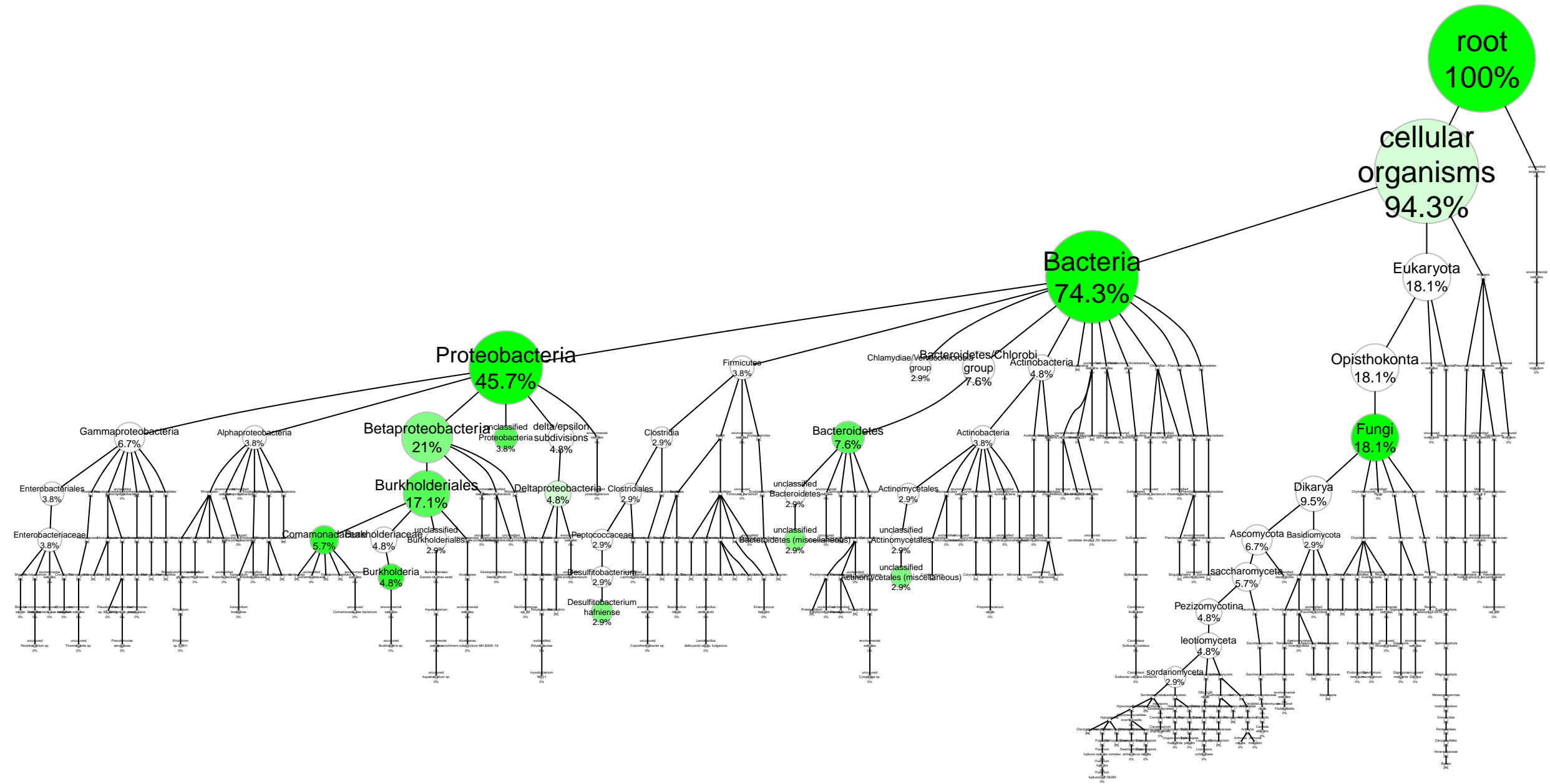
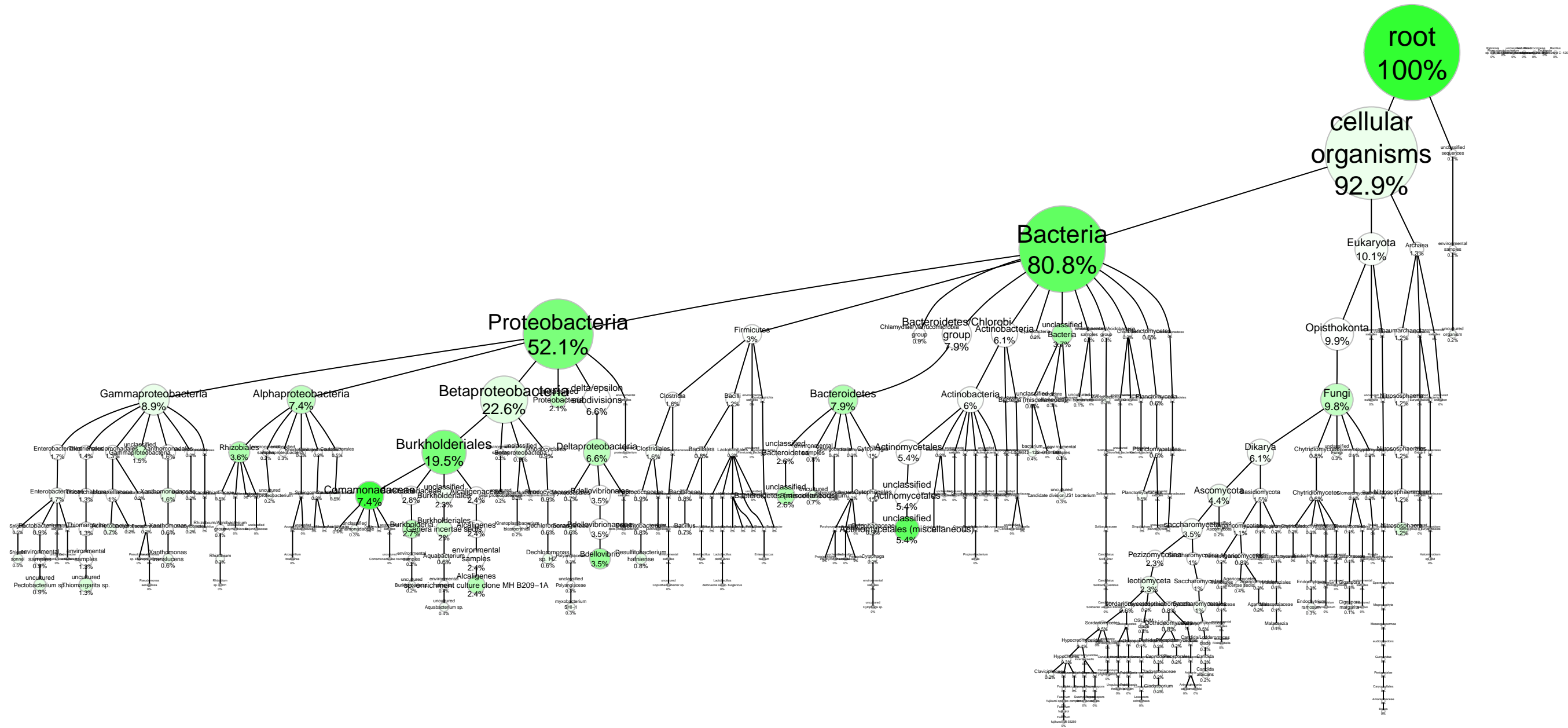
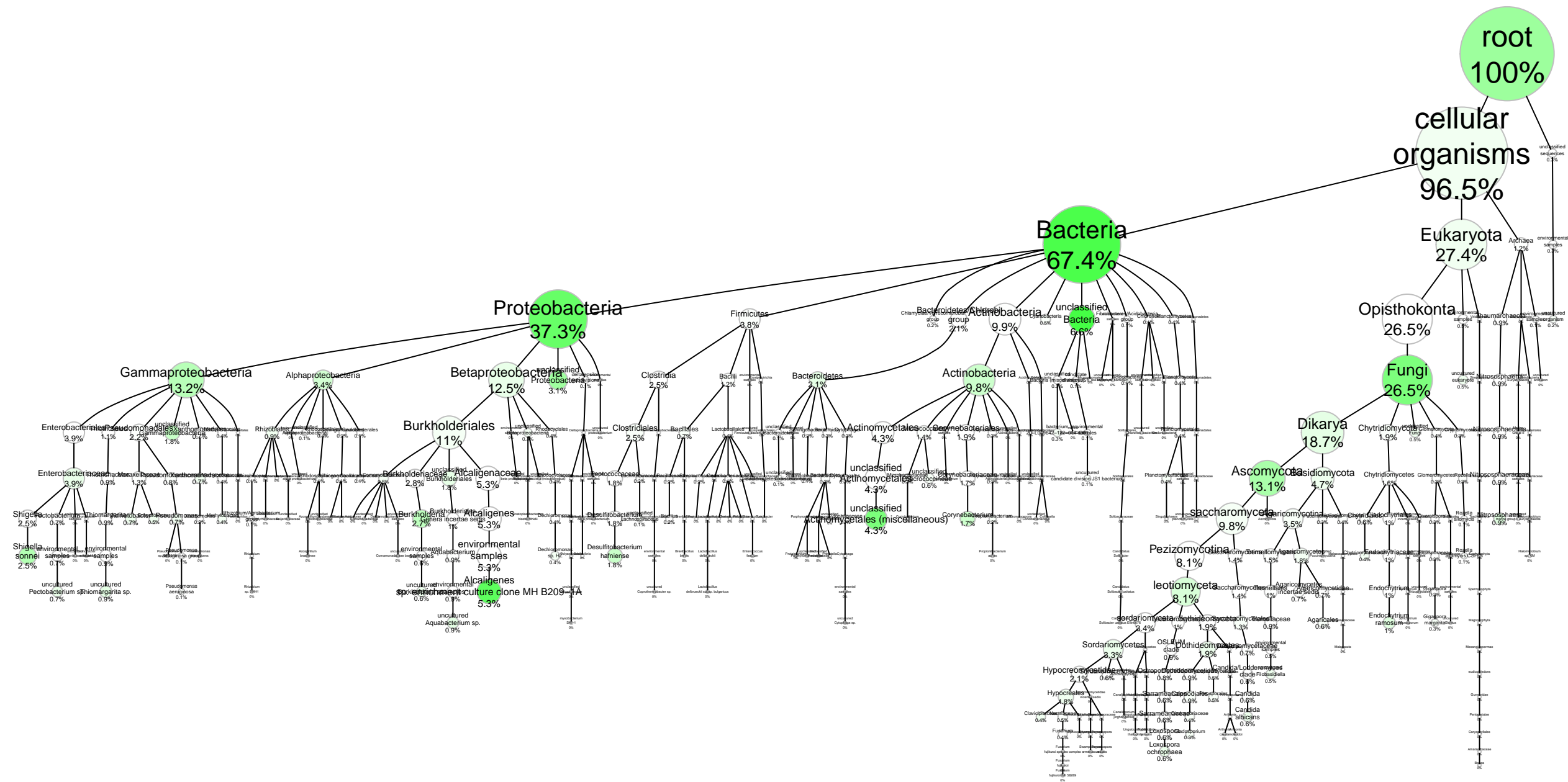


