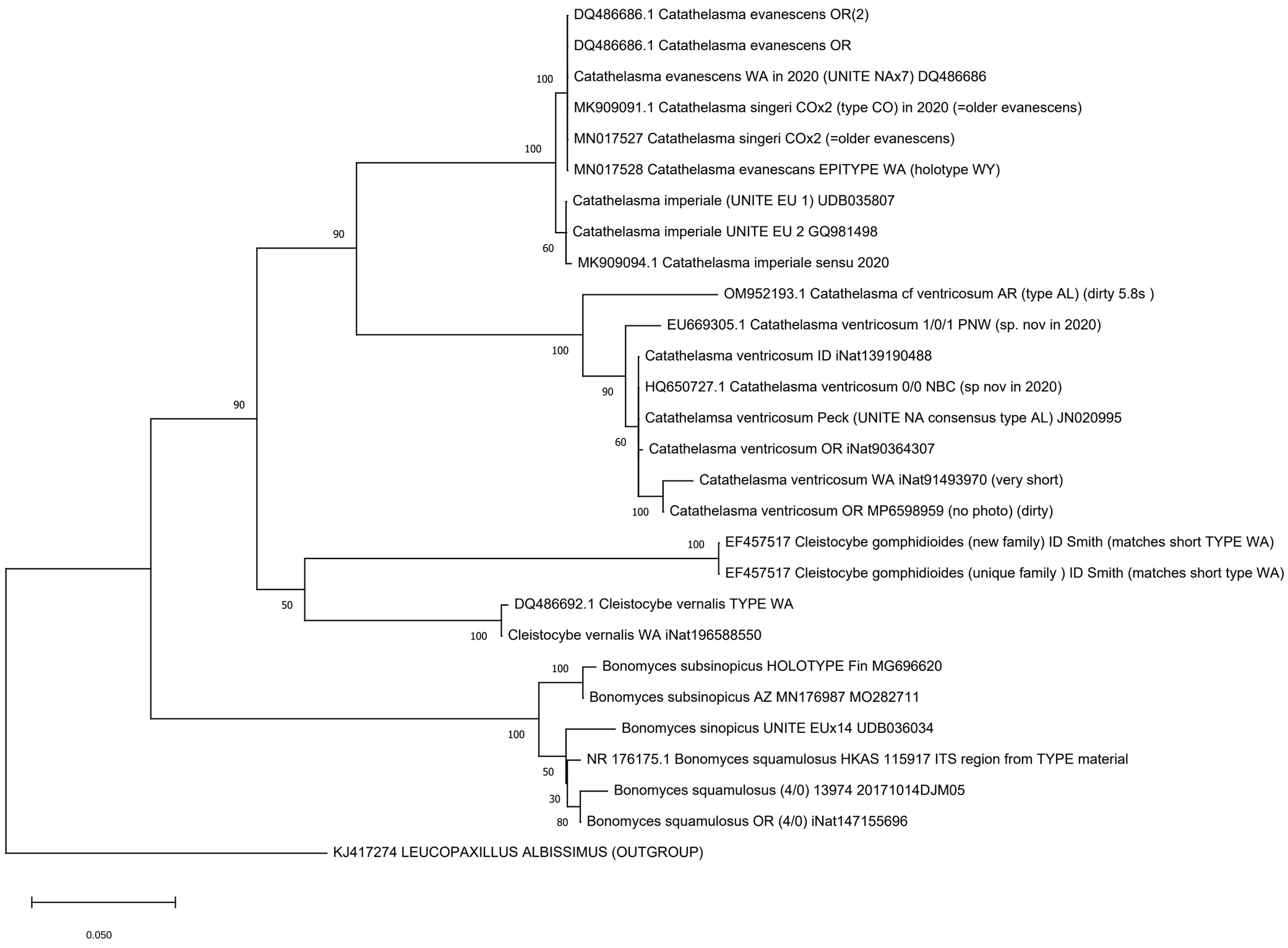
DQ486686.1 Catathelasma evanescens OR(2)
DQ486686.1 Catathelasma evanescens OR
Catathelasma evanescens WA in 2020 (UNITE NAx7) DQ486686
MK909091.1 Catathelasma singeri COx2 (type CO) in 2020 (=older evanescens)
MN017527 Catathelasma singeri COx2 (=older evanescens)
MN017528 Catathelasma evanescans EPITYPE WA (holotype WY)
Catathelasma imperiale (UNITE EU 1) UDB035807
Catathelasma imperiale UNITE EU 2 GQ981498
MK909094.1 Catathelasma imperiale sensu 2020
OM952193.1 Catathelasma cf ventricosum AR (type AL) (dirty 5.8s)
EU669305.1 Catathelasma ventricosum 1/0/1 PNW (sp. nov in 2020)
Catathelasma ventricosum ID iNat139190488
HQ650727.1 Catathelasma ventricosum 0/0 NBC (sp nov in 2020)
Catathelamsa ventricosum Peck (UNITE NA consensus type AL) JN020995
Catathelasma ventricosum OR iNat90364307
Catathelasma ventricosum WA iNat91493970 (very short)
Catathelasma ventricosum OR MP6598959 (no photo) (dirty)
EF457517 Cleistocybe gomphidioides (new family) ID Smith (matches short TYPE WA)
EF457517 Cleistocybe gomphidioides (unique family) ID Smith (matches short type WA)
DQ486692.1 Cleistocybe vernalis TYPE WA
Cleistocybe vernalis WA iNat196588550
Bonomyces subsinopicus HOLOTYPE Fin MG696620
Bonomyces subsinopicus AZ MN176987 MO282711
Bonomyces sinopicus UNITE EUx14 UDB036034
NR 176175.1 Bonomyces squamulosus HKAS 115917 ITS region from TYPE material
Bonomyces squamulosus (4/0) 13974 20171014DJM05
Bonomyces squamulosus OR (4/0) iNat147155696
KJ417274 LEUCOPAXILLUS ALBISSIMUS (OUTGROUP)
100
100
90
100
90
60
90
100
90
60
100
100
50
100
100
100
50
30
80
0.050