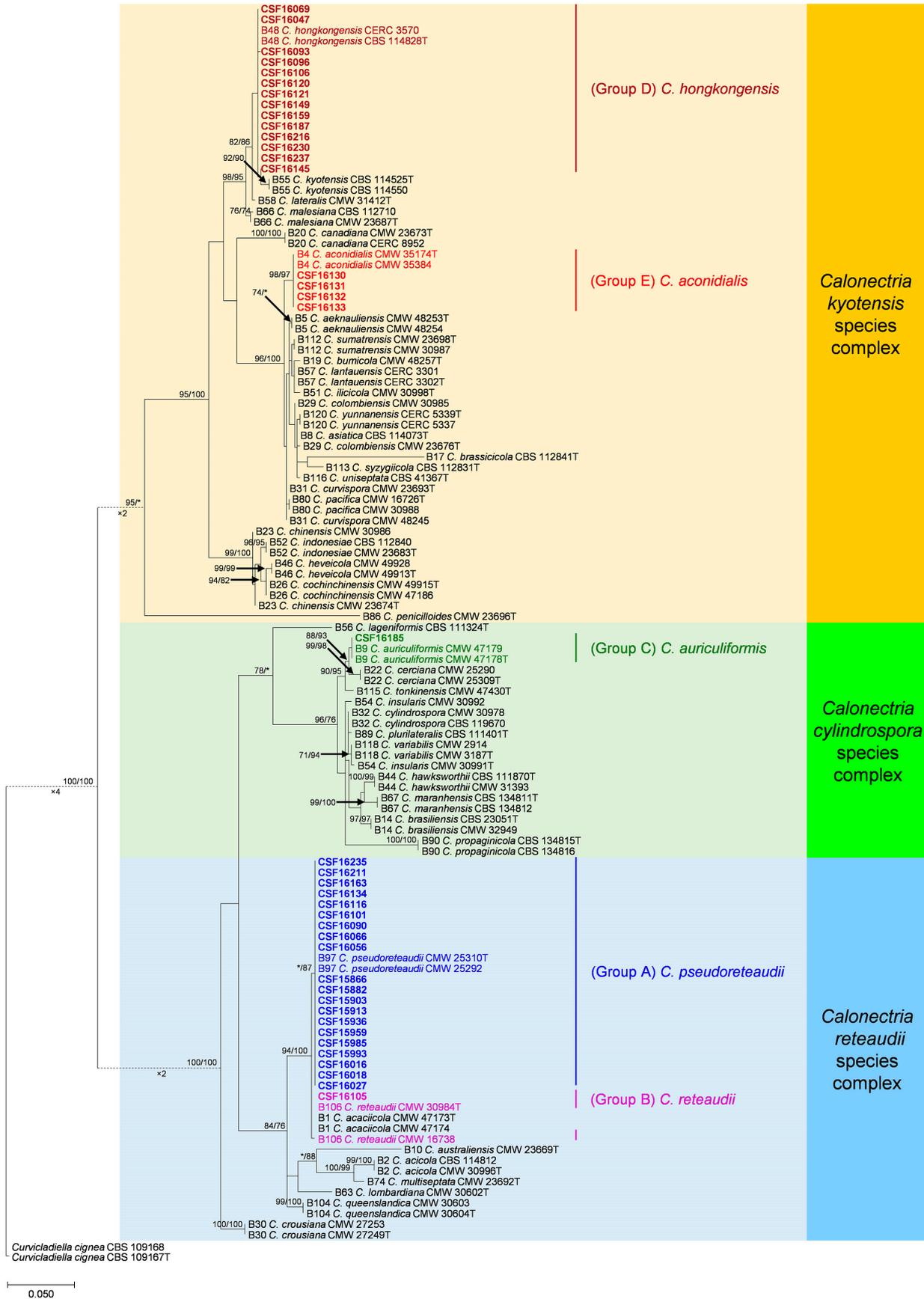
Supplementary Figure S2. Phylogenetic tree of *Calonectria* species based on maximum likelihood (ML) analyses of the *tub2* gene sequences in this study. Bootstrap support values $\geq 70\%$ are presented above the branches as follows: ML/MP. Bootstrap values $< 70\%$ and absent are marked with “*”. Isolates highlighted in five different colours and bold were obtained in this study. Ex-type isolates are marked with “T”. The “B” species codes are consistent with the recently published results in Liu and co-authors [1]. *Curviciadiella cignea* (CBS 109167 and CBS 109168) was used as an outgroup taxon.

