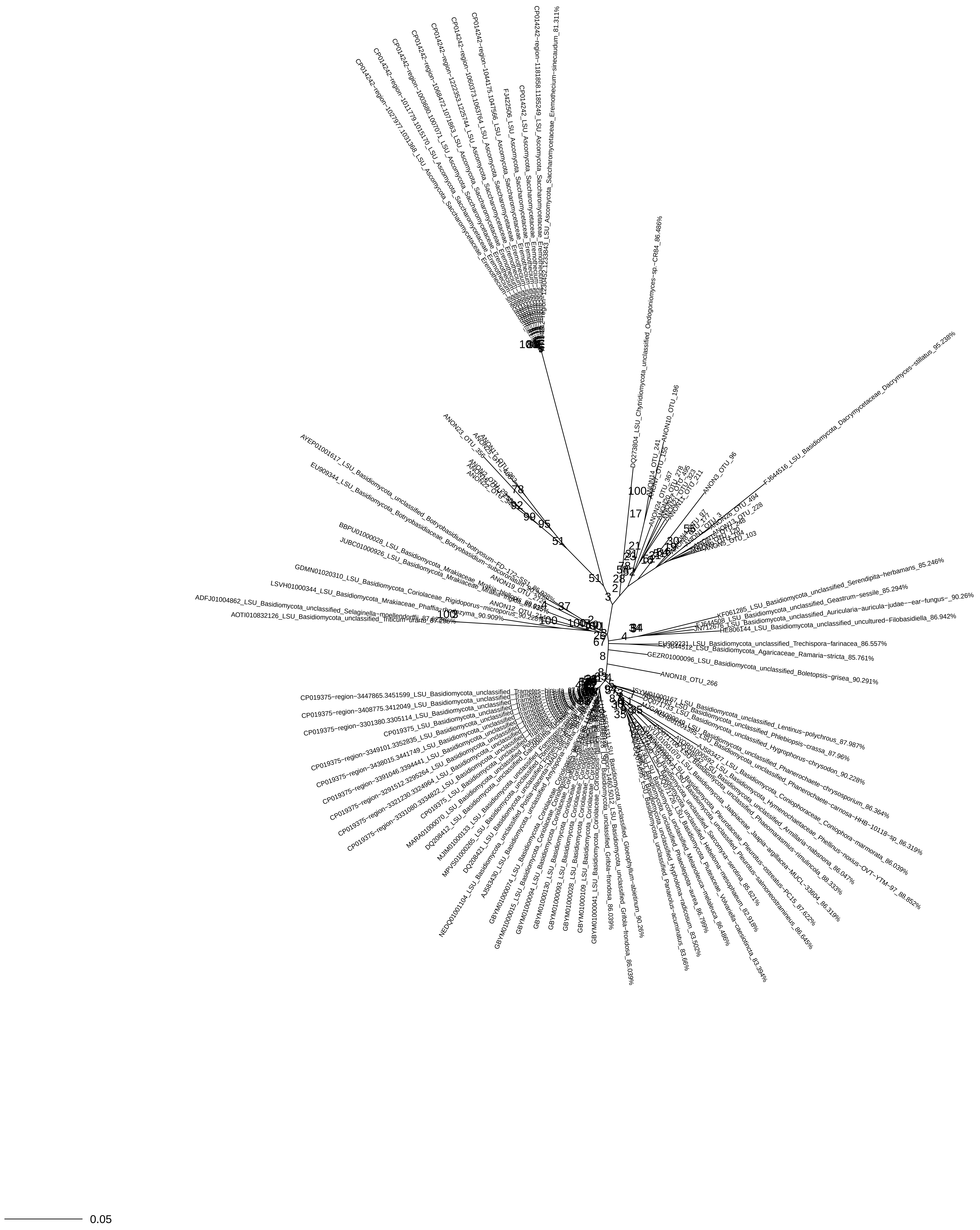
CP014242~region~1181858.1185249_LSU_Ascomycota_Saccharomycetaceae_Eremothecium~sinecaudum_81.311%
CP014242~region~1230452.1233843_LSU_Ascomycota_Saccharomycetaceae_Eremothecium~sinecaudum
CP014242_LSU_Ascomycota_Saccharomycetaceae_Eremothecium~sinecaudum
FJ422506_LSU_Ascomycota_Saccharomycetaceae_Eremothecium~sinecaudum
CP014242~region~1044175.1047566_LSU_Ascomycota_Saccharomycetaceae_Eremothecium~sinecaudum
CP014242~region~1060373.1063764_LSU_Ascomycota_Saccharomycetaceae_Eremothecium~sinecaudum
CP014242~region~1222353.1225744_LSU_Ascomycota_Saccharomycetaceae_Eremothecium~sinecaudum
CP014242~region~1068472.1071863_LSU_Ascomycota_Saccharomycetaceae_Eremothecium~sinecaudum
CP014242~region~1003680.1007071_LSU_Ascomycota_Saccharomycetaceae_Eremothecium~sinecaudum
CP014242~region~1011779.1015170_LSU_Ascomycota_Saccharomycetaceae_Eremothecium~sinecaudum
CP014242~region~1027977.1031368_LSU_Ascomycota_Saccharomycetaceae_Eremothecium~sinecaudum
DQ273804_LSU_Chytridiomycota_unclassified_Oedogoniomyces~sp.~CR84_86.486%
ANON10_OTU_196
ANON14_OTU_241
ANON7_OTU_155
ANON24_OTU_367
ANON20_OTU_278
ANON27_OTU_495
ANON21_OTU_323
ANON11_OTU_211
ANON3_OTU_96
FJ644516_LSU_Basidiomycota_Dacrymycetaceae_Dacrymyces~stillatus_95.238%
ANON4_OTU_97
ANON8_OTU_177
ANON1_OTU_3
ANON26_OTU_494
ANON13_OTU_228
ANON15_OTU_248
ANON6_OTU_120
ANON9_OTU_194
ANON5_OTU_103
KF061285_LSU_Basidiomycota_unclassified_Serendipita~herbamans_85.246%
FJ644508_LSU_Basidiomycota_unclassified_Geastrum~sessile_85.294%
JN712676_LSU_Basidiomycota_unclassified_Auricularia~auricula~judae~~ear~fungus~_90.26%
HE806144_LSU_Basidiomycota_unclassified_uncultured~Filobasidiella_86.942%
EU909231_LSU_Basidiomycota_unclassified_Trechispora~farinacea_86.557%
FJ644512_LSU_Basidiomycota_Agaricaceae_Ramaria~stricta_85.761%
