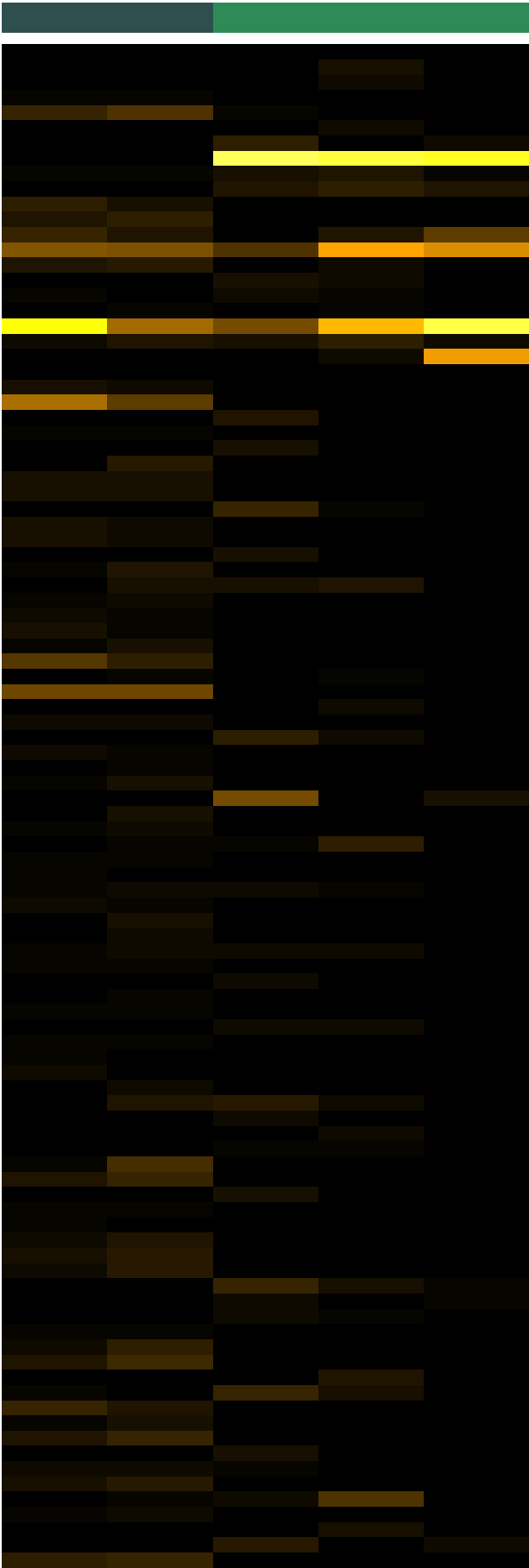




Group

SAA KO Normal

SAA KO BL HFD



- Peptococcaceae_[G-1] bacterium_MOT-146
- Helicobacter hepaticus
- Akkermansia muciniphila
- Bacteroides caecimuris
- Phocaeicola sartorii
- Lactococcus lactis
- Parabacteroides goldsteinii
- Erysipelatoclostridium [Clostridium] cocleatum
- Streptococcus danieliae
- Lactococcus cremoris
- Bacteroides acidifaciens
- Duncaniella freteri
- Lactobacillus gasseri
- Ligilactobacillus murinus
- Helicobacter ganmani
- Lachnospiraceae_[G-3] bacterium_MOT-168
- Adlercreutzia mucosicola
- Adlercreutzia muris
- Lactobacillus johnsonii
- Adlercreutzia caecimuris
- Limosilactobacillus reuteri
- Parabacteroides distasonis
- Faecalibaculum rodentium_nov_96.571%
- Prevotella shahii_nov_87.054%
- Anaerotruncus rubiinfantis_nov_91.506%
- Sporobacter termitidis_nov_82.970%
- Anaerotignum lactatifermentans_nov_95.577%
- Lachnospiraceae_[G-7] bacterium_MOT-172_nov_93.204%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.353%
- Duncaniella freteri_nov_90.152%
- Maiihella massiliensis_nov_90.377%
- Duncaniella freteri_nov_87.896%
- Parasutterella excrementihominis_nov_94.584%
- Blautia producta_nov_96.132%
- Lachnospiraceae_[G-11] bacterium_MOT-176_nov_92.885%
- Caproicibacter fermentans_nov_89.864%
- Lachnospiraceae_[G-12] bacterium_MOT-180_nov_89.942%
- Lachnospiraceae_[G-7] bacterium_MOT-172_nov_92.843%
- Duncaniella freteri_nov_87.759%
- Lachnospiraceae_[G-3] bacterium_MOT-168_nov_95.059%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_86.590%
- Fusicatenibacter saccharivorans_nov_90.514%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_89.768%
- Lachnospiraceae_[G-14] bacterium_MOT-182_nov_89.200%
- Duncaniella freteri_nov_89.077%
- Oscillospiraceae_[G-3] bacterium_MOT-150_nov_93.910%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.905%
- Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.841%
- Anaerotignum aminivorans_nov_92.585%
- Sporosolibacterium tautonense_nov_82.659%
- Ihubacter massiliensis_nov_94.402%
- Absiella tortuosum_nov_88.725%
- Faecalicatena fissicatena_nov_94.521%
- Lachnospiraceae_[G-14] bacterium_MOT-183_nov_97.967%
- Faecalicatena fissicatena_nov_94.129%
