

SUPPLEMENTARY MATERIAL FOR

Synopsis of *Leptosphaeriaceae* and Introduction of Four New Taxa from China

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LIST OF SUPPORTING INFORMATION

Figure S1. Phylogram generated from Bayesian inference analysis based on combined ITS, LSU, SSU and *tub2* sequence data.

Figure S2. Phylogram generated from maximum likelihood analysis based on combined ITS, LSU, SSU, *tub2*, and *rpb2* sequence data.

Figure S3. Phylogram generated from maximum likelihood analysis based on combined ITS, LSU, SSU and *rpb2* sequence data.

Figure S4: Phylogram generated from maximum likelihood analysis using *rpb2* sequence data.

Figure S5: Phylogram generated from maximum likelihood analysis using *tub2* sequence data.



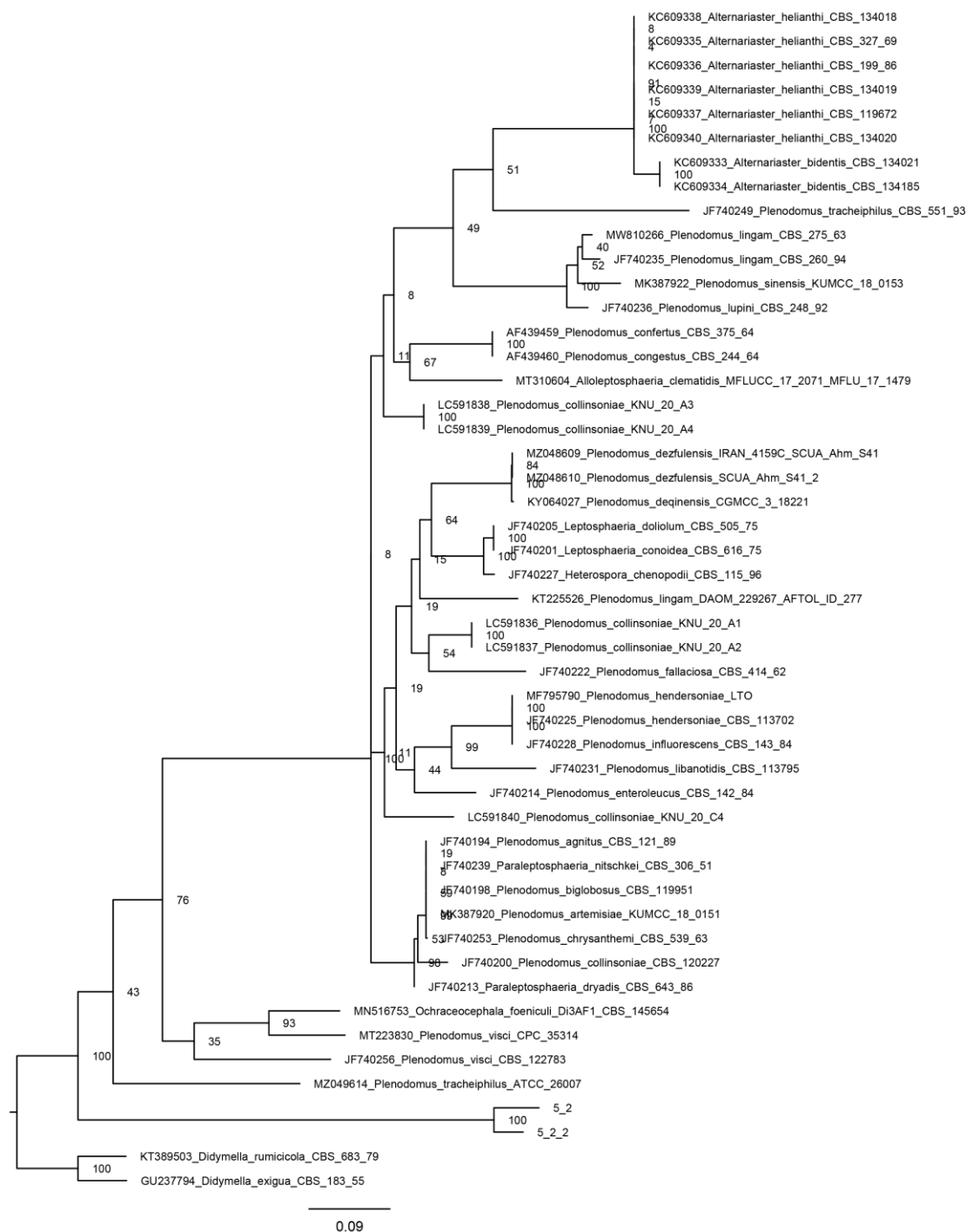


Figure S4: Phylogram generated from maximum likelihood analysis using *rpb2* sequence data. The strains are named following the accession number of the ITS region.