[illegible]

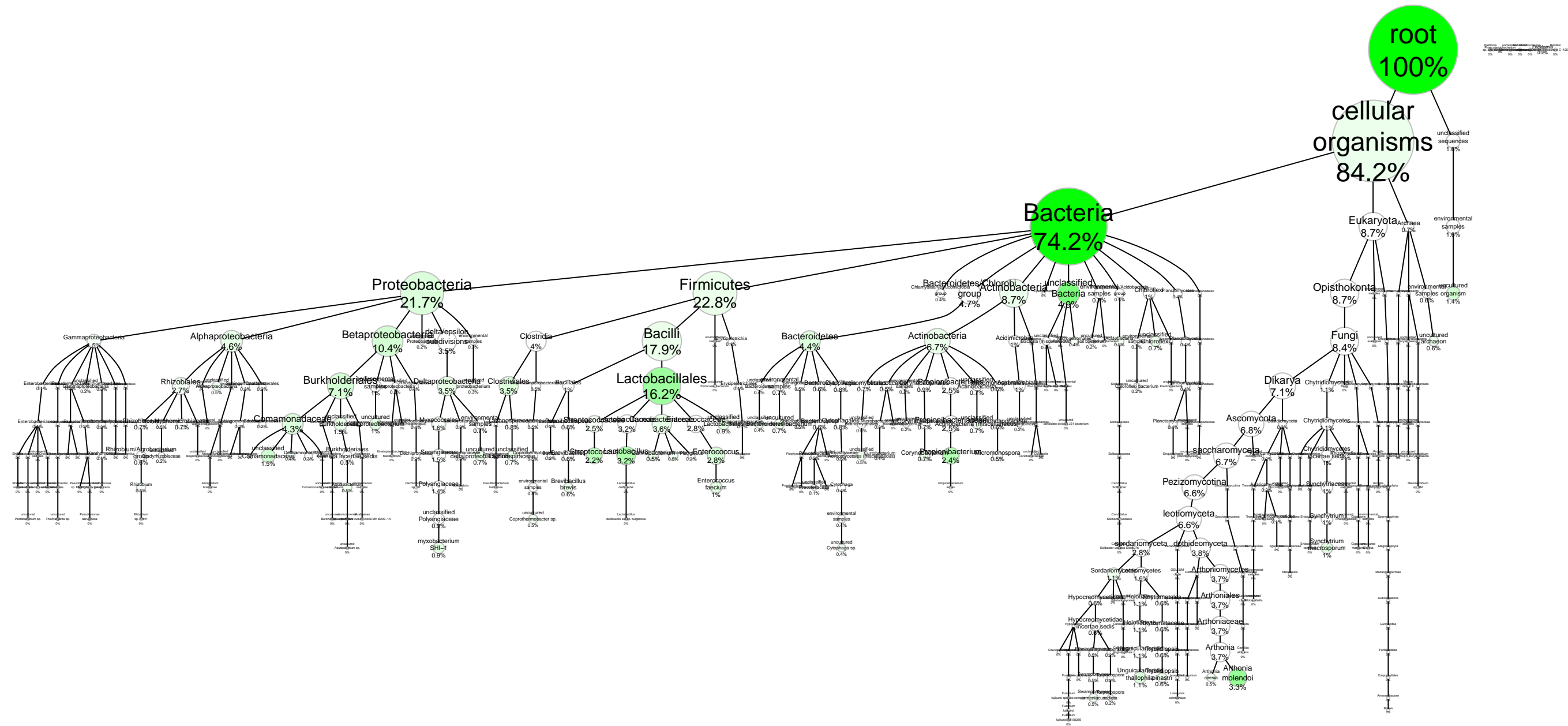
sp. C-125 0% 0% 0% 0% 0% 0% 0%



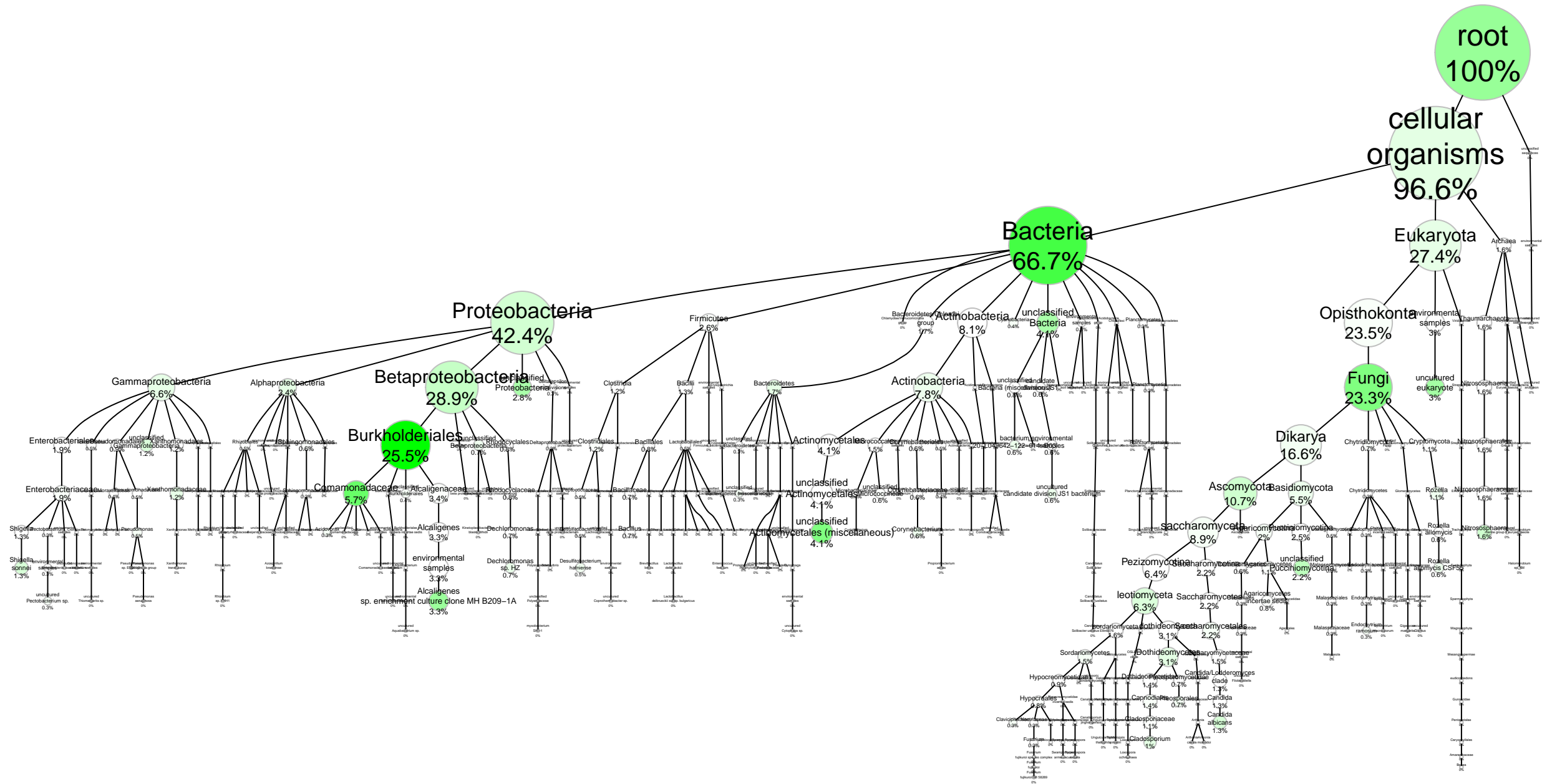
