



truncLen	0-0	270-270	280-280	290-290	300-300
Prevotella	0	0	0	0	0
Ruminococcus_2	0	0	0	0	0
Escherichia/Shigella	0	0	0	0	0
Bacteroides	0	0	0	0	0
Blautia	0	0	0	0	0
Agathobacter	0	0	0	0	0
Sutterella	0	0	0	0	0
Alloprevotella	0	0	0	0	0
Leptotrichia	0	0	0	0	0
Prevotella_7	0	0	0	0	0
Anaerobacillus	0	0	0	0	0
Erysipelotrichaceae_UCG-003	0	0	0	0	0
Collinsella	0	0	0	0	0
Olsenella	0	0	0	0	0
Bifidobacterium	0	0	0	0	0
Parvimonas	0	0	0	0	0
Parabacteroides	0	0	0	0	0
Ruminococcaceae_UCG-013	0	0	0	0	0
Alistipes	0	0	0	0	0
Bilophila	0	0	0	0	0
Ammonia	0	0	0	0	0
Butyrivibrio	0	0	0	0	0
Odoribacter	0	0	0	0	0
Rikenellaceae_RC9_gut_group	0	0	0	0	0
Subdoligranulum	0	0	0	0	0
Lachnospiraceae_UCG-010	0	0	0	0	0
Peptostreptococcus	0	0	0	0	0
Ruminococcaceae_UCG-004	0	0	0	0	0
Selenomonas_3	0	0	0	0	0
Ruminococcaceae_NK4A214_group	0	0	0	0	0
Fusobacterium	0	0	0	0	0
Paraprevotella	0	0	0	0	0
Prevotella_9	0	0	0	0	0
Peptococcus	0	0	0	0	0
Christensenellaceae_R-7_group	0	0	0	0	0
Ruminococcaceae_UCG-005	0	0	0	0	0
Ruminococcaceae_UCG-010	0	0	0	0	0
Victivallis	0	0	0	0	0
Slackia	0	0	0	0	0
Selenomonas	0	0	0	0	0
Dialister	0	0	0	0	0
Gemella	0	0	0	0	0
Lachnoanaerobaculum	0	0	0	0	0
Treponema_2	0	0	0	0	0
Ruminococcaceae_UCG-002	0	0	0	0	0
Streptococcus	0	0	0	0	0
Faecalibacterium	0	0	0	0	0
Ruminococcaceae_UCG-014	0	0	0	0	0
Flavonifractor	0	0	0	0	0
GCA-900066225	0	0	0	0	0
Family_XIII_AD3011_group	0	0	0	0	0
Desulfovibrio	0	0	0	0	0
Coprobacter	0	0	0	0	0
Delftia	0	0	0	0	0
Anaeroplasm	0	0	0	0	0
Lachnospira	0	0	0	0	0
Howardella	0	0	0	0	0
Brachyspira	0	0	0	0	0
Ruminococcus_1	0	0	0	0	0
Phascolarctobacterium	0	0	0	0	0
Lachnospiraceae_ND3007_group	0	0	0	0	0
Acidibacter	0	0	0	0	0
Ruminiclostridium_9	0	0	0	0	0
Solobacterium	0	0	0	0	0
DTU089	0	0	0	0	0
Barnesiella	0	0	0	0	0
Anaerotruncus	0	0	0	0	0
Roseburia	0	0	0	0	0
Cutibacterium	0	0	0	0	0
Distigma	0	0	0	0	0
Ruminiclostridium_6	0	0	0	0	0
Iyzerella	0	0	0	0	0
Butyricimonas	0	0	0	0	0
Acetanaerobacterium	0	0	0	0	0
Holdemanella	0	0	0	0	0
Stenotrophomonas	0	0	0	0	0
Fuscatenibacter	0	0	0	0	0
Sediminibacterium	0	0	0	0	0
Lachnoclostridium	0	0	0	0	0
Senegalimassilia	0	0	0	0	0
Ruminiclostridium_5	0	0	0	0	0
Oscillibacter	0	0	0	0	0
CAG-56	0	0	0	0	0
Mogibacterium	0	0	0	0	0
Fretibacterium	0	0	0	0	0
Lachnospiraceae_NK4A136_group	0	0	0	0	0
Anaerostipes	0	0	0	0	0
Dorea	0	0	0	0	0
Coprococcus_2	0	0	0	0	0
Lachnospiraceae_UCG-001	0	0	0	0	0
Marvinbryantia	0	0	0	0	0
Lachnospiraceae_UCG-003	0	0	0	0	0
Lachnospiraceae_UCG-004	0	0	0	0	0
Coprococcus_1	0	0	0	0	0
Coprococcus_3	0	0	0	0	0
Sphingomonas	0	0	0	0	0
Oribacterium	0	0	0	0	0
Ralstonia	0	0	0	0	0
Haemophilus	0	0	0	0	0
Caproiciproducens	0	0	0	0	0
Catabacter	0	0	0	0	0
Adlercreutzia	0	0	0	0	0
Falsochrobacterium	0	0	0	0	0
Lachnospiraceae_FCS020_group	0	0	0	0	0
Candidatus_Soleaferrea	0	0	0	0	0
Parasutterella	0	0	0	0	0
Ruminococcaceae_UCG-007	0	0	0	0	0
Enterorhabdus	0	0	0	0	0
Microbacterium	0	0	0	0	0