



- Otu340_Actinomycetales100%
- Otu337_Verrucomicrobiaceae100%
- Otu334_Porphyromonadaceae97%
- Otu331_Lachnospiraceae100%
- Otu328_ClostridialesIncertaeSedisXI99%
- Otu325_Sphingobacteriaceae100%
- Otu322_Lachnospiraceae69%
- Otu319_Burkholderiaceae100%
- Otu316_Rhodocyclaceae14%
- Otu313_Lachnospiraceae90%
- Otu310_Lachnospiraceae100%
- Otu307_Lachnospiraceae78%
- Otu304_Ruminococcaceae90%
- Otu301_Lactobacillaceae91%
- Otu298_Lachnospiraceae96%
- Otu295_Bradyrhizobiaceae99%
- Otu292_Luteivirga1%
- Otu289_Lachnospiraceae100%
- Otu286_Lachnospiraceae100%
- Otu283_Lachnospiraceae80%
- Otu280_Lactobacillaceae90%
- Otu277_Lachnospiraceae100%
- Otu274_Lachnospiraceae97%
- Otu271_Lachnospiraceae100%
- Otu268_Actinomycetales100%
- Otu265_Lachnospiraceae99%
- Otu262_Ruminococcaceae98%
- Otu259_Lachnospiraceae97%
- Otu256_Pseudomonadaceae100%
- Otu253_Acetobacteraceae100%
- Otu250_Ruminococcaceae100%
- Otu247_Lachnospiraceae100%
- Otu244_Lachnospiraceae99%
- Otu241_Peptostreptococcaceae25%
- Otu238_Sphingomonadaceae100%
- Otu235_Lachnospiraceae59%
- Otu232_Porphyromonadaceae97%
- Otu229_Lachnospiraceae42%
- Otu226_Coriobacteriales98%
- Otu223_Coriobacteriales100%
- Otu220_Lachnospiraceae99%
- Otu217_Lachnospiraceae97%
- Otu214_Lachnospiraceae100%
- Otu211_Planctomycetaceae9%
- Otu208_Lachnospiraceae100%
- Otu205_Porphyromonadaceae87%
- Otu202_BacillalesIncertaeSedisX100%
- Otu199_Ruminococcaceae99%
- Otu196_Eubacteriaceae46%
- Otu193_Comamonadaceae63%
- Otu190_Lachnospiraceae88%
- Otu187_ClostridialesIncertaeSedisXIII16%
- Otu184_Carnobacteriaceae100%
- Otu181_Lachnospiraceae98%
- Otu178_Lachnospiraceae93%
- Otu175_Lachnospiraceae100%
- Otu172_Lachnospiraceae100%
- Otu169_Lachnospiraceae18%
- Otu166_Ruminococcaceae99%
- Otu163_Lachnospiraceae96%
- Otu160_Rhodobacteraceae100%
- Otu157_Lachnospiraceae99%
- Otu154_Erysipelotrichaceae98%
- Otu151_Lachnospiraceae100%
- Otu148_ClostridialesIncertaeSedisXIII4%
- Otu145_Lachnospiraceae83%
- Otu142_Lachnospiraceae100%
- Otu139_Lachnospiraceae100%
- Otu136_Lachnospiraceae97%
- Otu133_Alteromonadalesincertaesedis20%
- Otu130_Lachnospiraceae100%
- Otu127_Rhodocyclaceae100%
- Otu124_Enterobacteriaceae100%
- Otu121_Ruminococcaceae98%
- Otu118_Defluviitaleaceae18%
- Otu115_Veillonellaceae100%
- Otu112_Ruminococcaceae98%
- Otu109_Lachnospiraceae94%
- Otu106_Ruminococcaceae95%
- Otu103_ClostridialesIncertaeSedisXI28%
- Otu100_Actinomycetales100%
- Otu97_ClostridialesIncertaeSedisXII30%
- Otu94_Hyphomonadaceae23%
- Otu91_Lachnospiraceae95%
- Otu88_Lachnospiraceae98%
- Otu85_Lachnospiraceae98%
- Otu82_Lachnospiraceae92%
- Otu79_ClostridialesIncertaeSedisXI16%
- Otu76_Eubacteriaceae46%
- Otu73_ClostridialesIncertaeSedisXIII8%
- Otu70_Coriobacteriales100%
- Otu67_Ruminococcaceae100%
- Otu64_ClostridialesIncertaeSedisXIII11%
- Otu61_Christensenellaceae5%
- Otu58_Staphylococcaceae100%
- Otu55_Lachnospiraceae100%
- Otu52_Ruminococcaceae98%
- Otu49_Ruminococcaceae97%
- Otu46_Coriobacteriales97%
- Otu43_Lachnospiraceae100%
- Otu40_Lachnospiraceae98%
- Otu37_Lachnospiraceae96%
- Otu34_Coriobacteriales100%
- Otu31_ClostridialesIncertaeSedisXIII6%
- Otu28_Lachnospiraceae98%
- Otu25_Lachnospiraceae100%
- Otu22_Lachnospiraceae97%
- Otu19_Christensenellaceae13%
- Otu16_Lachnospiraceae100%
- Otu13_Ruminococcaceae75%
- Otu10_Lachnospiraceae38%
- Otu7_Lachnospiraceae100%
- Otu4_Lactobacillaceae100%
- Otu1_Coriobacteriales99%