

Species

- SP311 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 Anaerosporeobacter 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_fruementi_oris_panis_reuteri_reuteri_c ...(8 species)
- SPP8 Enterocloster bolteae_clostridioformis
- SPPN1 Oscillospiraceae_multigenus multispecies_sppn1_2_nov_93.676%
- SPPN102 Lachnospiraceae_multigenus multispecies_sppn102_4_nov_94.510%
- SPPN106 Lachnospiraceae_multigenus multispecies_sppn106_2_nov_94.882%
- SPPN115 Lachnospiraceae_multigenus multispecies_sppn115_4_nov_89.764%
- SPPN124 Lachnospiraceae_multigenus multispecies_sppn124_2_nov_96.063%
- SPPN13 Eubacteriales_multifamily_multigenus multispecies_sppn13_2_nov_87.008%
- SPPN133 Lachnospiraceae_multigenus multispecies_sppn133_2_nov_96.063%
- SPPN143 Lachnospiraceae_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%