GEZR01000096_LSU_Basidiomycota_unclassified_Boletopsis~grisea_90.291%
ANON18_OTU_266
JSYW01000167_LSU_Basidiomycota_unclassified_Lentinus~polychrous_87.987%
DQ071733_LSU_Basidiomycota_unclassified_Phlebiopsis~crassa_87.96%
MJGA01000049_LSU_Basidiomycota_unclassified_Hygrophorus~chrysodon_90.228%
AEHB01001395_LSU_Basidiomycota_unclassified_Phanerochaete~chrysosporium_86.364%
Phanerochaete~carnosa~HHB~10118~sp_86.319%
AJ583427_LSU_Basidiomycota_Coniophoraceae_Coniophora~marmorata_86.039%
AYOR01000592_LSU_Basidiomycota_Hymenochaetaceae_Phellinus~noxius~OVT~YTM~97_88.852%
Armillaria~nabsnona_86.047%
AYUL01001070_LSU_Basidiomycota_unclassified_Phaeomarasmius~rimulincola_88.333%
Jaapiaceae_Jaapia~argillacea~MUCL~33604_86.319%
Pleurotaceae_Pleurotus~ostreatus~PC15_87.622%
Pleurotus~salmoneostramineus_86.645%
Sarcomyxa~serotina_85.621%
Hebeloma~mesophaeum_82.918%
DQ071726_LSU_Basidiomycota_Pluteaceae_Volvariella~caesiotincta_83.394%
Melanoleuca~melaleuca_86.486%
Phaeolepiota~aurea_86.799%
Hypholoma~radicosum_83.502%
DQ071695_LSU_Basidiomycota_unclassified_Panaeolus~acuminatus_83.66%
Basidiomycota_unclassified_Gloeophyllum~abietinum_90.26%
region~1460.5012_LSU_Basidiomycota_unclassified_Grifola~frondosa_86.039%
GBYM01000041_LSU_Basidiomycota_unclassified_Grifola~frondosa_86.039%
GBYM01000109_LSU_Basidiomycota_Coriolaceae_Coriolopsis~gallica
GBYM01000028_LSU_Basidiomycota_Coriolaceae_Coriolopsis~gallica
GBYM01000093_LSU_Basidiomycota_Coriolaceae_Coriolopsis~gallica
GBYM01000130_LSU_Basidiomycota_Coriolaceae_Coriolopsis~gallica
GBYM01000094_LSU_Basidiomycota_Coriolaceae_Coriolopsis~gallica
GBYM01000015_LSU_Basidiomycota_Coriolaceae_Coriolopsis~gallica
GBYM01000074_LSU_Basidiomycota_Coriolaceae_Coriolopsis~gallica
AJ583430_LSU_Basidiomycota_unclassified_Amyloporia~xantha
