





Genus

- [Eubacterium] eligens group*
- [Eubacterium] xylanophilum group*
- [Ruminococcus] gauvreauii group*
- [Ruminococcus] torques group*
- Acidaminococcus*
- Agathobacter*
- Akkermansia*
- Alistipes*
- Anaerostipes*
- Bacteroides*
- Bifidobacterium*
- Bilophila*
- Blautia*
- Butyrimonas*
- Catenibacterium*
- Christensenellaceae R-7 group*
- Collinsella*
- Coprococcus 3*
- Desulfovibrio*
- Dorea*
- Erysipelotrichaceae UCG-003*
- Escherichia-Shigella*
- Faecalibacterium*
- Flavonifractor*
- Fusicatenibacter*
- gut metagenome*
- Intestinibacter*
- Lachnospiraceae NK4A136 group*
- Megasphaera*
- Methanobrevibacter*
- Odoribacter*
- Oscillibacter*
- Parabacteroides*
- Paraprevotella*
- Parasutterella*
- Phascolarctobacterium*
- Prevotella 9*
- Lachnoclostridium*
- Roseburia*
- Ruminiclostridium 5*
- Ruminiclostridium 6*
- Ruminococcaceae NK4A214 group*
- Ruminococcaceae UCG-002*
- Ruminococcaceae UCG-003*
- Ruminococcaceae UCG-005*
- Ruminococcus 1*
- Ruminococcus 2*
- Subdoligranulum*
- Tyzzerella*
- UBA1819*
- uncultured*
- uncultured Thermoanaerobacterales bacterium*