	Otu731_Porphyromonadaceae70%
	Otu726_Flavobacteriaceae100% Otu721_ClostridialesIncertaeSedisXIII100% Otu716_Actinomycetales100% Otu711_Lactobacillaceae93%
	Otu706_Burkholderialesincertaesedis99% Otu701_Ruminococcaceae98%
	Otu696_Lachnospiraceae100% Otu691_Alteromonadalesincertaesedis20% Otu686_Lachnospiraceae100%
	Otu681_Acidaminococcaceae100% Otu676_Actinomycetales100% Otu671 Coriobacteriales97%
	Otu666_Lachnospiraceae100% Otu661_Planococcaceae66% Otu656 Caulobacteraceae100%
	Otu651_Lachnospiraceae99% Otu646_Lachnospiraceae99% Otu641_Porphyromonadaceae74%
	Otu636_Lachnospiraceae98% Otu631_Ruminococcaceae88% Otu626_Bacillaceae185%
	Otu621_Planococcaceae78% Otu616_Ruminococcaceae77%
	Otu611_Actinomycetales100% Otu606_Lachnospiraceae99% Otu601_Porphyromonadaceae95%
	Otu596_CandidatusCarsonella14% Otu591_Ruminococcaceae99% Otu586_Erysipelotrichaceae98%
	Otu581_ClostridialesIncertaeSedisXII29% Otu576_Ruminococcaceae100% Otu571_Lachnospiraceae100%
	Otu566_Ruminococcaceae99% Otu561_Ruminococcaceae92% Otu556_Lachnospiraceae23%
	Otu551_ClostridialesIncertaeSedisXII18% Otu546_Lachnospiraceae97% Otu541_Erysipelotrichaceae96%
	Otu536_Coriobacteriales100% Otu531_ClostridialesIncertaeSedisIII100% Otu526_Ruminococcaceae79%
	Otu521_Orbaceae33% Otu516_Actinomycetales100% Otu511_Planococcaceae90%
	Otu506_Lachnospiraceae36% Otu501_ClostridialesIncertaeSedisXI100% Otu496_Clostridiaceae410%
	Otu490_Ruminococcaceae89% Otu486_Lachnospiraceae55% Otu481_Lachnospiraceae98%
	Otu476_Phyllobacteriaceae96% Otu471_Lachnospiraceae100%
	Otu466_ClostridialesIncertaeSedisXIII10% Otu461_ClostridialesIncertaeSedisXII16% Otu456_Lachnospiraceae90%
	Otu451_Lachnospiraceae42% Otu446_ClostridialesIncertaeSedisXII20% Otu441_Ruminococcaceae86%
	Otu436_Peptoniphilaceae100% Otu431_Lachnospiraceae95% Otu426_Lachnospiraceae54%
	Otu421_Ruminococcaceae82% Otu416_Ruminococcaceae22% Otu411_Clostridiaceae317%
	Otu406_Lachnospiraceae100% Otu401_Peptostreptococcaceae25% Otu396_Lachnospiraceae100%
	Otu391_Ruminococcaceae100% Otu386_Actinomycetales100% Otu381_Lachnospiraceae100%
	Otu376_Lachnospiraceae98% Otu371_Lachnospiraceae91% Otu366_Planctomycetaceae96%
	Otu361_ClostridialesIncertaeSedisXI12% Otu356_Ruminococcaceae59% Otu351_Ruminococcaceae100%
	Otu346_ClostridialesIncertaeSedisXII35% Otu341_Lachnospiraceae97% Otu336_Lachnospiraceae100%
	Otu331_Desulfomicrobiaceae42% Otu326_Lachnospiraceae99% Otu321_Lachnospiraceae100%
	Otu316_ClostridialesIncertaeSedisXII40% Otu311_Bdellovibrionaceae65% Otu306_Proteinivoraceae7%
	Otu301_Ruminococcaceae97% Otu296_Ruminococcaceae94% Otu291_Erysipelotrichaceae92%
	Otu286_Erysipelotrichaceae100% Otu281_Bacillalesincertaesedis14% Otu276_Lachnospiraceae86%
	Otu271_Lachnospiraceae87% Otu266_Porphyromonadaceae100% Otu261_Lachnospiraceae100%
	Otu256_Streptococcaceae100% Otu251_Ruminococcaceae94% Otu246_Lachnospiraceae92%
	Otu241_Lachnospiraceae100% Otu236_Lachnospiraceae99% Otu231_Erysipelotrichaceae32%
	Otu226_Lachnospiraceae99% Otu221_Lachnospiraceae99% Otu216_Bdellovibrionaceae58%
	Otu210_Coriobacteriales100% Otu201_Coriobacteriales100% Otu206_Coriobacteriales100% Otu201_ClostridialesIncertaeSedisXII19%
	Otu196_Lachnospiraceae100% Otu191_Ruminococcaceae98%
	Otu186_Coriobacteriales100% Otu181_Lachnospiraceae99% Otu176_Lachnospiraceae100%
	Otu171_Porphyromonadaceae70% Otu166_Lachnospiraceae22% Otu161_Eubacteriaceae46%
	Otu156_Rikenellaceae100% Otu151_Lachnospiraceae98% Otu146_Lachnospiraceae100%
	Otu141_Lachnospiraceae99% Otu136_Christensenellaceae13% Otu131_Rikenellaceae98%
	Otu126_Ruminococcaceae100% Otu121_Porphyromonadaceae98% Otu116_Lachnospiraceae99%
	Otu111_Lachnospiraceae100% Otu106_Defluviitaleaceae29% Otu101_Lachnospiraceae98%
	Otu96_Porphyromonadaceae97% Otu91_Catabacteriaceae28% Otu86_Lachnospiraceae91%
	Otu81_Lachnospiraceae99% Otu76_Ruminococcaceae88% Otu71_Lachnospiraceae100% Otu65_Lachnospiraceae77%
	Otu66_Lachnospiraceae97% Otu61_Ruminococcaceae90% Otu56_Porphyromonadaceae99%
are rittle table of the detected of the common of the following of the first of the common of the common of the	Otu51_Porphyromonadaceae98% Otu46_Ruminococcaceae89% Otu41_Ruminococcaceae100%
in the first of the control of the c	Otu36_Lactobacillaceae100% Otu31_Ruminococcaceae100% Otu26_Lachnospiraceae100%
<u> </u>	Otu21_Peptococcaceae147% Otu16_Lachnospiraceae96% Otu11_Lachnospiraceae100%
8885887788758758758758758758758758758758	Otu6_Bacteroidaceae100% Otu1_Planococcaceae100%
Internace Inte	1
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