

Species

●	SP31 Lachnospiraceae_[G-11] bacterium_MOT-178
●	SP35 Muribaculum intestinale
●	SP45 Turicibacter sanguinis
●	SP46 Acutalibacter muris
●	SP49 Oscillospiraceae_[G-4] bacterium_MOT-151
●	SP5 Lachnospiraceae_[G-1] bacterium_MOT-166
●	SP50 Lachnospiraceae_[G-5] bacterium_MOT-170
●	SP54 Anaerostipes caccae
●	SP55 Erysipelotrichaceae_[G-1] bacterium_MOT-189
●	SP6 Anaerotignum lactatifermentans
●	SP65 Acetatifactor muris
●	SP7 Romboutsia timonensis
●	SP70 Lachnospiraceae_[G-11] bacterium_MOT-176
●	SP72 Lachnospiraceae_[G-2] bacterium_MOT-167
●	SP74 Adlercreutzia muris
●	SP75 Parasutterella excrementihominis
●	SP76 Alistipes sp._MOT-127
●	SP93 Lawsonibacter asaccharolyticus
●	SP96 Lachnospiraceae_[G-3] bacterium_MOT-168
●	SP97 Erysipelatoclostridium [Clostridium] cocleatum
●	SP98 Faecalibaculum rodentium
●	SP99 Peptococcaceae_[G-1] bacterium_MOT-146
●	SPN100 Acetitomaculum ruminis_nov_90.551%
●	SPN101 Acetatifactor muris_nov_94.118%
●	SPN102 Muribaculum intestinale_nov_93.676%
●	SPN103 Roseburia intestinalis_nov_92.913%
●	SPN104 Acetatifactor muris_nov_97.638%
●	SPN106 Muribaculaceae_[G-2] bacterium_MOT-104_nov_90.551%
●	SPN107 Lachnospira eligens_nov_87.109%
●	SPN108 Lachnospiraceae_[G-7] bacterium_MOT-172_nov_93.307%
●	SPN109 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_92.969%
●	SPN112 Peptostreptococcaceae_[G-2] bacterium_HMT_091_nov_81.600%
●	SPN120 Muribaculaceae_[G-1] bacterium_MOT-129_nov_93.676%
●	SPN130 Muribaculum intestinale_nov_96.443%
●	SPN140 Prevotella sp._MOT-128_nov_93.307%
●	SPN16 Eubacteriales_[G-3] bacterium_MOT-163_nov_96.457%
●	SPN161 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_97.638%
●	SPN163 Muribaculaceae_[G-1] bacterium_MOT-129_nov_93.676%
●	SPN164 Desulfovibrio fairfieldensis_nov_96.457%
●	SPN175 Kineothrix alysoides_nov_97.255%

●	SPN388 Lawsonibacter asaccharolyticus_nov_94.094%
●	SPN39 Lachnoclostridium [Clostridium] populeti_nov_93.359%
●	SPN398 Butyribacter intestini_nov_95.294%
●	SPN402 Merdimonas faecis_nov_96.457%
●	SPN406 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_97.255%
●	SPN417 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_94.531%
●	SPN427 Breznakia pachnodaee_nov_92.157%
●	SPN447 Papillibacter cinnamivorans_nov_89.370%
●	SPN46 Oscillibacter ruminantium_nov_94.902%
●	SPN464 Muribaculum intestinale_nov_93.676%
●	SPN465 Muribaculum intestinale_nov_93.676%
●	SPN47 Lachnoclostridium [Clostridium] scindens_nov_93.725%
●	SPN477 Kineothrix alysoides_nov_97.244%
●	SPN498 Kineothrix alysoides_nov_97.255%
●	SPN501 Eisenbergiella massiliensis_nov_96.457%
●	SPN509 Lachnospiraceae_[G-11] bacterium_MOT-178_nov_97.638%
●	SPN530 Anaerotignum lactatifermentans_nov_95.276%
●	SPN541 Merdimonas faecis_nov_96.484%
●	SPN552 Eubacteriales_[G-4] bacterium_MOT-164_nov_97.244%
●	SPN56 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_93.701%
●	SPN562 Eubacterium coprostanoligenes_nov_94.902%
●	SPN573 Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.095%
●	SPN582 Anaerotruncus colihominis_nov_94.466%
●	SPN591 Sporobacter termitidis_nov_94.094%
●	SPN595 Acetivibrio alkalicellulosi_nov_88.189%
●	SPN608 Lachnoclostridium [Clostridium] scindens_nov_93.701%
●	SPN61 Lachnospiraceae_[G-9] bacterium_MOT-174_nov_93.359%
●	SPN629 Hungatella xylanolytica_nov_96.063%
●	SPN63 Hungatella effluvii_nov_97.638%
●	SPN639 Anaerotruncus rubiinfantis_nov_93.307%
●	SPN651 Paraeggerthella hongkongensis_nov_92.520%
●	SPN659 Eubacterium ventriosum_nov_95.669%
●	SPN667 Muribaculum intestinale_nov_95.686%
●	SPN668 Lawsonibacter asaccharolyticus_nov_97.638%
●	SPN69 Lachnospira multipara_nov_87.451%
●	SPN690 Butyrvibrio crossotus_nov_95.294%
●	SPN694 Acetatifactor muris_nov_94.118%
●	SPN700 Lachnoclostridium pacaense_nov_97.638%
●	SPN711 Lachnoclostridium [Clostridium] scindens_nov_92.969%
●	SPN719 Butyribacter intestini_nov_93.333%

