



SUPPLEMENTAL FIGURE 1. A species maximum likelihood phylogeny based on comparisons between the *LSU* regions of the *Absidia* species that are most closely related to *A. urquhartii*. For *A. urquhartii*, the strain names are provided, with their GenBank accessions in table 1. For other *Absidia* species, GenBank accessions are in grey font. The tree is rooted with *Chlamydoabsidia* (= *Absidia*) *padenii*. The numbers near the nodes are percentage values from bootstrapping (100 replicates).