Clostridiaceae bacterium DJF LS40 human gut metagenome Clostridium sp. Culture Jar-8 Lubacterium siraeum V10Sc8a Ruminiclostridium bacterium coprostanoligenes group Angelakisella massiliensis Ruminiclostridium Buman gut metagenome [Clostridium] cocleatum Clostridiales bacterium 30-4c Angelakisella Ruminococcaceae UCG-004 Erysipelatoclostridium Anaerotruncus sp. G3(2012) Flavonifractor gut metagenome Anaerotruncus Ruminococcaceae **Ervsipelotrichaceae** Ruminococcaceae UCectobacillus apodemi DSM 16634 = JCM 16172 Oscillibacter Ruminococcaceae UCG-010Ruminiclostridiumbacterium enrichment culture Erysipelotrichales Lactobacillus metagenome Ruminiclostridium Clostridium saudiense Lactobacillaceae [Clostridium] leptum Clostridiaceae 1 Erysipelotrichia Ak Akkermansia sp. UNK.MGS-1 Akkermansia Akkermansiaceae Romboutsia ilealis Lactobacillales Verrucomicrobiales Clostridiales Romboutsia Peptostreptococcaceae Bacilli Verrucomicrobiae Lachnospiraceae bacterium DW67 Firmicutes Verrucomicrobia metagenome Deferribacterales Clostridiales bacterium CISOA090006657 Clostridia Deferribacteres Deferribacte Clostridiaceae bacterium SH021 Clostridiaceae bacterium Zagget Bacteria Deferribacteres Mucispirillum Clostridium sp. Clone-44 Shuttleworthia human gut metagenome Proteobacteria Roseburia Tyzzerella Hungatella Bacteroidetes Lachnospiraceae Bacterium AJ110@Aterobacteriales metagenome Lachnospiraceae bacterium AJ110@Aterobacteriales
Lachnospiraceae bacteriumaenospiraceae lurcspiraceae NK4A136 grgup
Marvinbryanttiachnospiraceae UCG-006metagenome

Marvinbryanttiachnospiraceae UCG-006metagenome Nodes metagenome [Eubacterium] fissicatena grainenbetgiellmospiraceae bacteriu Bradtorordalleiscum aestivum (bread wheat) 0.00 -5.00UC5-1-2E3 mouse gut metagenome Acetatifactor Clostridiales bacterium CICASTO Lam sp. ASF502achnospiraceae bacterium G11 rannerellaceae 1.42 -3.33Salmonella enterica subsp. enterica Eisenbergiella tayi ratio Eubacteriaceae bacterium UC5.1–2E3 Parabacteroides -1.675.67 Blautia [Clostridium asparagiforme] DSM 15981 Lachnospiraceae bacterium 5 Costricum fusiformis human gut metagenome 0.00 12.80 2 Clostridiales bacterium VE202-@chnoclostridium [Clostridium] symbiosum ATCC 14940 22.70 1.67 Lachnospiraceae bacterium DW3 metagenome 35.40 3.33 Clostridium sp. M62/1 mouse gut metagenome 5.00 51.00