ASV639_Weissella ASV34_Lachnospiraceae NK4A136 group ASV72_Enter ASVCt566_Rothia ASV84_CAG56 ASV43_Roseburia ASV13_Collinsella ASV94_Rikenellaceae RC9 gut group ASV239_Lachnospiraceae FCS020 group ASV2_Agathobacter ASV791_Lachnospiraceae NK4B4 group ASV5_Fusicatenibacter ASV32_[Ruminococcus] torques group ASV672_[Eubacterium] brachy group ASV3_Faecalibacterium ASV33_Coprococcus ASV24_Subdoligranulum ASV1907_Paludicola ASV875_Eggerthella Holdemania ASV788_Anaerotruncus ASV201_[Eubacterium] ventriosum group ASV28_Parabacteroides ASV39_Anaerostipes ASV12_Bacteroides ASV21_Alistipes ASV576_[Eubacterium] fissicatena group ASV218_Prevotella ASS/4430 FISIOSTFIGILIEN innocuum group ASV1635_Oscillospira ASV394_UBA1819 ASV71_Erysipelotrichaceae UCG003 ASV495_Candidatus Soleaferrea ASV346_Lachnospiraceae NK3A20 group ASV142_Odoribacter ASV178_[Ruminococcus] gnavus group ASV2277_Actinomyces ASV299_Mitsuokella ASV424_UCG009 ASV14 Prevotella 9 ASV31_Catenibacterium ASV106_Alloprevotella ASV23_UCG002 ASV130_Megasphaera ASV53_ChristenseneffSV295RMethanobrevibacter ASV136_UCG003 ASV336_Tyzzerella ASV29_Holdemanella ASV40_Akkermansia ASV125_UCG005 ASV123_NK4A214 group ASV642_Allisonella ASV453_Peptococcus ASV256_Family XIII AD3011 group ASV67_Lachnoclostridium ASV386 Lactobacillus ASV220_Bilophila ASV288_Limosilactobacillus ASV279_Haemophilus ASV70_Lachnospiraceae ND3007 group ASV11_Romboutsia ASV116_Butyricicoccus ASV712_Veillonella ASV78_Lachnospira ASV25_Clostridium sensu stricto 1 ASV59_Dialister ASV135_Terrisporobacter_{V168}_Phascolarctobacterium ASV36_Intestinibacter ASV266_Coriobacteriaceae UCG003 ASV391_Olsenella ASV234_Lachnospiraceae UCG0269_Turicibacter

ASV639_Weissella ASV34_Lachnospiraceae NK4A136 group ASV72_Enter ASVCt566_Rothia ASV84_CAG56 ASV43_Roseburia ASV13_Collinsella ASV94_Rikenellaceae RC9 gut group ASV239_Lachnospiraceae FCS020 group ASV2_Agathobacter ASV9_Dorea ASV791_Lachnospiraceae NK4B4 group ASV5_Fusicatenibacter ASV32_[Ruminococcus] torques group ASV672_[Eubacterium] brachy group ASV3_Faecalibacterium ASV33_Coprococcus ASV1_Blautia ASV1907_Paludicola ASV24_Subdoligranulum ASV875_Eggerthella Holdemania ASV788_Anaerotruncus ASV201_[Eubacte<mark>rium]</mark> ventriosum group ASV28_Parabacteroides ASV39_Anaerostipes ASV985_Shuttlew69th950_Erysipelatoclostridium ASV576_[Eubacterium] fissicatena group ASV12 Bacteroldésipes ASV218_Prevotella ASSY4930 FISUSTIFICIUM innocuum group ASV1635_Oscillospira ASV394_UBA1819 ASV71_Erysipelotrichaceae UCG003 ASV495_Candidatus Soleaferrea ASV346_Lachnospiraceae NK3A20 group ASV142_Odoribacter ASV178_[Ruminococcus] gnavus group ASV299_Mitsuokella ASV2277_Actinomyces ASV424_UCG009 ASV14_Prevotella_9 ASV31_Catenibacterium ASV106_Alloprevotella ASV23_UCG002 ASV130_Megasphaera Christenseneflaceae R Methanobrevibacter ASV1<mark>36_UC</mark>G003 ASV29_Holdemanella ASV336_Tyzzerella ASV40_Akkermansia ASV125_UCG005 ASV123_NK4A214 group ASV642_Allisonella ASV453_Peptococcus ASV256_Family XIII AD3011 group ASV67_Lachnoclostridium ASV386_Lactobacillus ASV220_Bilophila ASV288_Limosilactobacillus ASV279_Haemophilus ASV/70_Lachnospiraceae ND3007 group ASV11_Romboutsia ASV116_Butyricicoccus ASV78_Lachnospira ASV712_Veillonella ASV25_Clostridium sensu stricto 1 ASV59_Dialister ASV135_Terrisporobacter_{V168}_Phascolarctobacterium ASV36_Intestinibacter ASV266_Coriobacteriaceae UCG003 ASV391_Olsenella ASV234_Lachnospiraceae UCG0269_Turicibacter