NEDQ01001104_LSU_Basidiomycota_unclassified_Postia~placenta~MAD~698~R
DQ208421_LSU_Basidiomycota_unclassified_Fomes~fomentarius
MPVS01000265_LSU_Basidiomycota_unclassified_Fomitopsis~pinicola
MJIM01000133_LSU_Basidiomycota_unclassified_Fomitopsis~palustris
DQ208412_LSU_Basidiomycota_unclassified_Ganoderma~lucidum
MARA01000070_LSU_Basidiomycota_unclassified_Polyporus~brumalis
CP019375~region~3331080.3334822_LSU_Basidiomycota_unclassified_Trametes
CP019375~region~3321230.3324964_LSU_Basidiomycota_unclassified_Trametes
CP019375~region~3291512.3295264_LSU_Basidiomycota_unclassified_Trametes
CP019375~region~3391046.3394441_LSU_Basidiomycota_unclassified_Trametes
CP019375~region~3438015.3441749_LSU_Basidiomycota_unclassified_Trametes
CP019375~region~3349101.3352835_LSU_Basidiomycota_unclassified_Trametes
CP019375_LSU_Basidiomycota_unclassified_Trametes
CP019375~region~3301380.3305114_LSU_Basidiomycota_unclassified_Trametes
CP019375~region~3408775.3412049_LSU_Basidiomycota_unclassified_Trametes
CP019375~region~3447865.3451599_LSU_Basidiomycota_unclassified_Trametes~hirsuta
AOTI010832126_LSU_Basidiomycota_unclassified_Triticum~urartu_87.286%
ADFJ01004862_LSU_Basidiomycota_unclassified_Selaginella~moellendorffii_87.622%
LSVH01000344_LSU_Basidiomycota_Mrakiaceae_Phaffia~rhodozyma_90.909%
GDMN01020310_LSU_Basidiomycota_Coriolaceae_Rigidoporus~microporus_90.228%
ANON12_OTU_214
JUBC01000926_LSU_Basidiomycota_Mrakiaceae_Mrakia~frigida_89.935%
BBPU01000028_LSU_Basidiomycota_Mrakiaceae_Mrakia~blollopis_89.935%
ANON19_OTU_272
EU909344_LSU_Basidiomycota_Botryobasidiaceae_Botryobasidium~subcoronatum_90.26%
AYEP01001617_LSU_Basidiomycota_unclassified_Botryobasidium~botryosum~FD~172~SS1_86.928%
ANON23_OTU_356
ANON25_OTU_466
ANON17_OTU_263
ANON2_OTU_29
ANON16_OTU_258
ANON22_OTU_349
100
17
78
92
99
95
51
51
3
2
28
21
37
100
100
67
4
8
35
0.05