



Group

SAA KO Normal
SAA KO HFD

Lachnospiraceae_[G-14] bacterium_MOT-183_nov_97.967%
Sporobacter termitidis_nov_82.970%
Parabacteroides merdae_nov_93.182%
Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.905%
Lachnospiraceae_[G-7] bacterium_MOT-172_nov_92.843%
Duncaniella frateri_nov_89.077%
Desulfovibrio fairfieldensis_nov_89.168%
Bariatrikus massiliensis_nov_93.037%
Lachnoclostridium [Clostridium] polysaccharolyticum_nov_90.751%
Duncaniella frateri_nov_90.152%
Duncaniella frateri_nov_87.896%
Parasutterella excrementihominis_nov_94.584%
Duncaniella frateri_nov_87.759%
Lachnospiraceae_[G-3] bacterium_MOT-168_nov_95.059%
Anaerotignum aminivorans_nov_92.585%
Lachnospiraceae_[G-11] bacterium_MOT-176_nov_92.885%
Sporosalibacterium tautonense_nov_82.659%
Lachnospiraceae_[G-7] bacterium_MOT-172_nov_94.831%
Lachnoclostridium [Clostridium] polysaccharolyticum_nov_93.050%
Lachnospiraceae_[G-11] bacterium_MOT-176_nov_94.798%
Lachnospiraceae_[G-7] bacterium_MOT-172_nov_93.204%
Lachnospiraceae_[G-12] bacterium_MOT-180_nov_89.942%
Absiella tortuosum_nov_88.725%
Lachnospiraceae_[G-12] bacterium_MOT-179_nov_94.737%
Adlercreutzia caecimuris_nov_92.644%
Fusicatenibacter saccharivorans_nov_90.514%
Eisenbergiella massiliensis_nov_88.292%
Desulfovibrio multispecies_sppn4_2_nov_96.275%
Oscillospiraceae_[G-4] bacterium_MOT-151_nov_93.491%
Lachnospiraceae_[G-14] bacterium_MOT-182_nov_92.245%
Oscillospiraceae_[G-4] bacterium_MOT-151_nov_94.477%
Lacrimispora xylanolytica_nov_93.969%
Faecalcalenata orotica_nov_92.218%
Parabacteroides distasonis_nov_97.323%
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.861%
Adlercreutzia caecimuris_nov_94.000%
Ihubacter massiliensis_nov_94.767%
Lachnoclostridium [Clostridium] scindens_nov_88.247%
Parabacteroides goldsteinii
Adlercreutzia muris
Muribaculaceae_[G-1] bacterium_MOT-129_nov_87.308%
Peptococcaceae_[G-1] bacterium_MOT-146
Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.050%
Akkermansia muciniphila
Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.059%
Faecalibaculum rodentium_nov_96.571%
Bariatrikus massiliensis_nov_93.230%
Adlercreutzia caecimuris_nov_95.382%
Eisenbergiella massiliensis_nov_88.697%
Adlercreutzia mucosicola
Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.083%
Lachnoclostridium [Clostridium] aminophilum_nov_89.961%
Lacrimispora xylanolytica_nov_92.992%
Mucispirillum schaedleri_nov_93.307%
Caproicibacter fermentans_nov_89.824%
Streptococcus danieliae
Sphingomonas echinoides
Herbaspirillum huttiense
Moraxella osloensis
Pelomonas saccharophila
Sphingobium limneticum
Sphingomonas carotinifaciens
Planococcus massiliensis_nov_96.992%
Acinetobacter iwoffii
Erwinia billingiae
Comamonas sediminis
Flavobacterium branchiicola_nov_96.282%
Acinetobacter radioresistens
Prevotella multispecies_sppn3_2_nov_89.792%
Roseburia hominis_nov_92.471%
Lachnospiraceae_[G-1] bacterium_MOT-166_nov_95.661%
Duncaniella frateri
Bacteroides uniformis_nov_95.594%
Clostridium collagenovorans_nov_80.952%
Lachnospiraceae_[G-11] bacterium_MOT-177_nov_96.267%
Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.050%
Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.643%
Lacrimispora xylanolytica_nov_94.314%
Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.353%
Helicobacter ganmani
Blautia faecicola_nov_89.709%
Bacteroides acidifaciens
Eubacterium xylanophilum_nov_91.149%
Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.590%
Phocaeicola sartorii
Adlercreutzia caecimuris
Mailhella massiliensis_nov_90.377%
Kocuria indica
Lactococcus cremoris
Lactococcus lactis
Lactobacillus gasseri
Limosilactobacillus reuteri
Faecalibaculum rodentium_nov_96.571%
Eubacterium ventriosum_nov_92.843%
Prevotella shahii_nov_87.242%
Muribaculaceae_[G-1] bacterium_MOT-129_nov_89.768%
Ligilactobacillus murinus
Lactobacillus johnsonii
Erysipelatoclostridium [Clostridium] cocleatum
Actinidia eriantha

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

Samples

Species