

Figure S1: Phylogenetic estimation of relationships between OMF identified as A) *Ceratobasidium* and B) *Tulasnella*. Sample names match those in Table S1. Trees were constructed using FastTree 2.1.11 (fastest setting, GTR model, rate categories of sites = 20) in Geneious Prime to illustrate similarities between sequenced OMF. Numbers at nodes are Fasttree support values, which range from <0.01 (very weakly supported) to 1 (strongly supported). The scale bar at the bottom of each tree indicates expected number of substitutions.

Figure S1-A: *Ceratobasidium*



Figure S1-B. *Tulasnella*

