



Group

SAA KO Normal

SAA KO BL HFD



- Peptococcaceae_[G-1]
- Phocaeicola
- Parabacteroides
- Erysipelatoclostridium
- Streptococcus
- Lactococcus
- Duncaniella
- Ligilactobacillus
- Helicobacter
- Lachnospiraceae_[G-3]
- Lactobacillus
- Adlercreutzia
- Limosilactobacillus
- Faecalibaculum
- Prevotella
- Anaerotruncus
- Lachnospiraceae_[G-7]
- Maiihella
- Parasutterella
- Caproicibacter
- Lachnospiraceae_[G-12]
- Fusicatenibacter
- Muribaculaceae_[G-1]
- Oscillospiraceae_[G-3]
- Muribaculaceae_[G-2]
- Anaerotignum
- Ihubacter
- Sporosolibacterium
- Absiella
- Lachnoclostridium
- Lachnospiraceae_[G-6]
- Pseudoflavonifractor
- Oscillospiraceae_[G-4]
- Oscillospiraceae_[G-2]
- Lachnospiraceae_[G-11]
- Blautia
- Bacteroides
- Lacrimispora
- Eisenbergiella
- Bacteroidetes_[G-3]
- Roseburia
- Lachnospiraceae_[G-14]
- Faecalicatena
- Eubacterium
- Lachnospiraceae_[G-1]
- Bariatricus
- Clostridium
- Mucispirillum
- Desulfovibrio
- Longibaculum

Genus

F10949.S07

F10949.S08

F10949.S14

F10949.S15

F10949.S16

Samples