Supplementary Figure S3. Phylogenetic tree of *Calonectria* species based on maximum likelihood (ML) analyses of the *cmdA* gene sequences in this study. Bootstrap support values $\geq 70\%$ are presented above the branches as follows: ML/MP. Bootstrap values $< 70\%$ and absent are marked with “*”. Isolates highlighted in five different colours and bold were obtained in this study. Ex-type isolates are marked with “T”. The “B” species codes are consistent with the recently published results in Liu and co-authors [1]. *Curviciadiella cignea* (CBS 109167 and CBS 109168) was used as an outgroup taxon.

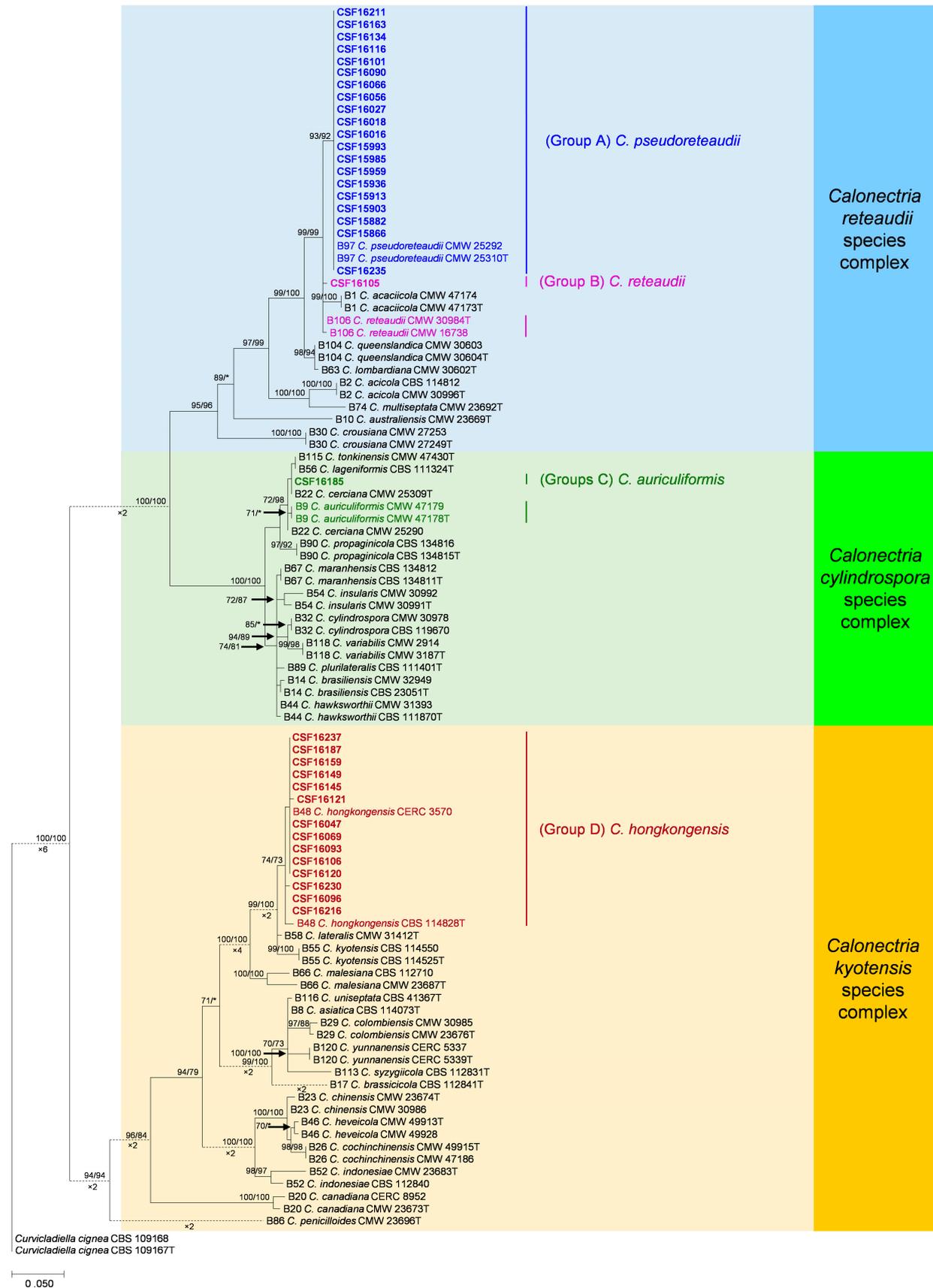
Supplementary Figure S4. Phylogenetic tree of *Calonectria* species based on maximum likelihood (ML) analyses of the *his3* gene sequences in this study. Bootstrap support values $\geq 70\%$ are presented above the branches as follows: ML/MP. Bootstrap values $< 70\%$ and absent are marked with “*”. Isolates highlighted in five different colours and bold were obtained in this study. Ex-type isolates are marked with “T”. The “B” species codes are consistent with the recently published results in Liu and co-authors [1]. *Curviciadiella cignea* (CBS 109167 and CBS 109168) was used as an outgroup taxon.

