

allcohorts

- g__Bacteroides
- g__Bifidobacterium
- g__Butyrivibrio
- g__Catenibacterium
- g__Clostridium.sensu.stricto.1
- g__Collinsella
- g__Dialister
- g__Eubacterium.siraeum.group
- g__Faecalibacterium
- g__Fusicatenibacter
- g__Holdemanella
- g__Lachnoclostridium
- g__Lachnospiraceae.NK4A136.group
- g__Marvinbryantia
- g__Megasphaera
- g__Mitsuokella
- g__NK4A214.group
- g__Phascolarctobacterium
- g__Rikenellaceae.RC9.gut.group
- g__Romboutsia
- g__Roseburia
- g__Ruminococcus
- g__Subdoligranulum
- g__Succinivibrio
- g__Treponema
- g__UCG.005

