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% Otu252_Ruffilliococcaceae95%
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%
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% ****3 /5 9/ 8 6> V10 /2 4 //