

Species

- SP1 Streptomyces aculeolatus
- SP10 Staphylococcus saprophyticus
- SP12 Triticum aestivum
- SP14 Corynebacterium stationis
- SP17 Stenotrophomonas maltophilia
- SP18 Cutibacterium acnes
- SP2 Ligilactobacillus murinus
- SP23 Lachnospiraceae_[G-11] bacterium_MOT-177
- SP24 Ligilactobacillus animalis
- SP25 Mammaliicoccus lentus
- SP26 Lachnospiraceae_[G-14] bacterium_MOT-185
- SP27 Corynebacterium ammoniagenes
- SP28 Streptococcus thermophilus
- SP33 Delftia acidovorans
- SP34 Cutibacterium granulosum
- SP35 Massilia aurea
- SP39 Actinidia eriantha
- SP40 Enterococcus gallinarum
- SP41 Moraxella osloensis
- SP42 Mollicutes_[G-1] bacterium_MOT-186
- SP45 Atopostipes sp._MOT-201
- SP47 Enterococcus faecalis
- SP49 Pelomonas saccharophila
- SP51 Limosilactobacillus reuteri
- SP53 Clostridium disporicum
- SP54 Leptothrix sp._HMT_025
- SP56 Lactobacillus johnsonii
- SP64 Anaerococcus sp._HMT_290
- SP65 Staphylococcus equorum
- SP68 Pseudomonas helleri
- SP7 Jeotgalicoccus halotolerans
- SP70 Stenotrophomonas [Pseudomonas] hibiscicola
- SP73 Dubosiella newyorkensis
- SP74 Eubacteriales_[G-2] bacterium_MOT-162
- SP76 Ralstonia sp._HMT_406
- SP78 Psychrobacter alimentarius
- SP8 Eubacteriales_[G-4] bacterium_MOT-164
- SP80 Bradyrhizobium pachyrhizi
- SP83 Akkermansia muciniphila
- SP84 Secundilactobacillus paracollinoides
- SP9 Staphylococcus ureilyticus
- SP93 Bifidobacterium pseudolongum
- SPN10 Oribacterium parvum_nov_89.770%
- SPN102 Fusobacterium varium_nov_96.696%
- SPN107 Duncaniella freteri_nov_93.293%
- SPN117 Enterococcus faecalis_nov_95.825%
- SPN118 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.867%
- SPN149 Duncaniella freteri_nov_89.775%
- SPN15 Fusicatenibacter saccharivorans_nov_90.526%
- SPN173 Alistipes senegalensis_nov_93.443%
- SPN178 Peptococcus sp._HMT_168_nov_84.866%
- SPN181 Duncaniella freteri_nov_89.718%
- SPN188 Oscillospiraceae_[G-6] bacterium_MOT-153_nov_91.631%
- SPN193 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.000%
- SPN208 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.719%
- SPN217 Fusobacterium perfoetens_nov_91.126%
- SPN219 Actinidia eriantha_nov_97.011%
- SPN29 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.423%
- SPN33 Duncaniella freteri_nov_88.934%
- SPN35 Yaniella halotolerans_nov_97.040%
- SPN40 Oscillospiraceae_[G-1] bacterium_MOT-147_nov_96.674%
- SPN43 Lachnoclostridium [Clostridium] aminophilum_nov_89.792%
- SPN51 Duncaniella freteri_nov_93.699%
- SPN52 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_92.917%
- SPN55 Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.105%
- SPN59 Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.887%
- SPN67 Parafannyhessea umbonata_nov_92.161%
- SPN82 Turicibacter sanguinis_nov_95.923%
- SPN83 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.198%
- SPP1 Staphylococcus saprophyticus_xylosus
- SPP2 Sphingomonas aquatilis_melonis
- SPP3 Lactacaseibacillus_Lactobacillus casei_rhamnosus
- SPPN3 Faecalicatena multispecies_sppn3_2_nov_92.067%