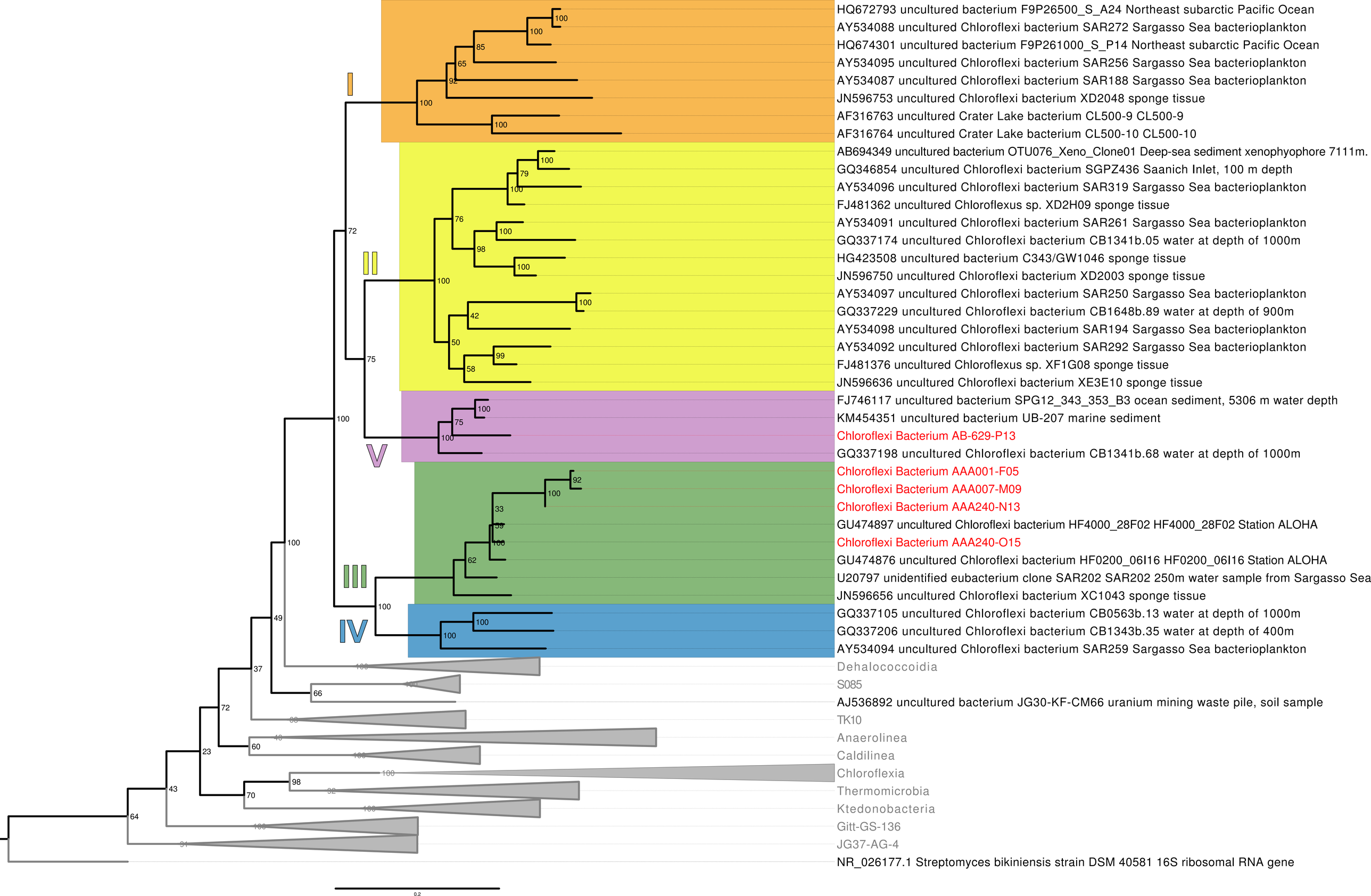
HQ672793 uncultured bacterium F9P26500_S_A24 Northeast subarctic Pacific Ocean
AY534088 uncultured Chloroflexi bacterium SAR272 Sargasso Sea bacterioplankton
HQ674301 uncultured bacterium F9P261000_S_P14 Northeast subarctic Pacific Ocean
AY534095 uncultured Chloroflexi bacterium SAR256 Sargasso Sea bacterioplankton
AY534087 uncultured Chloroflexi bacterium SAR188 Sargasso Sea bacterioplankton
JN596753 uncultured Chloroflexi bacterium XD2048 sponge tissue
AF316763 uncultured Crater Lake bacterium CL500-9 CL500-9
AF316764 uncultured Crater Lake bacterium CL500-10 CL500-10
AB694349 uncultured bacterium OTU076_Xeno_Clone01 Deep-sea sediment xenophyophore 7111m.
GQ346854 uncultured Chloroflexi bacterium SGPZ436 Saanich Inlet, 100 m depth
AY534096 uncultured Chloroflexi bacterium SAR319 Sargasso Sea bacterioplankton
FJ481362 uncultured Chloroflexus sp. XD2H09 sponge tissue
AY534091 uncultured Chloroflexi bacterium SAR261 Sargasso Sea bacterioplankton
GQ337174 uncultured Chloroflexi bacterium CB1341b.05 water at depth of 1000m
HG423508 uncultured bacterium C343/GW1046 sponge tissue
JN596750 uncultured Chloroflexi bacterium XD2003 sponge tissue
AY534097 uncultured Chloroflexi bacterium SAR250 Sargasso Sea bacterioplankton
GQ337229 uncultured Chloroflexi bacterium CB1648b.89 water at depth of 900m
AY534098 uncultured Chloroflexi bacterium SAR194 Sargasso Sea bacterioplankton
AY534092 uncultured Chloroflexi bacterium SAR292 Sargasso Sea bacterioplankton
FJ481376 uncultured Chloroflexus sp. XF1G08 sponge tissue
JN596636 uncultured Chloroflexi bacterium XE3E10 sponge tissue
FJ746117 uncultured bacterium SPG12_343_353_B3 ocean sediment, 5306 m water depth
KM454351 uncultured bacterium UB-207 marine sediment
Chloroflexi Bacterium AB-629-P13
GQ337198 uncultured Chloroflexi bacterium CB1341b.68 water at depth of 1000m
Chloroflexi Bacterium AAA001-F05
Chloroflexi Bacterium AAA007-M09
Chloroflexi Bacterium AAA240-N13
GU474897 uncultured Chloroflexi bacterium HF4000_28F02 HF4000_28F02 Station ALOHA
Chloroflexi Bacterium AAA240-O15
GU474876 uncultured Chloroflexi bacterium HF0200_06I16 HF0200_06I16 Station ALOHA
U20797 unidentified eubacterium clone SAR202 SAR202 250m water sample from Sargasso Sea
JN596656 uncultured Chloroflexi bacterium XC1043 sponge tissue
GQ337105 uncultured Chloroflexi bacterium CB0563b.13 water at depth of 1000m
GQ337206 uncultured Chloroflexi bacterium CB1343b.35 water at depth of 400m
AY534094 uncultured Chloroflexi bacterium SAR259 Sargasso Sea bacterioplankton
Dehalococcoidia
S085
AJ536892 uncultured bacterium JG30-KF-CM66 uranium mining waste pile, soil sample
TK10
Anaerolinea
Caldilinea
Chloroflexia
Thermomicrobia
Ktedonobacteria
Gitt-GS-136
JG37-AG-4
NR_026177.1 Streptomyces bikiniensis strain DSM 40581 16S ribosomal RNA gene
I
II
III
IV
V
0.2