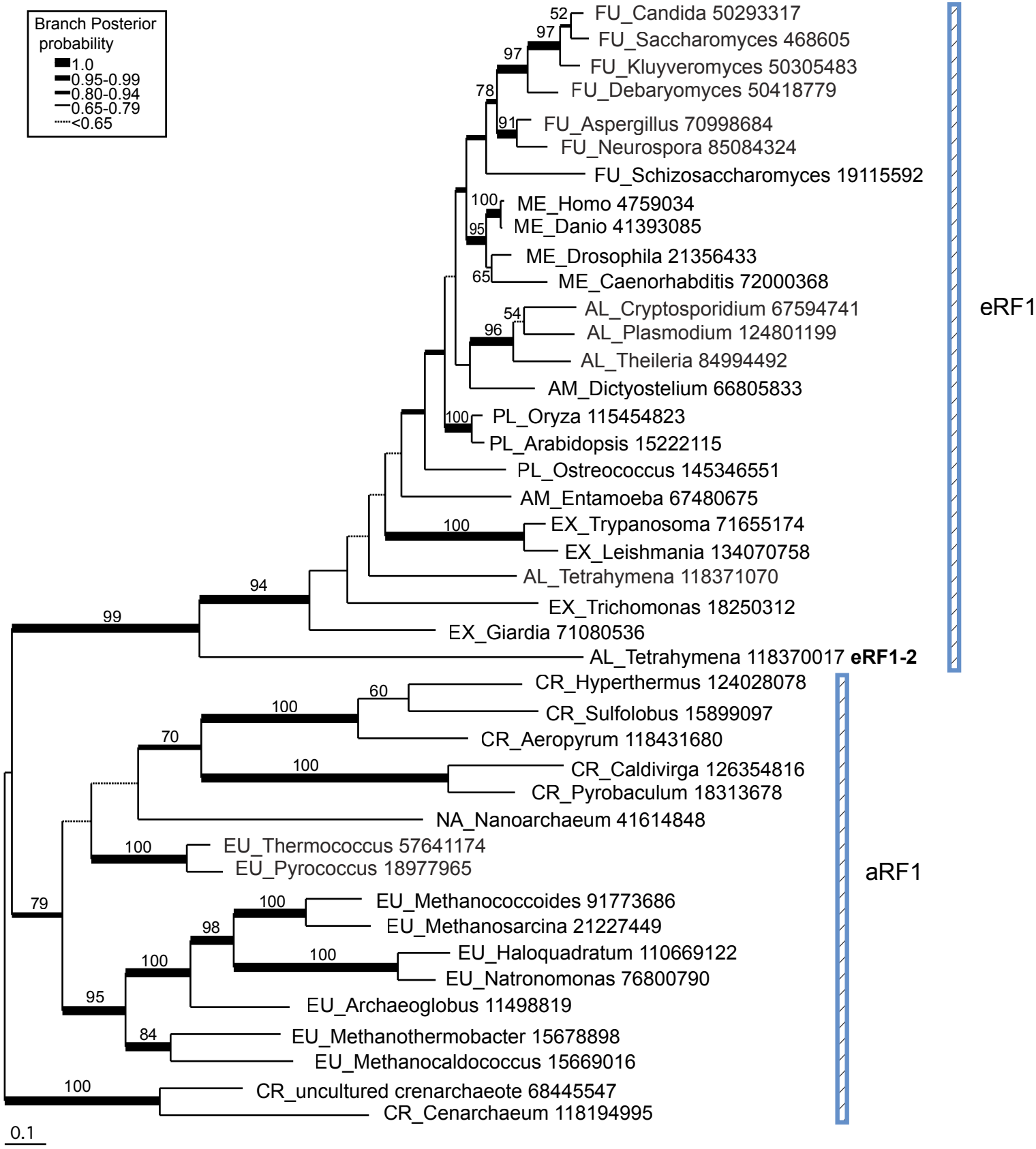
Branch Posterior probability
1.0
0.95-0.99
0.80-0.94
0.65-0.79
<0.65
FU_Candida 50293317
FU_Saccharomyces 468605
FU_Kluyveromyces 50305483
FU_Debaryomyces 50418779
FU_Aspergillus 70998684
FU_Neurospora 85084324
FU_Schizosaccharomyces 19115592
ME_Homo 4759034
ME_Danio 41393085
ME_Drosophila 21356433
ME_Caenorhabditis 72000368
AL_Cryptosporidium 67594741
AL_Plasmodium 124801199
AL_Theileria 84994492
AM_Dictyostelium 66805833
PL_Oryza 115454823
PL_Arabidopsis 15222115
PL_Ostreococcus 145346551
AM_Entamoeba 67480675
EX_Trypanosoma 71655174
EX_Leishmania 134070758
AL_Tetrahymena 118371070
EX_Trichomonas 18250312
EX_Giardia 71080536
AL_Tetrahymena 118370017 eRF1-2
eRF1
CR_Hyperthermus 124028078
CR_Sulfolobus 15899097
CR_Aeropyrum 118431680
CR_Caldivirga 126354816
CR_Pyrobaculum 18313678
NA_Nanoarchaeum 41614848
EU_Thermococcus 57641174
EU_Pyrococcus 18977965
EU_Methanococcoides 91773686
EU_Methanosarcina 21227449
EU_Haloquadratum 110669122
EU_Natronomonas 76800790
EU_Archaeoglobus 11498819
EU_Methanothermobacter 15678898
EU_Methanocaldococcus 15669016
CR_uncultured crenarchaeote 68445547
CR_Cenarchaeum 118194995
aRF1
0.1