



Otu340_Actinomycetales	100%
Otu337_Verrucomicrobiaceae	100%
Otu334_Porphyromonadaceae	97%
Otu331_Lachnospiraceae	100%
Otu328_ClostridialesIncertaeSedisXI	99%
Otu325_Sphingobacteriaceae	100%
Otu322_Lachnospiraceae	69%
Otu319_Burkholderiaceae	100%
Otu316_Rhodocyclaceae	14%
Otu313_Lachnospiraceae	90%
Otu310_Lachnospiraceae	100%
Otu307_Lachnospiraceae	78%
Otu304_Ruminococcaceae	90%
Otu301_Lactobacillaceae	91%
Otu298_Lachnospiraceae	96%
Otu295_Bradyrhizobiaceae	99%
Otu292_Luteivirga	1%
Otu289_Lachnospiraceae	100%
Otu286_Lachnospiraceae	100%
Otu283_Lachnospiraceae	80%
Otu280_Lactobacillaceae	90%
Otu277_Lachnospiraceae	100%
Otu274_Lachnospiraceae	97%
Otu271_Lachnospiraceae	100%
Otu268_Actinomycetales	100%
Otu265_Lachnospiraceae	99%
Otu262_Ruminococcaceae	98%
Otu259_Lachnospiraceae	97%
Otu256_Pseudomonadaceae	100%
Otu253_Acetobacteraceae	100%
Otu250_Ruminococcaceae	100%
Otu247_Lachnospiraceae	100%
Otu244_Lachnospiraceae	99%
Otu241_Peptostreptococcaceae	25%
Otu238_Sphingomonadaceae	100%
Otu235_Lachnospiraceae	59%
Otu232_Porphyromonadaceae	97%
Otu229_Lachnospiraceae	42%
Otu226_Coriobacteriales	98%
Otu223_Coriobacteriales	100%
Otu220_Lachnospiraceae	99%
Otu217_Lachnospiraceae	97%
Otu214_Lachnospiraceae	100%
Otu211_Planctomycetaceae	9%
Otu208_Lachnospiraceae	100%
Otu205_Porphyromonadaceae	87%
Otu202_BacillalesIncertaeSedisX	100%
Otu199_Ruminococcaceae	99%
Otu196_Eubacteriaceae	46%
Otu193_Comamonadaceae	63%
Otu190_Lachnospiraceae	88%
Otu187_ClostridialesIncertaeSedisXIII	16%
Otu184_Carnobacteriaceae	100%
Otu181_Lachnospiraceae	98%
Otu178_Lachnospiraceae	93%
Otu175_Lachnospiraceae	100%
Otu172_Lachnospiraceae	100%
Otu169_Lachnospiraceae	18%
Otu166_Ruminococcaceae	99%
Otu163_Lachnospiraceae	96%
Otu160_Rhodobacteraceae	100%
Otu157_Lachnospiraceae	99%
Otu154_Erysipelotrichaceae	98%
Otu151_Lachnospiraceae	100%
Otu148_ClostridialesIncertaeSedisXIII	4%
Otu145_Lachnospiraceae	83%
Otu142_Lachnospiraceae	100%
Otu139_Lachnospiraceae	100%
Otu136_Lachnospiraceae	97%
Otu133_Alteromonadalesincertaesedis	20%
Otu130_Lachnospiraceae	100%
Otu127_Rhodocyclaceae	100%
Otu124_Enterobacteriaceae	100%
Otu121_Ruminococcaceae	98%
Otu118_Defluviitaleaceae	18%
Otu115_Veillonellaceae	100%
Otu112_Ruminococcaceae	98%
Otu109_Lachnospiraceae	94%
Otu106_Ruminococcaceae	95%
Otu103_ClostridialesIncertaeSedisXI	28%
Otu100_Actinomycetales	100%
Otu97_ClostridialesIncertaeSedisXII	30%
Otu94_Hyphomonadaceae	23%
Otu91_Lachnospiraceae	95%
Otu88_Lachnospiraceae	98%
Otu85_Lachnospiraceae	98%
Otu82_Lachnospiraceae	92%
Otu79_ClostridialesIncertaeSedisXI	16%
Otu76_Eubacteriaceae	46%
Otu73_ClostridialesIncertaeSedisXIII	8%
Otu70_Coriobacteriales	100%
Otu67_Ruminococcaceae	100%
Otu64_ClostridialesIncertaeSedisXIII	11%
Otu61_Christensenellaceae	5%
Otu58_Staphylococcaceae	100%
Otu55_Lachnospiraceae	100%
Otu52_Ruminococcaceae	98%
Otu49_Ruminococcaceae	97%
Otu46_Coriobacteriales	97%
Otu43_Lachnospiraceae	100%
Otu40_Lachnospiraceae	98%
Otu37_Lachnospiraceae	96%
Otu34_Coriobacteriales	100%
Otu31_ClostridialesIncertaeSedisXIII	6%
Otu28_Lachnospiraceae	98%
Otu25_Lachnospiraceae	100%
Otu22_Lachnospiraceae	97%
Otu19_Christensenellaceae	13%
Otu16_Lachnospiraceae	100%
Otu13_Ruminococcaceae	75%
Otu10_Lachnospiraceae	38%
Otu7_Lachnospiraceae	100%
Otu4_Lactobacillaceae	100%
Otu1_Coriobacteriales	99%