



Fig. S3. Maximum Likelihood (ML) phylogenetic tree of the SSU sequence alignment of the new genus *Endophytium* and related meristematic and meristematic-like fungi in *Dothideomycetes*. New taxa are in bold face. Thickened branches indicate full statistical support in all analyses. Confidence values for ML-BS $\geq 70\%$ (UFboot2/RAxML) are included near the nodes and the "-" indicates statistical support below the threshold values. The scale bar represents the expected number of changes per site. The tree was rooted to *Xylaria* (*Xylariales*, *Sordariomycetes*) and *Sordaria* (*Sordariales*, *Sordariomycetes*). Families and orders are shown in coloured blocks and indicated to the right of the tree.