



Group

- WT Normal
- WT HFD
- SAA KO Normal
- SAA KO HFD
- SAA KO HFD

Species

Parabacteroides distasonis 80.3614%

Oscillospiraceae [G-2] bacterium_MOT-149_nov_93.861%

Lacrimispora xylanolytica_nov_91.992%

Lachnospiraceae [G-6] bacterium_MOT-171_nov_94.083%

Adlercreutzia mucosicola

Lachnospiraceae [G-14] bacterium_MOT-182_nov_92.245%

Oscillospiraceae [G-4] bacterium_MOT-151_nov_94.477%

Eisenbergiella massiliensis_nov_88.292%

Lachnospiraceae [G-14] bacterium_MOT-182_nov_92.245%

Lachnospiraceae [G-3] bacterium_MOT-168_nov_94.059%

Lachnospiraceae [G-6] bacterium_MOT-171_nov_95.644%

Lachnospiraceae [G-14] bacterium_MOT-182_nov_92.245%

Eubacteriales [G-2] bacterium_MOT-162

Phocaea massiliensis_nov_90.297%

Oscillospiraceae [G-6] bacterium_MOT-153

Sporobacter thermophilus_nov_82.970%

Eubacteriales [G-4] bacterium_MOT-164

Lachnospiraceae [G-3] bacterium_MOT-168_nov_94.653%

Lachnospiraceae [G-7] bacterium_MOT-172_nov_91.945%

Lancefieldella parvula_nov_91.552%

Lachnospiraceae [G-14] bacterium_MOT-182_nov_92.245%

Lachnospiraceae [G-3] bacterium_MOT-168

Oscillospiraceae [G-3] bacterium_MOT-150_nov_93.910%

Parabacteroides distasonis

Longibaculum muris_nov_91.211%

Lachnospiraceae [G-11] bacterium_MOT-176_nov_94.798%

Olsenella phocaeensis_nov_92.172%

Lachnospiraceae [G-11] bacterium_MOT-176_nov_95.543%

Adlercreutzia muris

Bacteroidetes [G-3] bacterium_HMT_436_nov_85.575%

Parabacteroides distasonis_nov_97.514%

Longibaculum muris_nov_93.910%

Phocaea vulgaris

Lactococcus lactis

Eubacterium ventriosum_nov_92.843%

Eubacterium ramulus_nov_91.296%

Lawsonibacter asaccharolyticus_nov_91.571%

Lachnospiraceae [G-14] bacterium_MOT-185_nov_92.105%

Lachnospiraceae [G-14] bacterium_MOT-185_nov_92.105%

Oscillospiraceae [G-2] bacterium_MOT-149_nov_95.050%

Turicibacter sanguinis_nov_95.817%

Clostridium disporicum

Eisenbergiella massiliensis_nov_90.805%

Akkermansia muciniphila

Eubacteriales [G-1] bacterium_MOT-159

Lachnospiraceae [G-14] bacterium_MOT-185

Leifsonia kafiensis_nov_84.158%

Anaeroplasmabacterium_nov_87.352%

Faecalicatena fissicatena_nov_93.580%

Prevotella multispecies_sppn3_2_nov_89.792%

Roseburia hominis_nov_92.471%

Bacteroides uniformis_nov_95.594%

Lachnospiraceae [G-1] bacterium_MOT-166_nov_95.661%

Duncaniella freteri

Clostridium collagenovorans_nov_80.952%

Helicobacter ganmani

Eubacterium xylanophilum_nov_91.149%

Phocaea sartorii

Bacteroides acidifaciens

Lachnospiraceae [G-6] bacterium_MOT-171_nov_95.050%

Lachnospiraceae [G-11] bacterium_MOT-177_nov_96.267%

Lachnospiraceae [G-7] bacterium_MOT-172_nov_93.204%

Desulfotomaculum fairfieldensis_nov_89.168%

Bariatricus massiliensis_nov_93.037%

Lachnospiraceae [G-14] bacterium_MOT-185_nov_92.353%

Duncaniella freteri_nov_90.152%

Lacrimispora xylanolytica_nov_94.314%

Lachnospiraceae [G-6] bacterium_MOT-171_nov_94.643%

Blautia faecicola_nov_89.709%

Eisenbergiella massiliensis_nov_88.697%

Faecalicatena fissicatena_nov_94.521%

Faecalicatena orotica_nov_92.218%

Lachnospiraceae [G-6] bacterium_MOT-171_nov_93.307%

Comamonas sediminis

Flavobacterium branchiicola_nov_96.282%

Acinetobacter radioresistens

Kocuria indica

Lactococcus cremoris

Adlercreutzia caecimuris

Streptococcus danieliae

Caproicibacter fermentans_nov_89.824%

Mucispirillum schaedleri_nov_93.307%

Maihiella massiliensis_nov_90.377%

Iubacter massiliensis_nov_94.767%

Prevotella shahii_nov_87.242%

Muribaculaceae [G-1] bacterium_MOT-129_nov_89.768%

Lactobacillus gasseri

Limosilactobacillus reuteri

Neisseria shayegani

Faecalibaculum rodentium

Muribaculaceae [G-1] bacterium_MOT-129_nov_86.590%

Ligilactobacillus murinus

Faecalibaculum rodentium_nov_96.571%

Romboutsia ilealis

Actinidia eriantha

Lactobacillus johnsonii

Sphingomonas deserti

Erysipelatoclostridium [Clostridium] coelestem

Species