



- Faecalibacterium prausnitzii
- Ligilactobacillus salivarius
- Intestinibacter bartlettii
- Ligilactobacillus animalis
- Streptococcus lutetiensis
- Methanobrevibacter smithii
- Lactobacillus amylovorus
- Lactobacillus johnsonii
- Limosilactobacillus reuteri
- Prevotella copri
- Streptococcus equinus
- Sarcina ventriculi
- Bacteroidetes_[G-7] bacterium_HMT_911_nov_90.164%
- Coprococcus eutactus_nov_91.649%
- Prevotella copri_nov_97.959%
- Prevotella copri_nov_97.755%
- Prevotella saccharolytica_nov_90.389%
- Phascolarctobacterium faecium_nov_89.505%
- Sporobacter termitidis_nov_87.794%
- Faecalicatena fissicatena_nov_96.660%
- Prevotella brevis_nov_89.184%
- Treponema pectinovorum_nov_87.755%
- Coprococcus eutactus_nov_93.133%
- Prevotella sp._MOT-128_nov_86.992%
- Bacteroidetes_[G-3] bacterium_HMT_280_nov_89.278%
- Ethanoligenens harbinense_nov_87.609%
- Lachnospiraceae_[G-9] bacterium_MOT-174_nov_91.494%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_91.875%
- Prevotella copri_nov_94.501%
- Paludibacter propioniciigenes_nov_85.685%
- Prevotella maculosa_nov_90.061%
- Oribacterium sinus_nov_91.213%
- Ruminococcus champanellensis_nov_90.323%
- Anaerocolumna cellulositytica_nov_87.657%
- Butyrificoccus pullicaecorum_nov_87.810%
- Prevotella stercorea_nov_96.296%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.306%
- Ihubacter massiliensis_nov_92.917%
- Syntrophotalea carbinolica_nov_83.497%
- Prevotella copri_nov_95.732%
- Bacteroidetes_[G-7] bacterium_HMT_911_nov_89.528%
- Eubacterium ruminantium_nov_91.260%
- Mogibacterium neglectum_nov_89.855%
- Bacteroidetes_[G-3] bacterium_HMT_899_nov_87.730%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.340%
- Prevotella copri_nov_97.551%
- Treponema sp._HMT_490_nov_87.247%
- Magaeibacillus indolicus_nov_87.238%
- Treponema sp._HMT_264_nov_83.534%
- Ruminococcus champanellensis_nov_93.952%

Species

- F25123.S05
- F25123.S04
- F25123.S03
- F25123.S02
- F25123.S01
- F25123.S10
- F25123.S09
- F25123.S08
- F25123.S07
- F25123.S06

Samples