Cyanobacteria

Euryarchaeota

Chloroflexi

TACK+A

CPR

DPANN

Actinobacteria

Proteobacteria

Caldatribacteriota

Deltaproteobacteria

Poribacteria

Omnitrophica

Verrucomicrobia

Aminicenantes

Planctomycetes

Cloacimonadota

FBC

0.2

Bootstrap inferred with SH−aLRT test
· 99.9 − 100
· 90 − 99.9
< 90

basal branch
archaeal BS MAG
bacterial BS MAG
archaeal reference genome
bacterial reference genome