

Species

- SP103 Porphyromonas sp._MOT-131
- SP104 Pediococcus pentosaceus
- SP106 Streptococcus acidominimus
- SP11 Streptococcus danieliae
- SP110 Eubacteriales_[G-4] bacterium_MOT-164
- SP114 Enterococcus faecalis
- SP12 Corynebacterium stationis
- SP121 Mammaliicoccus lentus
- SP129 Bifidobacterium pseudolongum
- SP133 Mollicutes_[G-1] bacterium_MOT-186
- SP135 Lactococcus lactis
- SP138 Lactobacillus johnsonii
- SP14 Streptococcus thoraltensis
- SP149 Stenotrophomonas [Pseudomonas] hibiscicola
- SP15 Staphylococcus ureilyticus
- SP168 Weissella cibaria
- SP17 Staphylococcus saprophyticus
- SP172 Anaerococcus sp._HMT_290
- SP2 Triticum aestivum
- SP22 Muribacter sp._MOT-143
- SP23 Rodentibacter pneumotropicus
- SP25 Streptococcus thermophilus
- SP29 Moraxella osloensis
- SP3 Enterococcus gallinarum
- SP33 Limosilactobacillus reuteri
- SP34 Staphylococcus equorum
- SP36 Massilia aurea
- SP37 Psychrobacter alimentarius
- SP38 Dubosiella newyorkensis
- SP41 Atopostipes sp._MOT-201
- SP42 Ligilactobacillus animalis
- SP43 Lachnospiraceae_[G-14] bacterium_MOT-185
- SP44 Ligilactobacillus murinus
- SP46 Eubacteriales_[G-2] bacterium_MOT-162
- SP47 Cutibacterium granulosum
- SP48 Corynebacterium mastitidis
- SP50 Cutibacterium acnes
- SP52 Clostridium disporicum
- SP54 Stenotrophomonas maltophilia
- SP56 Streptomyces aculeolatus
- SP57 Lachnospiraceae_[G-11] bacterium_MOT-177
- SP64 Pseudomonas helleri
- SP65 Lactocaseibacillus rhamnosus
- SP69 Jeotgaliococcus halotolerans
- SP71 Levilactobacillus brevis
- SP72 Gemella sp._MOT-033
- SP73 Corynebacterium ammoniagenes
- SP75 Secundilactobacillus paracollinoides
- SP77 Leptothrix sp._HMT_025
- SP8 Weissella paramesenteroides
- SP84 Delftia acidovorans
- SP87 Carnobacteriaceae_[G-1] bacterium_MOT-197
- SP90 Rothia nasimurium
- SP92 Pelomonas saccharophila
- SP96 Ralstonia sp._HMT_406
- SP97 Actinidia eriantha
- SP99 Akkermansia muciniphila
- SPN109 Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.105%
- SPN110 Duncaniella freteri_nov_93.293%
- SPN122 Turicibacter sanguinis_nov_95.923%
- SPN151 Enterococcus faecalis_nov_95.825%
- SPN159 Fusobacterium varium_nov_96.696%
- SPN188 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.867%
- SPN21 Oribacterium parvum_nov_89.770%
- SPN226 Duncaniella freteri_nov_89.775%
- SPN272 Alistipes senegalensis_nov_93.443%
- SPN276 Erysipelotrichaceae_[G-1] bacterium_MOT-189_nov_88.798%
- SPN283 Peptococcus sp._HMT_168_nov_84.866%
- SPN286 Duncaniella freteri_nov_89.718%
- SPN29 Fusicatenibacter saccharivorans_nov_90.526%
- SPN293 Oscillospiraceae_[G-6] bacterium_MOT-153_nov_91.631%
- SPN3 Actinidia eriantha_nov_97.011%
- SPN302 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.000%
- SPN311 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.719%
- SPN319 Fusobacterium perfoetens_nov_91.126%
- SPN47 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.423%
- SPN54 Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.887%
- SPN66 Duncaniella freteri_nov_88.934%
- SPN72 Duncaniella freteri_nov_93.699%
- SPP10 Latilactobacillus curvatus_graminis
- SPP11 Sphingomonas aquatilis_melonis
- SPP9 Staphylococcus saprophyticus_xylosus