Micrococcineae



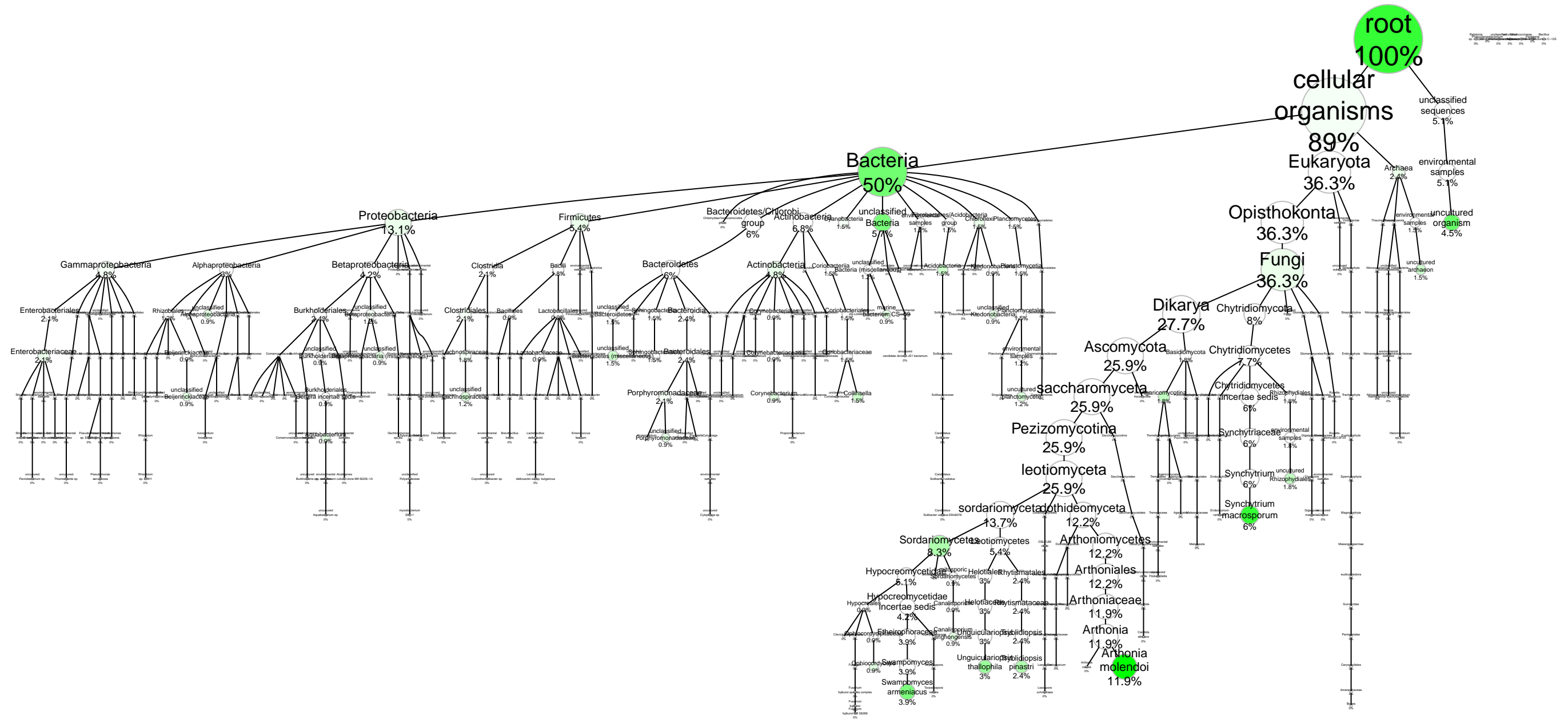
Phaeocystis unclassified, Microcystis, Bacillus
