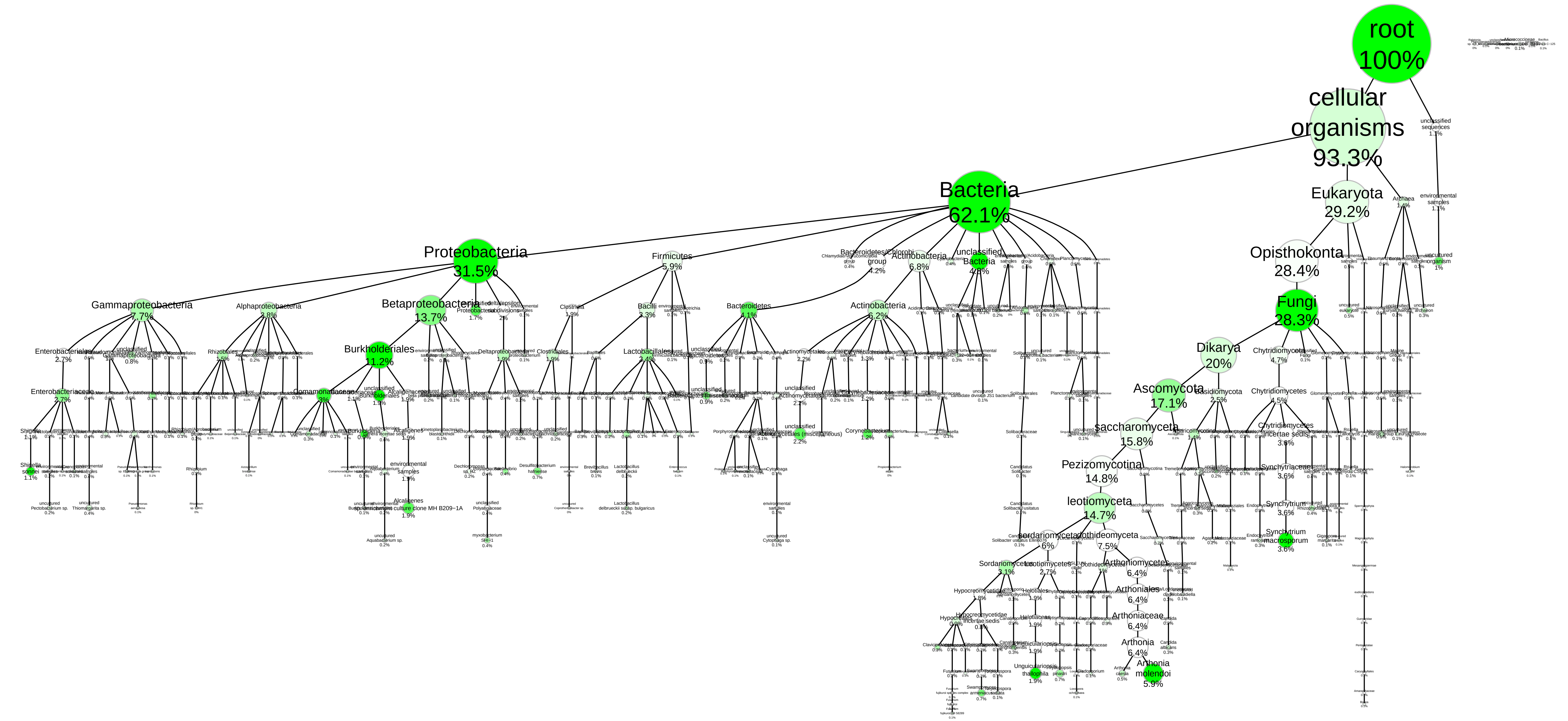

root
100%
cellular organisms
93.3%
unclassified sequences
1.1%
Bacteria
62.1%
Eukaryota
29.2%
Archaea
1.4%
environmental samples
1.1%
uncultured organism
1%
Proteobacteria
31.5%
Firmicutes
5.9%
Bacteroidetes/Chlorobi group
4.2%
Actinobacteria
6.8%
unclassified Bacteria
4.8%
Opisthokonta
28.4%
Fungi
28.3%
Dikarya
20%
Ascomycota
17.1%
Basidiomycota
2.5%
Chytridiomycota
4.7%
Chytridiomycetes
4.5%
Chytridiomycetes incertae sedis
3.6%
Synchytriaceae
3.6%
Synchytrium
3.6%
Synchytrium macrosporum
3.6%
saccharomyceta
15.8%
Pezizomycotina
14.8%
leotiomyceta
14.7%
sordariomyceta
6%
dothideomyceta
7.5%
Sordariomycetes
3.1%
Leotiomycetes
2.7%
Arthoniomycetes
6.4%
Arthoniales
6.4%
Arthoniaceae
6.4%
Arthonia
6.4%
Arthonia molendoi
5.9%
Hypocreomycetidae
1.8%
Helotiales
1.9%
Unguiculariopsis
1.9%
Unguiculariopsis thallophila
1.9%
Gammaproteobacteria
7.7%
Alphaproteobacteria
3.8%
Betaproteobacteria
13.7%
Burkholderiales
11.2%
Comamonadaceae
3%
Alcaligenaceae
1.9%
Alcaligenes
1.9%
environmental samples
1.9%
Alcaligenes sp. enrichment culture clone MH B209-1A
1.9%
unclassified Burkholderiales
1.9%
Enterobacteriales
2.7%
Enterobacteriaceae
2.7%
Shigella
1.1%
Shigella sonnei
1.1%
Rhizobiales
1.5%
Deltaproteobacteria
1.9%
Clostridia
1.9%
Clostridiales
1.9%
Bacilli
3.3%
Lactobacillales
2.4%
Bacteroidetes
4.1%
Actinobacteria
6.2%
Actinomycetales
2.2%
unclassified Actinomycetales
2.2%
unclassified Actinomycetales (miscellaneous)
2.2%