Species

Ruminococcoides intestinale
Agathobacter rectalis
Fusicatenibacter saccharivorans
Agathobacter faecis
Bifidobacterium pseudocatenulatum
Blautia_A wexlerae
Bacteroides stercoris
Prevotella copri_B
Bacteroides uniformis
Dialister invisus
Anaerostipes hadrus
Dorea_A longicatena
Methanocatella smithii
Bifidobacterium adolescentis
Faecalibacterium prausnitzii
Gemmiger qucibialis
Mediterraneibacter faecis
Dialister sp900545785
Alistipes putredinis
Anthropogastromicrobium aceti
Hominimerdicola aceti
Bifidobacterium longum
Ruminococcoides intestinalis
Mediterraneibacter lactaris
Phocaeicola dorei
Blautia_A fusiformis
Faecalibacterium duncaniae
Anaerobutyricum hallii
Aphodomorpha intestinavium
Bariatricus comes
Hominisplanchenecus faecis
Blautia_A obeum
Dialister sp000434475
Faecalibacterium longum
CAG−217 sp000436335
Prevotella sp003447235
Phocaeicola vulgatus
CAG−269 sp000431335
CAG−273 sp003534295
Klebsiella pneumoniae
Lachnospira rogosae
Bifidobacterium bifidum
Coprococcus aceti
Roseburia intestinalis
Roseburia inulinivorans
Faecalibacterium sp900539945
Parabacteroides merdae
CAG−273 sp000437855
Butyribacter intestini
Ruminococcus_C callidus
Hominenteromicrobium mulieris
Butyribacter sp003529475
Faecalibacterium taiwanense
Porcipelethomonas sp900538575
Blautia_A faecis
Lachnospira sp000437735
Alloprevotella sp900539625
Hominicoprocola fusiformis
Holdemanella sp900556915
Bifidobacterium animalis
Lachnospira eligens_A
Phascolarctobacterium faecium
Eubacterium_G ventriosum
Bacteroides massiliensis_D
Methanosphaera stadtmanae
Anaerostipes amylophilus
Coprococcus eutactus
CAG−590 sp900544905
UBA11524 sp000437595
Anaerotardibacter sp959607715
Enterocloster sp000431375
Phocaeicola plebeius
Bacteroides eggerthii
Otoolea fessa
Blautia_A sp900066335
Monoglobus sp. MAG085
Hominilimicola fabiformis
CAG−177 sp003514385
Barnesiella intestinihominis
Anaerotardibacter sp000436075
Parabacteroides distasonis
Coprococcus_A intestinihominis
Prevotella copri_G
Collinsella sp. MAG140
Succiniclasticum sp900314855
Ruminococcus_C sp000980705
Sutterella wadsworthensis_A
PeH17 sp000435055
SFEL01 sp004557245
Roseburia_C amylophila
Acetatifactor intestinalis
Gallintestinimicrobium propionicum
Agathobaculum butyriciproducens
Ruminococcus_C sp000433635
Mediterraneibacter torques
Anaerotardibacter sp000435675
Faecalitalea cylindroides
Catenibacterium mitsuokai
Akkermansia muciniphila
Prevotella copri_K
Eubacterium_R sp000434995
Mesosutterella multiformis
Phocaeicola coprophilus
Merdicola sp018371855
Evtepia gabavorous
Paraprevotella clara
CAG−302 sp000431795
Blautia_A intestinihominis
CAG−1193 sp900550145
Phocaeicola sp000432735
Fimivicinus sp900544375
Faecalibacillus intestinalis
Eubacterium_R sp000433975
Limivivens faecihominis
Gemmiger formicilis
Eubacterium_F sp003491505
Vescimonas sp000435555
Phocaeicola barnesiae
Romboutsia timonensis
Butyrivibrio_A sp000431815
CAG−103 sp900543625
Oscillibacter segnis
Vescimonas sp000435975
Anaerobutyricum soehngenii
Clostridium saudiense
CAG−417 sp000432835
Phocaeicola sp. MAG088
Megasphaera massiliensis
Eubacterium_I ramulus
Ligilactobacillus ruminis
Roseburia hominis
Maccoya intestinihominis
Faecalibacillus faecis
Pilosibacter fragilis
Coprococcus ammoniilyticus
Parasutterella excrementihominis
Collinsella sp. MAG118
Gemmiger sp900540595
Wujia hominis
Agathobaculum hominis
CAG−269 sp001916035
Slackia isoflavoniconvertens
CAG−177 sp000431775
Oliverpabstia faecicola
Hominicoprocola sp900317525
Blautia stercoris
Bacteroides cellulosilyticus
Adlercreutzia rubneri
CAG−103 sp934589185
Agathobacter sp000434275
Bacteroides fluxus
QAMH01 sp900544245
Alistipes communis
Gemmiger sp900540775
Oliverpabstia intestinalis
Faecalibacterium prausnitzii_I
Senegalimassilia anaerobia
Fusicatenibacter_A intestinipullorum
Cryptobacteroides sp000432515
Limisoma sp000437795
Faecalibacterium hattorii
Onthocola_B sp900544625
Phocaeicola plebeius_A
Limivicinus sp900549645
Sellimonas intestinalis
Angelakisella sp900547385
Hominenteromicrobium sp900549945
Clostridium_A leptum
Ventricola sp900542395
Blautia_A sp003474435
Coprococcus sp000154245
Vescimonas coprocola
Collinsella sp900555225
Fimenecus sp000432435
Collinsella sp. MAG165
CAG−103 sp000432375
CAG−245 sp000435175
Faecousia sp000434635
F23−B02 sp945613045
Eubacterium_G segne
Akkermansia massiliensis
Ruminococcus sp900540005
Enorma massiliensis
Apopatocola sp003522945
Vescimonas sp900545495
Eubacterium_F sp900539115
Vescimonas sp900554275
Limisoma sp900544305
Brotaphodocola catenula
Acutalibacter timonensis
Aphodovivens avicola
UMGS1338 sp900550805
Butyricicoccus_A intestinisimiae
CAG−103 sp900757655