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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%
Otu70_Lubacteriaceae40 %
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%
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