ASV5_Fusicatenibacter

ASV21_Alistipes

ASV299_Mitsuokella

ASV130_Megasphaera

ASV346_Lachnospiraceae NK3A20 group

ASV218_Prevotella

ASV28_Parabacteroides

ASV11_Romboutsia ASV25_Clostridium sensu stricto 1 ASV25_Clostri<mark>dium sensu stricto 1</mark> ASV36_Intestinibacter ASV72_Enterococcus ASV135_Terrisporobacter ASV234_Lachnospiraceae UCG010 ASV234_Lachnospiraceae UCG010 ASV2_Agathobacter ASV94_Rikenellaceae RC9 gut group ASV94_Rikenellaceae RC9 gut group ASV3_Faecalibacterium ASV3_Faecalibacterium ASV29_Holdemanella ASV29_Holdemanella ASV559_Adlercreutzia ASV559_Adlercreutzia ASV2277_Actinomyces ASV2277_Actinomyces ASV24_Subdoligran&&W43_Roseburia ASV24_Subdoligran&&May Roseburia ASV1635_Oscillospira ASV1635_Oscillospira ASV239_Lachnospiraceae FCS020 group ASV239_Lachnospiraceae FCS020 group ASV672_[Eubacterium] brachy group ASV31_Catenibacterium [Ruminococcus] torques group ASV672_[Eubacterium] brachy group ASV31_Catenibacterium_ ASV32_[Ruminococcus] torques group ASV5_Fusicatenibacter *SV1 Blautia ASV9 Dorea ASV13 Collinsella ASV985_Shuttleworthia ASV1_Blautia / ASV9_Dorea ASV13_Collinsella ASV985 Shuttleworthia ASV1 ASV576_[Eubacterium] fissicatena group ASV576_[Eubacterium] fissicatena group ASV788_Anaerotruncus ASV788_Anaerotruncus ASV71_Erysipelotrichaceae UCG003 ASV71_Erysipelotrichaceae UCG003 ASV875_Eggerthella ASV7_[Eubacterium] hallii graffV84_CAG56 ASV875_Eggerthella ASV7_[Eubacterium] hallii gr&\$V84_CAG56 ASV39_Anaerostipes ASV39_Anaerostipes ASV950 Strasip eletestatudium nocuum group ASV95A STrasip qletestastridiy mnocuum group ASV220_Bilophila ASV220_Bilophila ASV1907_Paludicolars V493_Flavonifractor ASV394_UBA1819 ASV1907_Paludicola/SV493_Flavonifractor ASV394_UBA1819 ASV12_Bacteroides ASV12_Bacteroides ASV453_Peptococcus ASV346_Lachnospiraceae NK3A20 group ASV453_Peptococcus ASV178_[Ruminococcus] gnavus group ASV178_[Ruminococcus] gnavus group ASV14_Prevotella_9 ASV791_Lachnospiraceae NK4B4 group ASV14_Prevotella_9 ASV791_Lachnospiraceae NK4B4 group ASV21_Alistipes ASV218_Prevotella ASV106_Alloprevotella ASV106_Alloprevotella ASV23_UCG002 ASV256_Family XIII AD3011 group ASV142_Odoribacter ASV23_UCG002 ASV142_Odoribacter ASV256_Family XIII AD3011 group A\$V28_Parabacteroides ASV125_UCG005 ASV125_UCG005 ASV136_UCG003 ASV136_UCG003 ASV40_Akkermansia ASV40_Akkermansia ASV266_Coriobacteriaceae UCG003 ASV266_Coriobacteriaceae UCG003 ASV299_Mitsuokella ASV123_NK4A214 group ASV123_NK4A214 group ASV53_Christensenellaceae R7 group ASV53_Christensenellaceae R7 group ASV391_Olsenella ASV391_Olsenella ASV295_Methanoprevibagite/clostridium ASV295_Methanopreviberteclostridium ASV130_Megasphaera ASV70_Lachnospiraceae ND3007 group ASV78_Lachnospira ASV70_Lachnospiraceae ND3007 group ASV78_Lachnospira ASV642_Allisonella ASV642_Allisonella ASV712_Veillonella ASV712_Veillonella ASV168_Phascolarctobacterium ASV168_Phascolarctobacterium ASV279_Haemophilus ASV279_Haemophilus ASV59_Dialister ASV59_Dialister

ASV288_Limos Factor See Lingtobacillus