Reference

1. Liu, Q.L.; Li, J.Q.; Wingfield, M.J.; Duong, T.A.; Wingfield, B.D.; Crous, P.W.; Chen, S.F. Reconsideration of species boundaries and proposed DNA barcodes for *Calonectria*. *Stud. Mycol.* **2020**, doi:10.1016/j.simyco.2020.08.001.

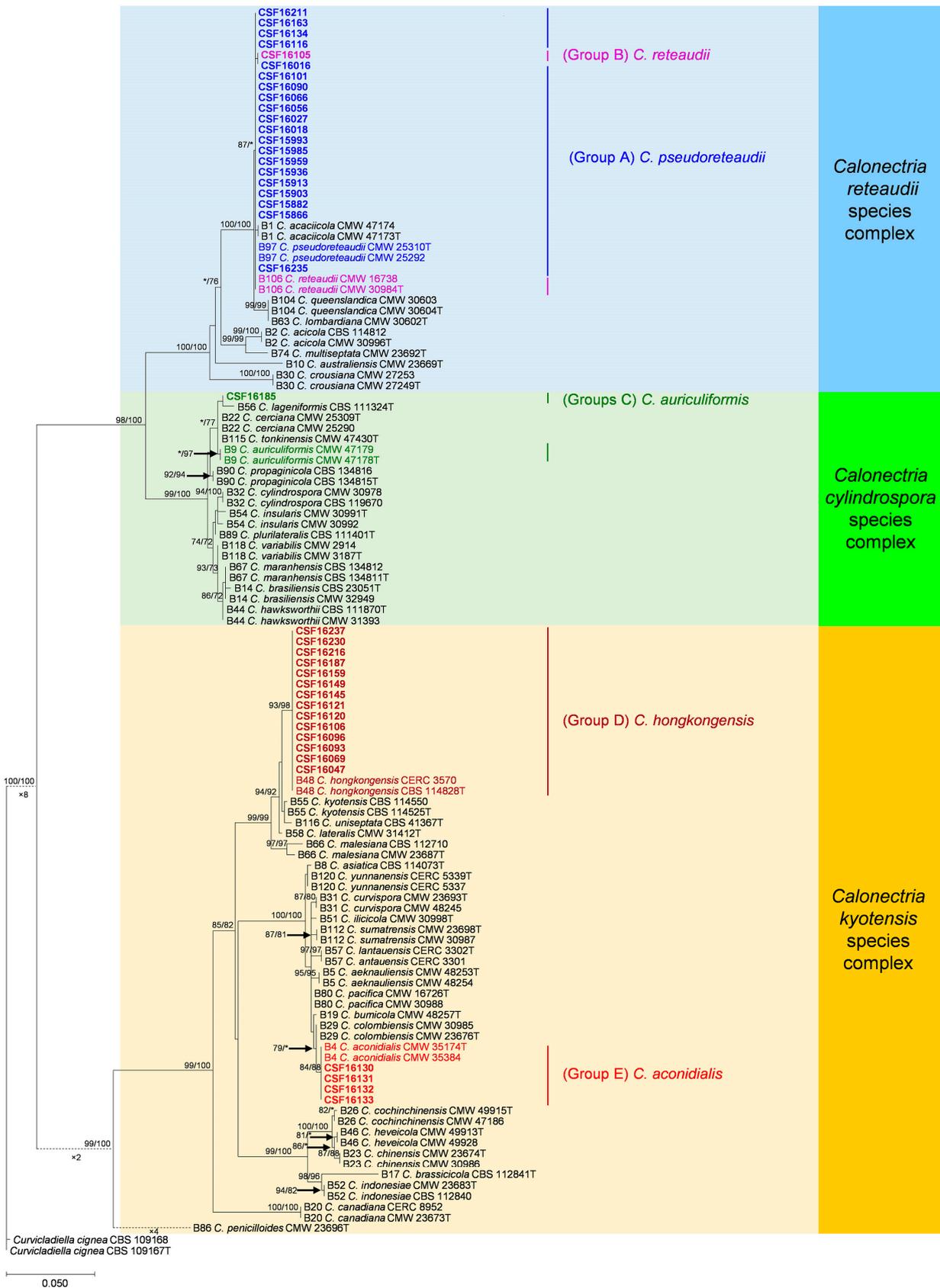


Supplementary Figure S1.



