





Genus

- g__[Eubacterium] eligens group*
- g__[Eubacterium] xylanophilum group*
- g__[Ruminococcus] gauvreauii group*
- g__[Ruminococcus] torques group*
- g__Acidaminococcus*
- g__Agathobacter*
- g__Akkermansia*
- g__Alistipes*
- g__Anaerostipes*
- g__Bacteroides*
- g__Bifidobacterium*
- g__Bilophila*
- g__Blautia*
- g__Butyricimonas*
- g__Catenibacterium*
- g__Christensenellaceae R-7 group*
- g__Collinsella*
- g__Coprococcus 3*
- g__Desulfovibrio*
- g__Dorea*
- g__Erysipelotrichaceae UCG-003*
- g__Escherichia-Shigella*
- g__Faecalibacterium*
- g__Flavonifractor*
- g__Fusicatenibacter*
- g__gut metagenome*
- g__Intestinibacter*
- g__Lachnospiraceae NK4A136 group*
- g__Megasphaera*
- g__Methanobrevibacter*
- g__Odoribacter*
- g__Oscillibacter*
- g__Parabacteroides*
- g__Paraprevotella*
- g__Parasutterella*
- g__Phascolarctobacterium*
- g__Prevotella 9*
- g__Roseburia*
- g__Ruminiclostridium 5*
- g__Ruminiclostridium 6*
- g__Ruminococcaceae NK4A214 group*
- g__Ruminococcaceae UCG-002*
- g__Ruminococcaceae UCG-003*
- g__Ruminococcaceae UCG-005*
- g__Ruminococcus 1*
- g__Ruminococcus 2*
- g__Subdoligranulum*
- g__Tyzzerella*
- g__UBA1819*
- g__uncultured*
- g__uncultured Thermoanaerobacterales bacterium*