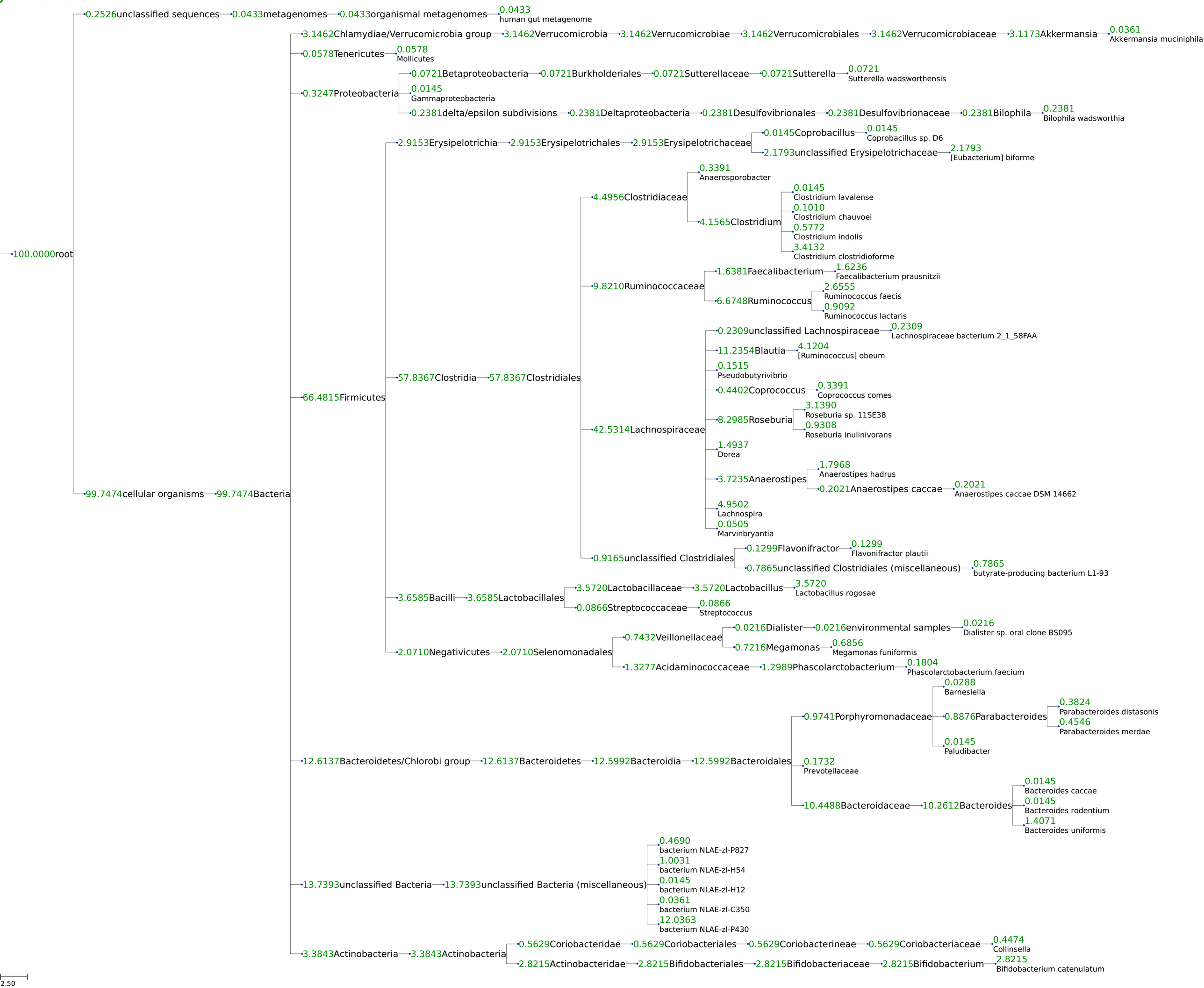
Jun 20 2015


100.0000 root
0.2526 unclassified sequences
0.0433 metagenomes
0.0433 organismal metagenomes
0.0433 human gut metagenome
99.7474 cellular organisms
99.7474 Bacteria
3.1462 Chlamydiae/Verrucomicrobia group
3.1462 Verrucomicrobia
3.1462 Verrucomicrobiae
3.1462 Verrucomicrobiales
3.1462 Verrucomicrobiaceae
3.1173 Akkermansia
0.0361 Akkermansia muciniphila
0.0578 Tenericutes
0.0578 Mollicutes
0.3247 Proteobacteria
0.0721 Betaproteobacteria
0.0721 Burkholderiales
0.0721 Sutterellaceae
0.0721 Sutterella
0.0721 Sutterella wadsworthensis
0.0145 Gammaproteobacteria
0.2381 delta/epsilon subdivisions
0.2381 Deltaproteobacteria
0.2381 Desulfovibrionales
0.2381 Desulfovibrionaceae
0.2381 Bilophila
0.2381 Bilophila wadsworthia
66.4815 Firmicutes
2.9153 Erysipelotrichia
2.9153 Erysipelotrichales
2.9153 Erysipelotrichaceae
0.0145 Coprobacillus
0.0145 Coprobacillus sp. D6
2.1793 unclassified Erysipelotrichaceae
2.1793 [Eubacterium] biforme
57.8367 Clostridia
57.8367 Clostridiales
4.4956 Clostridiaceae
0.3391 Anaerosporobacter
4.1565 Clostridium
0.0145 Clostridium lavalense
0.1010 Clostridium chauvoei
0.5772 Clostridium indolis
3.4132 Clostridium clostridioforme
9.8210 Ruminococcaceae
1.6381 Faecalibacterium
1.6236 Faecalibacterium prausnitzii
6.6748 Ruminococcus
2.6555 Ruminococcus faecis
0.9092 Ruminococcus lactaris
42.5314 Lachnospiraceae
0.2309 unclassified Lachnospiraceae
0.2309 Lachnospiraceae bacterium 2_1_58FAA
11.2354 Blautia
4.1204 [Ruminococcus] obeum
0.1515 Pseudobutyrivibrio
0.4402 Coprococcus
0.3391 Coprococcus comes
8.2985 Roseburia
3.1390 Roseburia sp. 11SE38
0.9308 Roseburia inulinivorans
1.4937 Dorea
