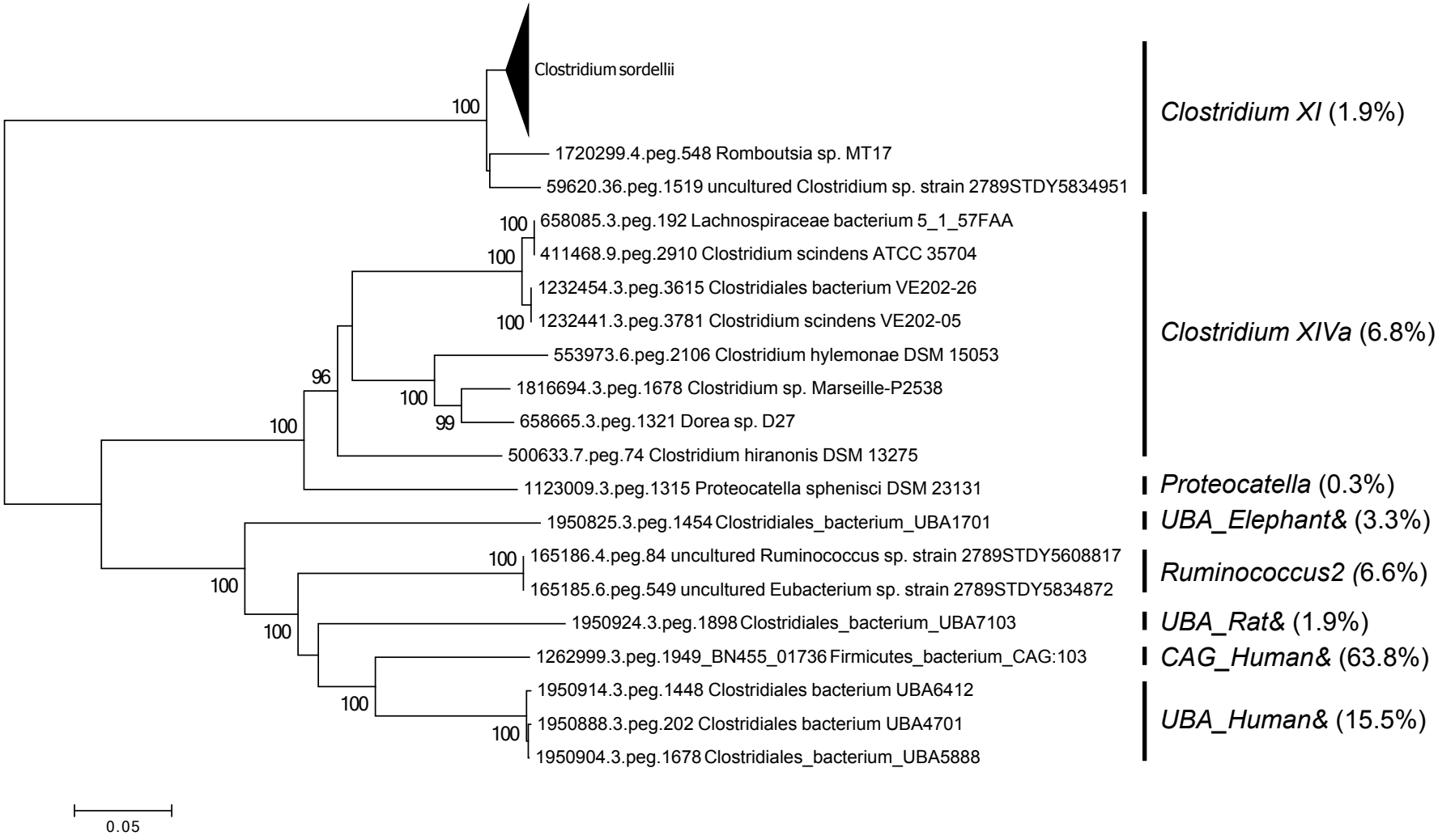
Clostridium sordellii
100
1720299.4.peg.548 Romboutsia sp. MT17
59620.36.peg.1519 uncultured Clostridium sp. strain 2789STDY5834951
Clostridium XI (1.9%)
100
658085.3.peg.192 Lachnospiraceae bacterium 5_1_57FAA
100
411468.9.peg.2910 Clostridium scindens ATCC 35704
1232454.3.peg.3615 Clostridiales bacterium VE202-26
100
1232441.3.peg.3781 Clostridium scindens VE202-05
553973.6.peg.2106 Clostridium hylemonae DSM 15053
96
100
1816694.3.peg.1678 Clostridium sp. Marseille-P2538
100
99
658665.3.peg.1321 Dorea sp. D27
500633.7.peg.74 Clostridium hiranonis DSM 13275
Clostridium XIVa (6.8%)
1123009.3.peg.1315 Proteocatella sphenisci DSM 23131
Proteocatella (0.3%)
1950825.3.peg.1454 Clostridiales_bacterium_UBA1701
UBA_Elephant& (3.3%)
100
165186.4.peg.84 uncultured Ruminococcus sp. strain 2789STDY5608817
100
165185.6.peg.549 uncultured Eubacterium sp. strain 2789STDY5834872
Ruminococcus2 (6.6%)
100
1950924.3.peg.1898 Clostridiales_bacterium_UBA7103
UBA_Rat& (1.9%)
1262999.3.peg.1949_BN455_01736 Firmicutes_bacterium_CAG:103
CAG_Human& (63.8%)
100
1950914.3.peg.1448 Clostridiales bacterium UBA6412
100
1950888.3.peg.202 Clostridiales bacterium UBA4701
1950904.3.peg.1678 Clostridiales_bacterium_UBA5888
UBA_Human& (15.5%)
0.05