



- Species
- Roseburia intestinalis
 - Bacteroides caccae
 - Blautia obeum
 - Fusicatenibacter saccharivorans
 - Agathobaculum butyriciproducens
 - Methanobrevibacter smithii
 - Parabacteroides distasonis
 - Ruminococcus callidus
 - Parabacteroides merdae
 - Alistipes putredinis
 - Lactobacillus rogosae
 - Eubacterium rectale
 - Romboutsia timonensis
 - Coprococcus eutactus
 - Prevotella copri
 - Bilophila wadsworthia
 - Dorea formicigenerans
 - Blautia luti
 - Lachnospira eligens
 - Lacnospira sp._SS3/4
 - Coprococcus comes
 - Ruminococcus bromii
 - Roseburia inulinivorans
 - Catenibacterium mitsuokai
 - Coprococcus catus
 - Eubacterium ramulus
 - Bifidobacterium bifidum
 - Prevotella stercorea
 - Eubacterium ventriosum
 - Lacnospira sp._str._L2_50
 - Alistipes finegoldii
 - Akkermansia muciniphila
 - Phocaeicola vulgatus
 - Eubacterium hallii
 - Faecalibacterium prausnitzii
 - Dorea longicatena
 - Subdoligranulum variabile
 - Collinsella aerofaciens
 - Blautia faecis
 - Bifidobacterium longum
 - Streptococcus thermophilus
 - Oscillibacter ruminantium_nov_94.144%
 - Oscillibacter valericigenes_nov_94.344%
 - Ruminococcaceae_[G-2]_bacterium_HMT_085_nov_93.946%
 - Prevotella stercorea_nov_97.403%
 - Ruminococcus callidus_nov_94.369%
 - Eubacterium coprostanoligenes_nov_97.059%
 - Kineothrix alysoides_nov_96.825%
 - Faecalibacterium prausnitzii_nov_97.964%
 - Clostridiales_[F-1][G-2]_bacterium_HMT_402_nov_89.038%
 - Oscillibacter ruminantium_nov_93.919%
 - Alloprevotella rava_nov_92.857%
 - Lachnospira pectinoschiza_nov_96.372%
 - Lachnospira eligens_nov_94.785%
 - unclassified_Ruminococcaceae_sp._str._D16_nov_93.243%
 - Butyrivibrio sp._HMT_090_nov_92.795%
 - Sporobacter termitidis_nov_90.562%
 - Paraprevotella clara_nov_97.609%
 - Prevotella colorans_nov_93.492%
 - Prevotellamassilia timonensis_nov_97.397%
 - Eubacterium coprostanoligenes_nov_92.777%
 - Asteroleplasma anaerobium_nov_80.300%
 - Succinivibrio dextrinosolvens_nov_97.079%
 - Eubacterium coprostanoligenes_nov_95.711%
 - Prevotella oralis_nov_92.641%
 - Dialister succinatiphilus_nov_96.567%
 - Duncaniella freteri_nov_87.069%
 - Ruminococcaceae_[G-2]_bacterium_HMT_085_nov_91.422%
 - Dialister succinatiphilus_nov_96.360%
 - Oscillibacter ruminantium_nov_95.701%
 - Fusicatenibacter saccharivorans_nov_97.279%
 - Lacrimispora xylanolytica_nov_97.285%
 - Clostridiales_[F-1][G-2]_bacterium_HMT_402_nov_91.011%
 - Monoglobus pectinilyticus_nov_90.112%
 - Erysipelatoclostridium [Clostridium] spiroforme_nov_93.333%
 - Lachnospira pacaense_nov_96.145%
 - Sporobacter termitidis_nov_92.584%
 - Ruminoclostridium methylpentosum_nov_89.462%
 - Ruminococcaceae_[G-2]_bacterium_HMT_085_nov_92.809%
 - Agathobaculum desmolans_nov_97.738%
 - Oscillibacter ruminantium_nov_92.601%
 - Holdemanella biformis_nov_97.854%
 - Bacteroides sp._str._4136_uniformis
 - Anaerostipes_Lacnospira_unclassified_Lachnospiraceae
 - Lactobacillus_Ligilactobacillus_ruminis
 - Roseburia faecalis_faecis
 - Blautia_Mediterraneibacter [Ruminococcus] gnavus_gnavus
 - Blautia_unclassified_Lachnospiraceae_sp._str._cont1.79_torques
 - Lacnospira sp._SM4/1_sp._str._M62/1
 - Bacteroides_Phocaeicola_dorei_sp._Oral_Taxon_D25_sp._str._219
 - Bifidobacterium adolescentis_faecale
 - Escherichia_Shigella_coli_dysenteriae_fergusonii_flexneri_sonnei
 - Blautia obeum_wexlerae
 - Parasutterella_Sutterella_excrementihominis_sp._str._cont1.66
 - Bacteroides_korensis_kribbi
 - Bacteroides_Phocaeicola_plebeius
 - Ruminococcus_multispecies_sppn1_2_nov_95.023%
 - Roseburia_multispecies_sppn106_3_nov_96.833%
 - Christensenella_multispecies_sppn2_2_nov_91.216%
 - Clostridium_multispecies_sppn27_2_nov_97.959%

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Samples