Phaeocystis unclassified, Microcystis, Bacillus
sp. cdk, Microcystis, Bacillus, Bacillus C-125

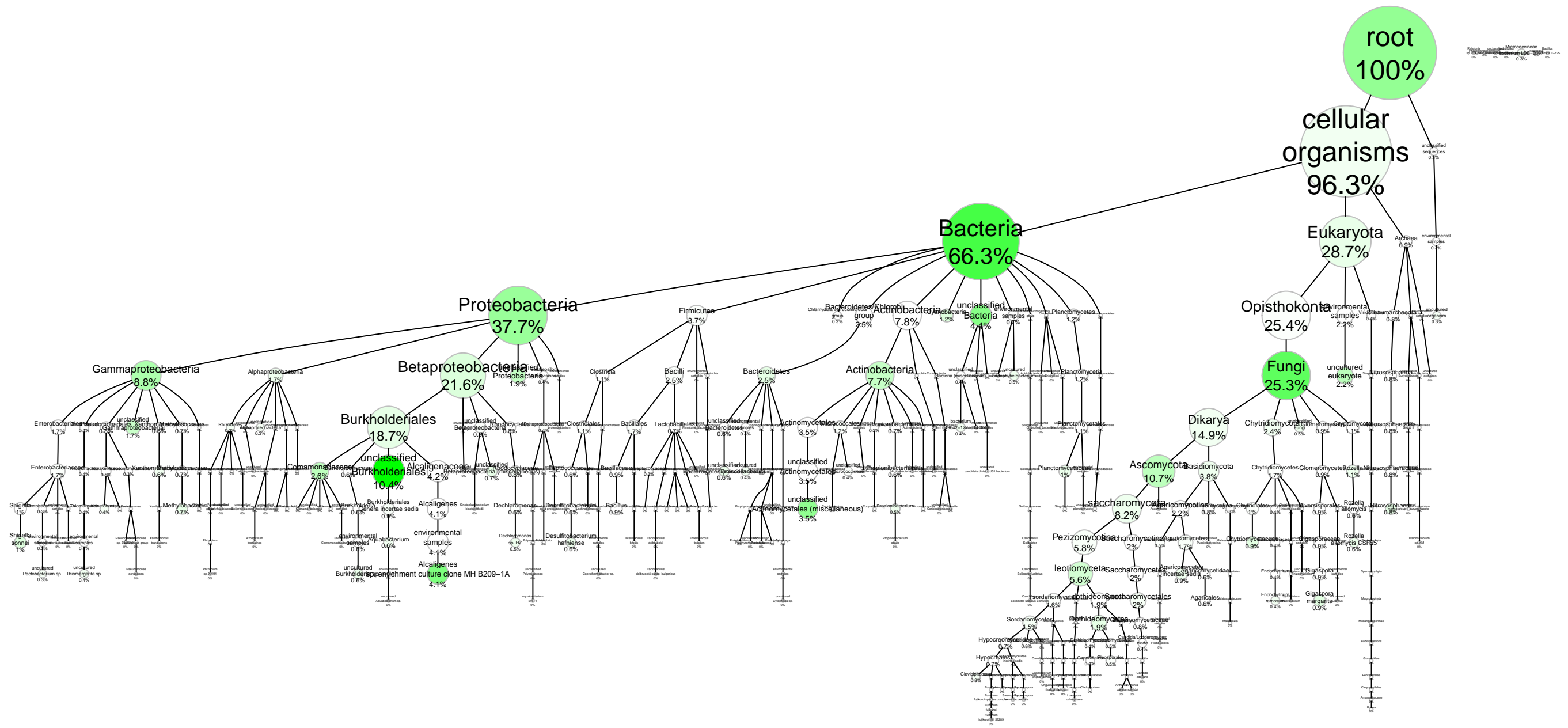


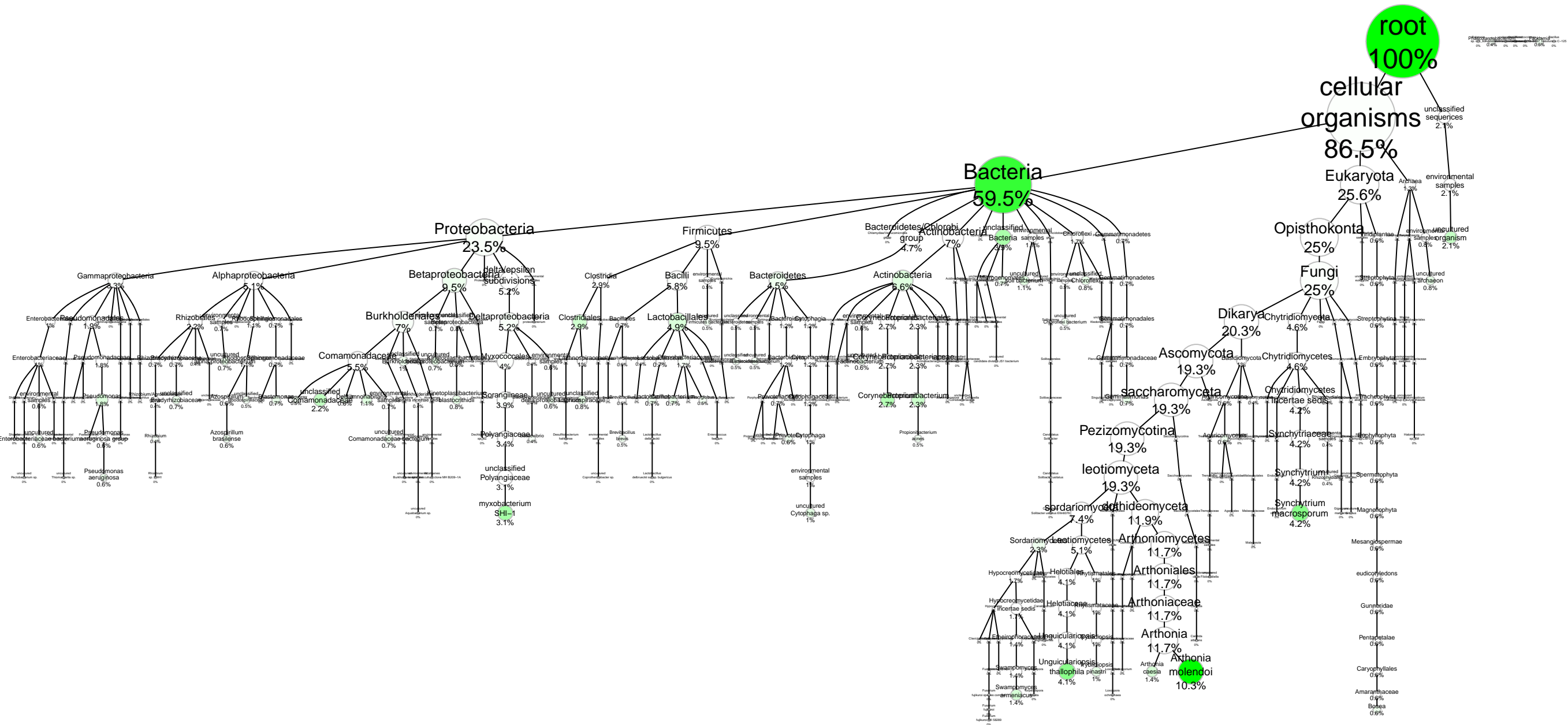
Genus	Relative Abundance (%)
Rhizobium	0.0%