- Lachnoclostridium [Clostridium] polysaccharolyticum_nov_90.751%
- Parabacteroides merdae_nov_93.182%
- Lachnospiraceae_[G-7] bacterium_MOT-172_nov_94.831%
- Adlercreutzia caecimuris_nov_92.644%
- Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.083%
- Lachnospiraceae_[G-14] bacterium_MOT-185_nov_94.726%
- Pseudoflavonifractor phocaeensis_nov_92.131%
- Phoceia massiliensis_nov_90.060%
- Duncaniella freteri_nov_89.013%
- Oscillospiraceae_[G-4] bacterium_MOT-151_nov_94.477%
- Duncaniella freteri_nov_87.453%
- Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351_nov_95.551%
- Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.544%
- Oscillospiraceae_[G-2] bacterium_MOT-149_nov_96.640%
- Lachnospiraceae_[G-11] bacterium_MOT-176_nov_94.798%
- Lacrimispora xylanolytica_nov_93.969%
- Lachnospiraceae_[G-14] bacterium_MOT-182_nov_92.480%
- Lachnospiraceae_[G-11] bacterium_MOT-176_nov_94.981%
- Blautia faecicola_nov_89.709%
- Bacteroides uniformis_nov_95.594%
- Lacrimispora xylanolytica_nov_91.992%
- Lachnospiraceae_[G-12] bacterium_MOT-179_nov_94.737%
- Eisenbergiella massiliensis_nov_90.805%
- Lacrimispora xylanolytica_nov_94.314%
- Lachnospiraceae_[G-11] bacterium_MOT-177_nov_95.874%
- Lachnospiraceae_[G-6] bacterium_MOT-171_nov_94.643%
- Eisenbergiella massiliensis_nov_88.697%
- Bacteroidetes_[G-3] bacterium_HMT_436_nov_85.575%
- Lachnoclostridium [Clostridium] scindens_nov_88.247%
- Muribaculaceae_[G-1] bacterium_MOT-129_nov_87.308%
- Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.050%
- Roseburia hominis_nov_92.471%
- Lachnospiraceae_[G-14] bacterium_MOT-182_nov_92.245%
- Faecalicatena orotica_nov_92.218%
- Eubacterium xylanophilum_nov_91.149%
- Lachnoclostridium [Clostridium] polysaccharolyticum_nov_93.050%
- Lachnospiraceae_[G-1] bacterium_MOT-166_nov_95.661%
- Parabacteroides distasonis_nov_97.323%
- Bariatricus massiliensis_nov_93.037%
- Clostridium collagenovorans_nov_80.952%
- Mucispirillum schaedleri_nov_93.110%
- Desulfovibrio fairfieldensis_nov_89.168%
- Lachnospiraceae_[G-3] bacterium_MOT-168_nov_94.059%
- Longibaculum muris_nov_91.211%
- Prevotella multispecies_sppn4_2_nov_89.792%

Species

F10949.S07

F10949.S08

F10949.S14

F10949.S15

F10949.S16

Samples