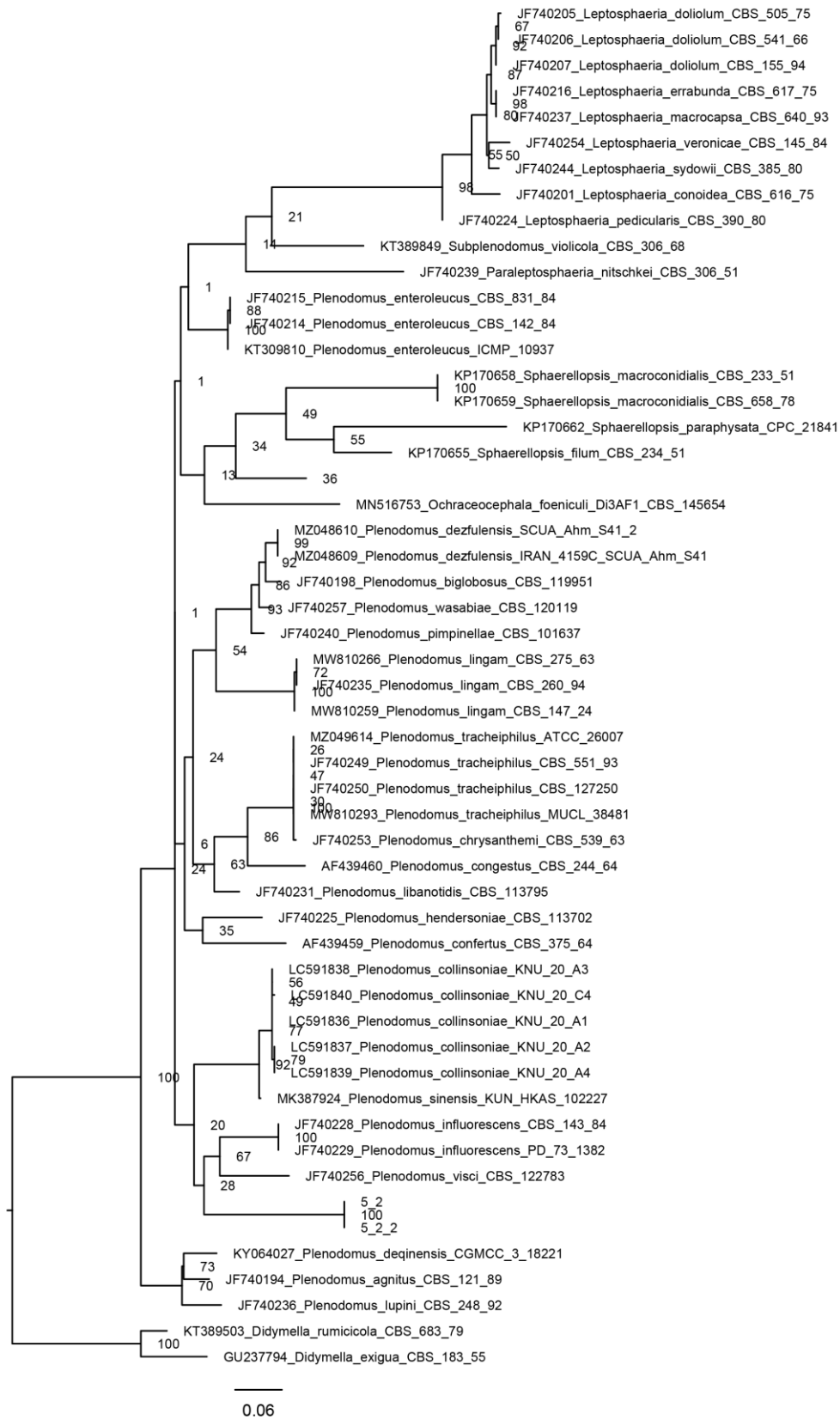


Figure S5: Phylogram generated from maximum likelihood analysis using *tub2* sequence data. The strains are named following the accession number of the ITS region.