uncultured bacterium	0.0%
Microcococcus	0.5%
Bacillus	0.5%
Bifidobacterium	0.0%



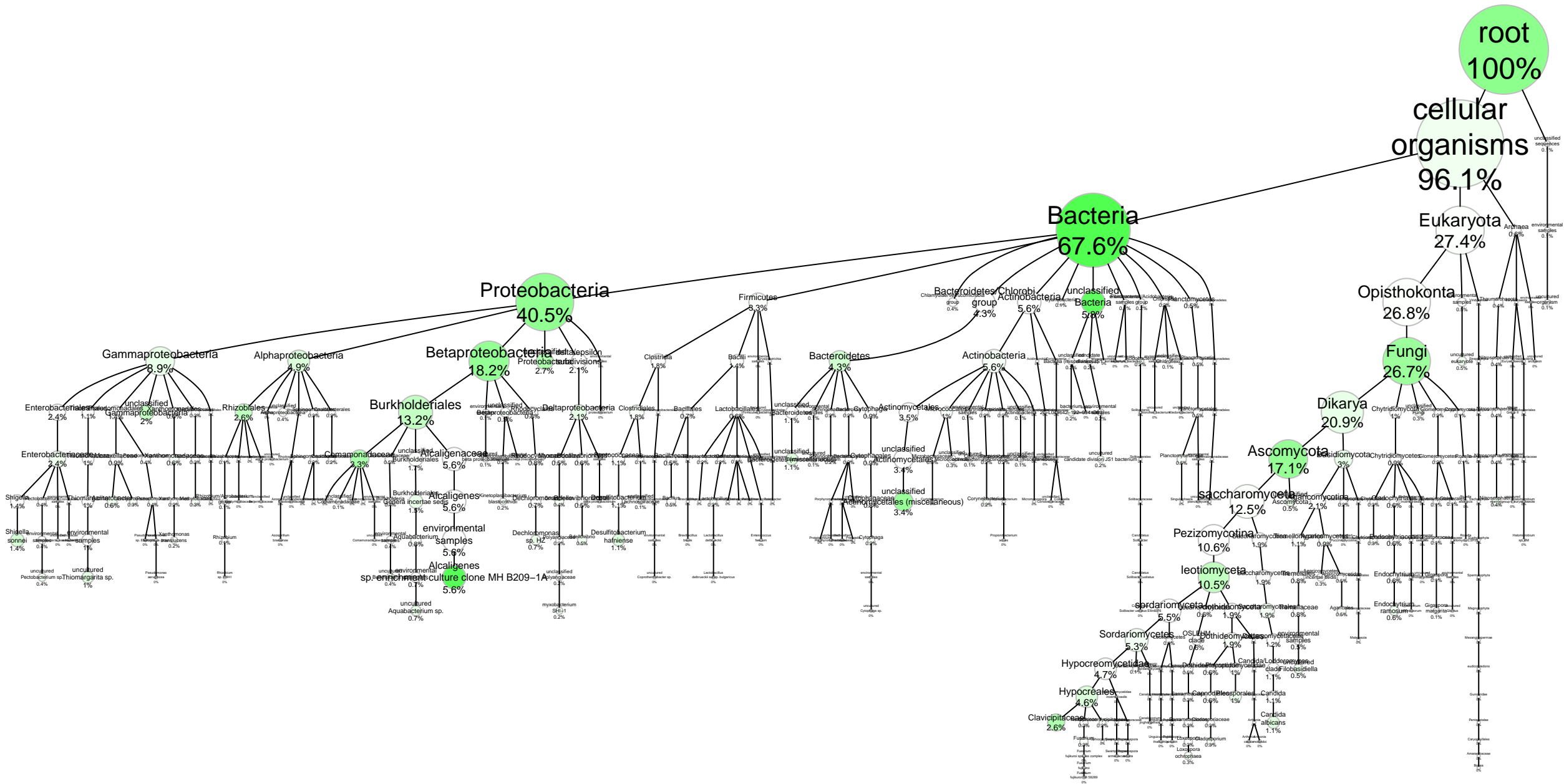
