



Group

SAA KO Normal

SAA KO BL HFD



- Neisseria shayegani
Adlercreutzia caecimuris
Comamonas sediminis
Akkermansia muciniphila
Herbaspirillum huttiense
Sphingomonas deserti
Helicobacter hepaticus
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
Erwinia billingiae
Sphingomonas carotinifaciens
Limosilactobacillus reuteri
Kocuria indica
Bacteroides acidifaciens
Pelomonas saccharophila
Parabacteroides goldsteinii
Helicobacter ganmani
Sphingomonas echinoides
Lachnospiraceae_[G-3] bacterium_MOT-168
Erysipelatoclostridium [Clostridium] cocleatum
Lactobacillus johnsonii
Lactobacillus gasseri
Lachnospiraceae_[G-1] bacterium_MOT-166_nov_95.661%
Bacteroidetes_[G-3] bacterium_HMT_436_nov_85.575%
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%
Desulfovibrio fairfieldensis_nov_89.168%
Longibaculum muris_nov_91.211%
Mucispirillum schaedleri_nov_93.307%
Anaerotignum lactatifermentans_nov_95.769%
Anaerotruncus rubiinfantis_nov_91.506%
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%
Mallihella 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%
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-14] bacterium_MOT-182_nov_89.200%
Lachnospiraceae_[G-11] bacterium_MOT-176_nov_94.798%
Duncaniella freteri_nov_89.077%
Faecalicatena fissicatena_nov_94.521%
Sporosolibacterium tautonense_nov_82.659%
Absiella 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%
Lacrimispora xylanolytica_nov_93.969%
Oscillospiraceae_[G-4] bacterium_MOT-151_nov_94.477%
Pseudoflavonifractor phocaeensis_nov_92.131%
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 branchiicola_nov_96.282%
Faecalicatena orotica_nov_92.218%
Lachnoclostridium [Clostridium] scindens_nov_88.247%
Oscillospiraceae_[G-3] bacterium_MOT-150_nov_93.910%
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%
Parabacteroides distasonis_nov_97.323%
Prevotella multispecies_sppn3_2_nov_89.792%

Species

F10949.S07

F10949.S08

F10949.S09

F10949.S14

F10949.S15

F10949.S16

F10949.S17

Samples