●	SPN843 Clostridiales_[F-1][G-1] bacterium_HMT_093_nov_89.804%
●	SPN85 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_96.457%
●	SPN87 Lachnoclostridium [Clostridium] scindens_nov_92.913%
●	SPN871 Muribaculum intestinale_nov_94.466%
●	SPN872 Senegalimassilia anaerobia_nov_91.339%
●	SPN88 Neglectibacter timonensis_nov_94.094%
●	SPN89 Neglectibacter timonensis_nov_96.850%
●	SPN892 Anaerotaenia torta_nov_88.583%
●	SPN895 Eubacteriales_[G-4] bacterium_MOT-164_nov_97.638%
●	SPN90 Christensenella minuta_nov_92.188%
●	SPN91 Muribaculum intestinale_nov_90.909%
●	SPN92 Muribaculaceae_[G-2] bacterium_MOT-104_nov_94.466%
●	SPN93 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.370%
●	SPN94 Hydrogenoanaerobacterium saccharovorans_nov_94.094%
●	SPN95 Oscillospiraceae_[G-4] bacterium_MOT-151_nov_97.638%
●	SPN97 Frisingicoccus caecimuris_nov_94.882%
●	SPN98 Lachnoclostridium [Clostridium] scindens_nov_91.406%
●	SPN99 Anaerosporbacter mobilis_nov_96.063%
●	SPP11 Ligilactobacillus animalis_apodemi_faecis_murinus
●	SPP12 Bacteroides faecichinchillae_faecis_thetaiotaomicron
●	SPP19 Lactobacillus gasseri_hominis_johnsonii_taiwanensis
●	SPP21 Bifidobacterium choerinum_pseudolongum
●	SPP23 Staphylococcus casei_cohnii_cohnii_subsp_cohnii_edaphicus_gallin ...(9 species)
●	SPP24 Blautia hansenii_hominis_marasmi
●	SPP25 Escherichia_Pseudescherichia_Shigella_boydii_coli_fergusonii_flexneri_marmotae_sonnei_vu ...(7 species)
●	SPP26 Adlercreutzia caecimuris_mucosicola
●	SPP3 Limosilactobacillus antri_caviae_fruentii_oris_panis_reuteri_reuteri_c ...(8 species)
●	SPP8 Enterocloster bolteaee_clostridioformis
●	SPPN1 multigenus multispecies_sppn1_2_nov_93.676%
●	SPPN102 multigenus multispecies_sppn102_4_nov_94.510%
●	SPPN106 multigenus multispecies_sppn106_2_nov_94.882%
●	SPPN115 multigenus multispecies_sppn115_4_nov_89.764%
●	SPPN124 multigenus multispecies_sppn124_2_nov_96.063%
●	SPPN13 multigenus multispecies_sppn13_2_nov_87.008%
●	SPPN133 multigenus multispecies_sppn133_2_nov_96.063%
●	SPPN143 multigenus multispecies_sppn143_2_nov_95.276%
●	SPPN175 Lacrimispora multispecies_sppn175_2_nov_97.244%
●	SPPN177 Enterocloster multispecies_sppn177_2_nov_94.488%
●	SPPN185 Lachnospiraceae_[G-12] multispecies_sppn185_2_nov_93.307%
●	SPPN21 Roseburia multispecies_sppn21_2_nov_95.276%