Table S9. List of commonOTUs identified by CD and CI methods



a One representative strain (from the CD method) and one representative clone (from the CI method) is presented for the 13 shared OTUs. b Abundance indicates the number of strains or clones for each OTU in the entire data set. c BLASTN score. d Accession number of the closest database match. EDFL, early diverging fungal lineages.