

Species

| | | |
|---|---|---|
| SP1 Eubacteriales_[G-4] bacterium_MOT-164 | SP57 Dubosiella newyorkensis | SPN181 Duncaniella freteri_nov_89.718% |
| SP10 Ligilactobacillus animalis | SP58 Eubacteriales_[G-2] bacterium_MOT-162 | SPN187 Oscillospiraceae_[G-6] bacterium_MOT-153_nov_91.631% |
| SP12 Streptomyces aculeolatus | SP6 Staphylococcus saprophyticus | SPN192 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.000% |
| SP14 Secundilactobacillus paracollinoides | SP61 Anaerococcus sp._HMT_290 | SPN204 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.719% |
| SP15 Mollicutes_[G-1] bacterium_MOT-186 | SP7 Cutibacterium acnes | SPN216 Fusobacterium perfoetens_nov_91.126% |
| SP18 Staphylococcus ureilyticus | SP72 Stenotrophomonas [Pseudomonas] hibiscicola | SPN218 Actinidia eriantha_nov_97.011% |
| SP19 Atopostipes sp._MOT-201 | SP73 Psychrobacter alimentarius | SPN29 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.423% |
| SP2 Actinidia eriantha | SP76 Akkermansia muciniphila | SPN33 Duncaniella freteri_nov_88.934% |
| SP20 Ralstonia sp._HMT_406 | SP77 Bradyrhizobium pachyrhizi | SPN35 Yarniella halotolerans_nov_97.040% |
| SP23 Cutibacterium granulosum | SP79 Pelomonas saccharophila | SPN39 Oscillospiraceae_[G-1] bacterium_MOT-147_nov_96.674% |
| SP26 Mammaliococcus lentus | SP8 Corynebacterium ammoniagenes | SPN43 Lachnoclostridium [Clostridium] aminophilum_nov_89.792% |
| SP27 Clostridium disporicum | SP80 Lachnospiraceae_[G-14] bacterium_MOT-185 | SPN49 Duncaniella freteri_nov_93.699% |
| SP28 Staphylococcus equorum | SP81 Pseudomonas helleri | SPN50 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_92.917% |
| SP29 Moraxella osloensis | SP88 Bifidobacterium pseudolongum | SPN59 Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.887% |
| SP3 Corynebacterium stationis | SP89 Enterococcus faecalis | SPN65 Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.105% |
| SP30 Limosilactobacillus reuteri | SP9 Triticum aestivum | SPN66 Parafannyhessea umbonata_nov_92.161% |
| SP31 Ligilactobacillus murinus | SP90 Lactobacillus johnsonii | SPN81 Turicibacter sanguinis_nov_95.923% |
| SP34 Leptothrix sp._HMT_025 | SPN101 Fusobacterium varium_nov_96.696% | SPN82 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.198% |
| SP38 Streptococcus thermophilus | SPN106 Duncaniella freteri_nov_93.293% | SPN9 Oribacterium parvum_nov_89.770% |
| SP4 Jeotgalicoccus halotolerans | SPN116 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.867% | SPP1 Staphylococcus saprophyticus_xylosus |
| SP42 Stenotrophomonas maltophilia | SPN117 Enterococcus faecalis_nov_95.825% | SPP3 Sphingomonas aquatilis_melonis |
| SP44 Enterococcus gallinarum | SPN148 Duncaniella freteri_nov_89.775% | SPP4 Lactocaseibacillus_Lactobacillus casei_rhamnosus |
| SP49 Delftia acidovorans | SPN16 Fusicatenibacter saccharivorans_nov_90.526% | SPPN3 Faecalicatena multispecies_sppn3_2_nov_92.067% |
| SP52 Lachnospiraceae_[G-11] bacterium_MOT-177 | SPN170 Alistipes senegalensis_nov_93.443% | |
| SP54 Massilia aurea | SPN177 Peptococcus sp._HMT_168_nov_84.866% | |