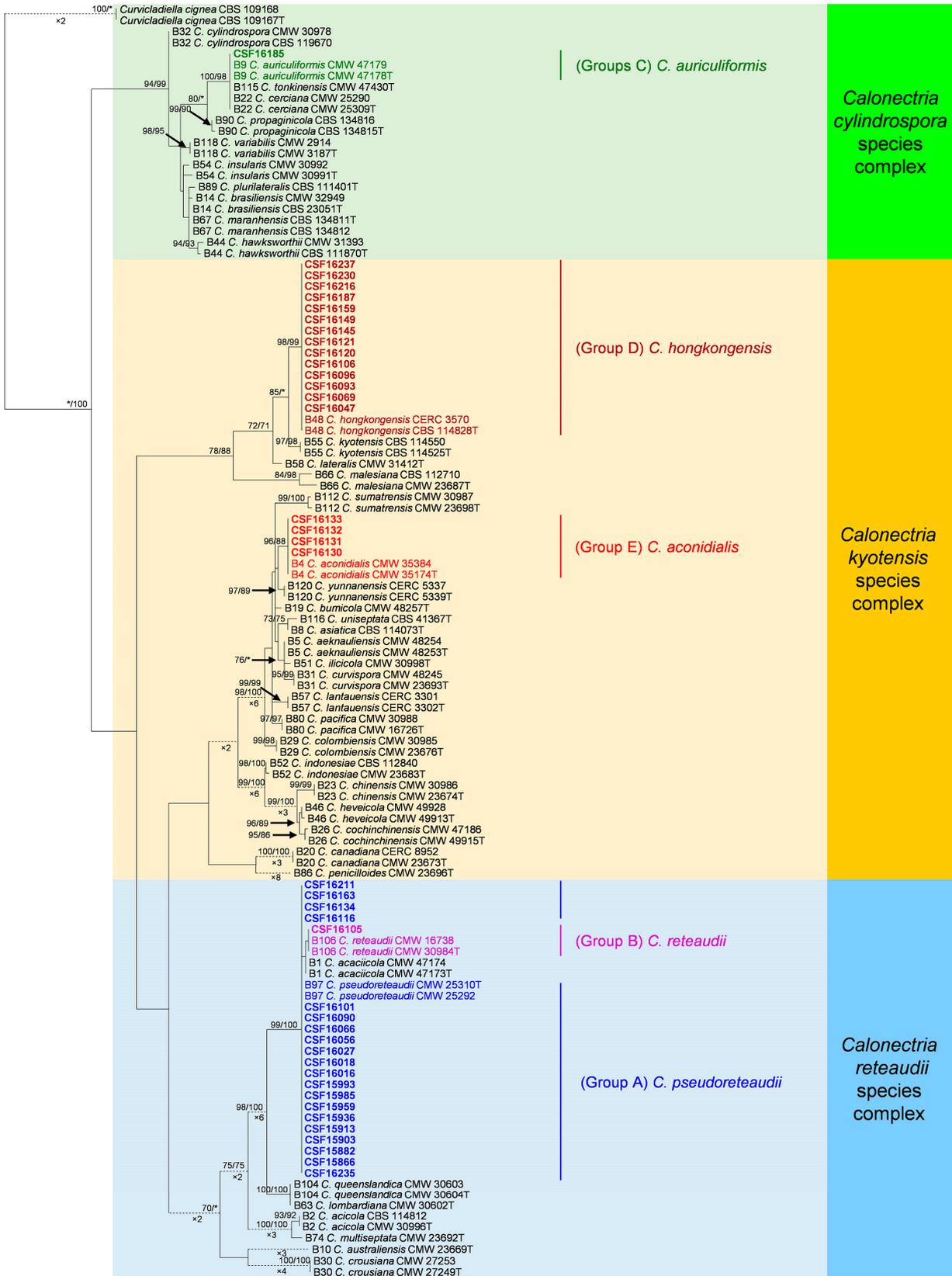
Supplementary Figure S2.

cmdA



Supplementary Figure S3.

his3



Supplementary Figure S4.