

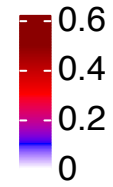
Samples

HB

NB

Bacteroides fragilis
Faecenecus gallistercoris
Faecimonas intestinavium
Lactobacillus crispatus
Escherichia coli
Anaerostipes butyraticus
Alloclastrodium intestingalinarum
Faecalibacterium faecigalinarum
Alistipes excrementavium
Faecalibacterium gallistercoris
Merdicola_sp001915925
Pelethenecus faecipullorum
Mediterraneibacter caccogalinarum
Ruthenibacterium merdavium
Gemmiger formicilis_B
Anaerotruncus colihominis
Clostridium_Q_saccharolyticum_A
Coproplasma stercorigalinarum
Mediterraneibacter avicola
Finenecus excrementavium
Coproplasma stercoravium
Choladocola avistercoris
Merdisoma faecalis
Scybalousia_sp900543675
Butyricoccus pullicaecorum
Coproplasma avistercoris
Caccovivens_sp930979925
Harysmithimonas_galli
Finenecus stercoravium
Merdicola faecigalinarum
Enterocloster excrementipullorum
Heteroclostridium caecigalinarum
Anaerotignum lactatifermentans
Merdibacter merdigalinarum
Anaerotignum merdipullorum
Onthovivinus excrementipullorum
Anaeromassilibacillus stercoravium
Negativibacillus faecipullorum
Enterocloster excrementigalinarum
Gallimonas_sp944367465
Fusicatenibacter_sp900543115
Blautia ornithocaccaceae
CAG-269_sp904419495
Caccovivinus merdipullorum
Thomasclostridium spiroformis
Gallimonas merdigalinarum
Ornithoclostridium excrementipullorum
Bifidobacterium pullorum_B
Butyricoccus_sp900604335
Ligilactobacillus salivarius
Ruthenibacterium avium
Caccousia avicola

Relative abundance



Taxonomic family

- Acutalibacteraceae
- Anaeroplasmataceae
- Anaerotignaceae
- Bacteroidaceae
- Bifidobacteriaceae
- Borkfalkiaceae
- Butyricococcaceae
- CAG-314
- CAG-508
- CAJFEE01
- Coprobacillaceae
- Enterobacteriaceae
- Erysipelotrichaceae
- Lachnospiraceae
- Lactobacillaceae
- Rikenellaceae
- Ruminococcaceae
- UBA1242
- UBA3700
- UBA660