3.7235 Anaerostipes
1.7968 Anaerostipes hadrus
0.2021 Anaerostipes caccae
0.2021 Anaerostipes caccae DSM 14662
4.9502 Lachnospira
0.0505 Marvinbryantia
0.9165 unclassified Clostridiales
0.1299 Flavonifractor
0.1299 Flavonifractor plautii
0.7865 unclassified Clostridiales (miscellaneous)
0.7865 butyrate-producing bacterium L1-93
3.6585 Bacilli
3.6585 Lactobacillales
3.5720 Lactobacillaceae
3.5720 Lactobacillus
3.5720 Lactobacillus rogosae
0.0866 Streptococcaceae
0.0866 Streptococcus
2.0710 Negativicutes
2.0710 Selenomonadales
0.7432 Veillonellaceae
0.0216 Dialister
0.0216 environmental samples
0.0216 Dialister sp. oral clone BS095
0.7216 Megamonas
0.6856 Megamonas funiformis
1.3277 Acidaminococcaceae
1.2989 Phascolarctobacterium
0.1804 Phascolarctobacterium faecium
12.6137 Bacteroidetes/Chlorobi group
12.6137 Bacteroidetes
12.5992 Bacteroidia
12.5992 Bacteroidales
0.9741 Porphyromonadaceae
0.0288 Barnesiella
0.8876 Parabacteroides
0.3824 Parabacteroides distasonis
0.4546 Parabacteroides merdae
0.0145 Paludibacter
0.1732 Prevotellaceae
10.4488 Bacteroidaceae
10.2612 Bacteroides
0.0145 Bacteroides caccae
0.0145 Bacteroides rodentium
1.4071 Bacteroides uniformis
13.7393 unclassified Bacteria
13.7393 unclassified Bacteria (miscellaneous)
0.4690 bacterium NLAE-zl-P827
1.0031 bacterium NLAE-zl-H54
0.0145 bacterium NLAE-zl-H12
0.0361 bacterium NLAE-zl-C350
12.0363 bacterium NLAE-zl-P430
3.3843 Actinobacteria
3.3843 Actinobacteria
0.5629 Coriobacteridae
0.5629 Coriobacteriales
0.5629 Coriobacterineae
0.5629 Coriobacteriaceae
0.4474 Collinsella
2.8215 Actinobacteridae
2.8215 Bifidobacteriales
2.8215 Bifidobacteriaceae
2.8215 Bifidobacterium
2.8215 Bifidobacterium catenulatum
2.50