



- Species
- Adlercreutzia caecimuris
 - Comamonas sediminis
 - Akkermansia muciniphila
 - Herbaspirillum huttiense
 - Acinetobacter radioresistens
 - Peptococcaceae_[G-1] bacterium_MOT-146
 - Duncaniella freteri
 - Sphingobium limneticum
 - Actinidia eriantha
 - Lactococcus cremoris
 - Lactococcus lactis
 - Moraxella osloensis
 - Ligilactobacillus murinus
 - Adlercreutzia mucosicola
 - Acinetobacter lwoffii
 - Streptococcus danieliae
 - Phocaeicola sartorii
 - Adlercreutzia muris
 - Erwinia billingiae
 - Sphingomonas carotinifaciens
 - Limosilactobacillus reuteri
 - Kocuria indica
 - Bacteroides acidifaciens
 - Pelomonas saccharophila
 - Parabacteroides goldsteinii
 - Helicobacter ganmani
 - Sphingomonas echinoides
 - Erysipelatoclostridium [Clostridium] cocleatum
 - Lactobacillus johnsonii
 - Lactobacillus gasseri
 - Lachnospiraceae_[G-1] bacterium_MOT-166_nov_95.661%
 - Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.059%
 - Prevotella shahii_nov_87.242%
 - Faecalibaculum rodentium_nov_96.571%
 - Clostridium collagenovorans_nov_80.952%
 - Bariatricus massiliensis_nov_93.037%
 - Eubacterium ventriosum_nov_92.843%
 - Desulfovibrio fairfieldensis_nov_89.168%
 - Bariatricus massiliensis_nov_93.230%
 - Mucispirillum schaedleri_nov_93.307%
 - Adlercreutzia caecimuris_nov_94.000%
 - Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.861%
 - Sporobacter termitidis_nov_82.970%
 - Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.353%
 - Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.590%
 - Lachnospiraceae_[G-12] bacterium_MOT-179_nov_94.737%
 - Maiihella massiliensis_nov_90.377%
 - Lachnospiraceae_[G-7] bacterium_MOT-172_nov_93.204%
 - Duncaniella freteri_nov_90.152%
 - Duncaniella freteri_nov_87.896%
 - Parasutterella excrementihominis_nov_94.584%
 - Muribaculaceae_[G-1] bacterium_MOT-129_nov_87.308%
 - Lachnospiraceae_[G-11] bacterium_MOT-176_nov_92.885%
 - Ihubacter massiliensis_nov_94.767%
 - Planococcus massiliensis_nov_96.992%
 - Lachnospiraceae_[G-12] bacterium_MOT-180_nov_89.942%
 - Duncaniella freteri_nov_87.759%
 - Lachnospiraceae_[G-7] bacterium_MOT-172_nov_92.843%
 - Lachnoclostridium [Clostridium] polysaccharolyticum_nov_93.050%
 - Caproicibacter fermentans_nov_89.824%
 - Fusicatenibacter saccharivorans_nov_90.514%
 - Lachnospiraceae_[G-3] bacterium_MOT-168_nov_95.059%
 - Lachnospiraceae_[G-11] bacterium_MOT-176_nov_94.798%
 - Duncaniella freteri_nov_89.077%
 - Adlercreutzia caecimuris_nov_95.382%
 - Oscillospiraceae_[G-4] bacterium_MOT-151_nov_93.491%
 - Sporosalbacterium tautonense_nov_82.659%
 - Abssiella tortuosum_nov_88.725%
 - Lachnospiraceae_[G-14] bacterium_MOT-183_nov_97.967%
 - Parabacteroides merdae_nov_93.182%
 - Lachnoclostridium [Clostridium] polysaccharolyticum_nov_90.751%
 - Lachnospiraceae_[G-7] bacterium_MOT-172_nov_94.831%
 - Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.905%
 - Adlercreutzia caecimuris_nov_92.644%
 - Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.083%
 - Anaerotignum aminivorans_nov_92.585%
 - Lachnoclostridium [Clostridium] aminophilum_nov_89.961%
 - Lacrimispora xylanolytica_nov_93.969%
 - Oscillospiraceae_[G-4] bacterium_MOT-151_nov_94.477%
 - Faecalibaculum rodentium_nov_96.571%
 - Blautia faecicola_nov_89.709%
 - Lacrimispora xylanolytica_nov_94.314%
 - Muribaculaceae_[G-1] bacterium_MOT-129_nov_89.768%
 - Bacteroides uniformis_nov_95.594%
 - Lacrimispora xylanolytica_nov_91.992%
 - Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.643%
 - Flavobacterium branchicola_nov_96.282%
 - Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.050%
 - Faecalicatena orotica_nov_92.218%
 - Lachnoclostridium [Clostridium] scindens_nov_88.247%
 - Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.050%
 - Eisenbergiella massiliensis_nov_88.292%
 - Lachnospiraceae_[G-11] bacterium_MOT-177_nov_96.267%
 - Roseburia hominis_nov_92.471%
 - Eisenbergiella massiliensis_nov_88.697%
 - Eubacterium xylanophilum_nov_91.149%
 - Lachnospiraceae_[G-14] bacterium_MOT-182_nov_92.245%
 - Prevotella multispecies_sppn3_2_nov_89.792%
 - Desulfovibrio multispecies_sppn4_2_nov_96.275%

F10949.S07
F10949.S08
F10949.S09
F10949.S10
F10949.S11
F10949.S12
F10949.S13

Samples