sample_3ml_Bioo_CGATGT_R1_001_fastq (total reads: 336)

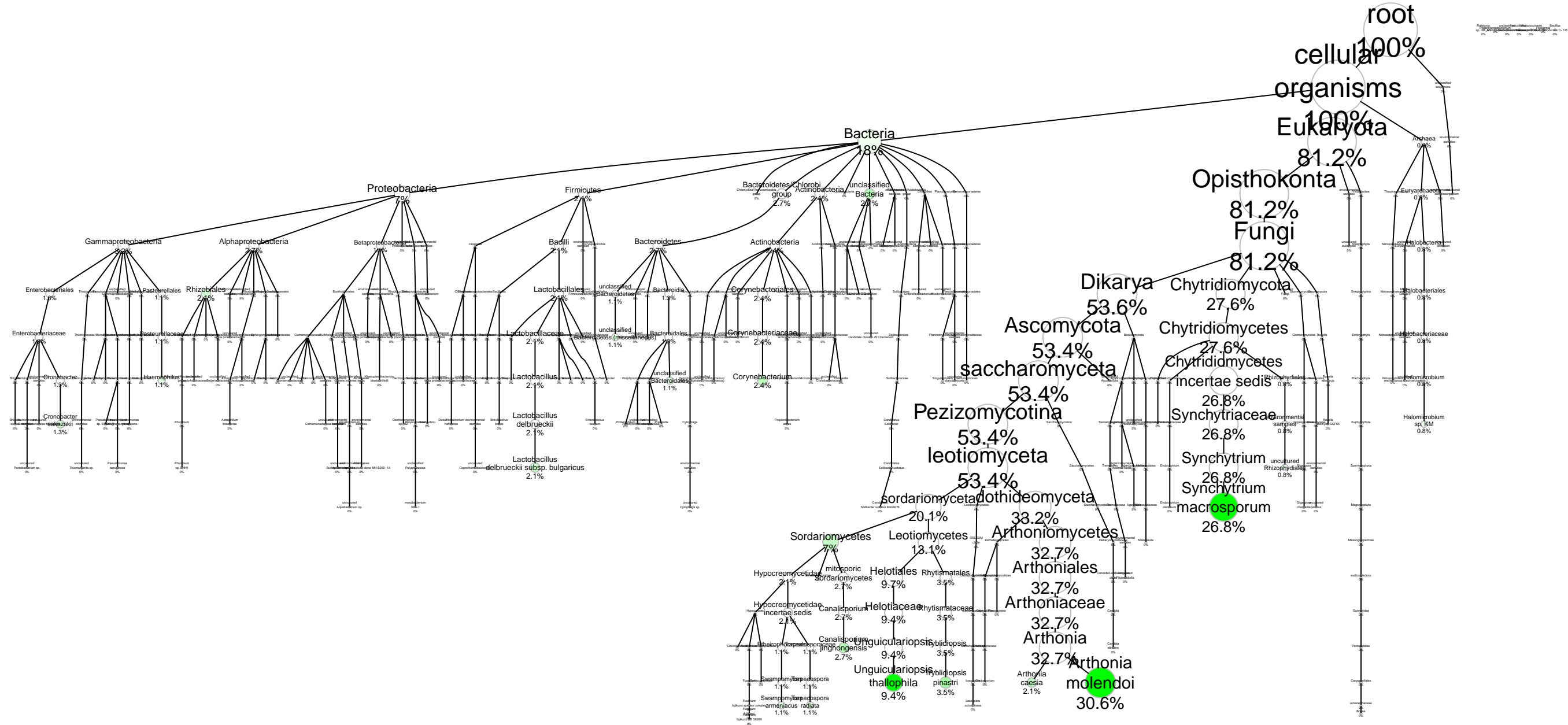




[illegible]

Micrococcineae Bacillus
0% 0% 0% 0% 0.3% 0.2%



[illegible]

sample_5ml_Illumina_CTCAGA_R1_001_fastq (total reads: 358)

