

Species

SP1 Streptomyces aculeolatus	SP54 Leptothrix sp._HMT_025	SPN178 Peptococcus sp._HMT_168_nov_84.866%
SP10 Staphylococcus saprophyticus	SP56 Lactobacillus johnsonii	SPN181 Duncaniella freteri_nov_89.718%
SP12 Triticum aestivum	SP64 Anaerococcus sp._HMT_290	SPN188 Oscillospiraceae_[G-6] bacterium_MOT-153_nov_91.631%
SP14 Corynebacterium stationis	SP65 Staphylococcus equorum	SPN193 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.000%
SP17 Stenotrophomonas maltophilia	SP68 Pseudomonas helleri	SPN208 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.719%
SP18 Cutibacterium acnes	SP7 Jeotgaliococcus halotolerans	SPN217 Fusobacterium perfoetens_nov_91.126%
SP2 Ligilactobacillus murinus	SP70 Stenotrophomonas [Pseudomonas] hibiscicola	SPN219 Actinidia eriantha_nov_97.011%
SP23 Lachnospiraceae_[G-11] bacterium_MOT-177	SP73 Dubosiella newyorkensis	SPN29 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.423%
SP24 Ligilactobacillus animalis	SP74 Eubacteriales_[G-2] bacterium_MOT-162	SPN33 Duncaniella freteri_nov_88.934%
SP25 Mammaliococcus lentus	SP76 Ralstonia sp._HMT_406	SPN35 Yaniella halotolerans_nov_97.040%
SP26 Lachnospiraceae_[G-14] bacterium_MOT-185	SP78 Psychrobacter alimentarius	SPN40 Oscillospiraceae_[G-1] bacterium_MOT-147_nov_96.674%
SP27 Corynebacterium ammoniagenes	SP8 Eubacteriales_[G-4] bacterium_MOT-164	SPN43 Lachnoclostridium [Clostridium] aminophilum_nov_89.792%
SP28 Streptococcus thermophilus	SP80 Bradyrhizobium pachyrhizi	SPN51 Duncaniella freteri_nov_93.699%
SP33 Delftia acidovorans	SP83 Akkermansia muciniphila	SPN52 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_92.917%
SP34 Cutibacterium granulosum	SP84 Secundilactobacillus paracollinoides	SPN55 Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.105%
SP35 Massilia aurea	SP9 Staphylococcus ureilyticus	SPN59 Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.887%
SP39 Actinidia eriantha	SP93 Bifidobacterium pseudolongum	SPN67 Parafannyhessea umbonata_nov_92.161%
SP40 Enterococcus gallinarum	SPN10 Oribacterium parvum_nov_89.770%	SPN82 Turicibacter sanguinis_nov_95.923%
SP41 Moraxella osloensis	SPN102 Fusobacterium varium_nov_96.696%	SPN83 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.198%
SP42 Mollicutes_[G-1] bacterium_MOT-186	SPN107 Duncaniella freteri_nov_93.293%	SPP1 Staphylococcus saprophyticus_xylosus
SP45 Atopostipes sp._MOT-201	SPN117 Enterococcus faecalis_nov_95.825%	SPP2 Sphingomonas aquatilis_melonis
SP47 Enterococcus faecalis	SPN118 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.867%	SPP3 Lacticaseibacillus_Lactobacillus casei_rhamnosus
SP49 Pelomonas saccharophila	SPN149 Duncaniella freteri_nov_89.775%	SPN3 Faecalicatena multispecies_sppn3_2_nov_92.067%
SP51 Limosilactobacillus reuteri	SPN15 Fusicatenibacter saccharivorans_nov_90.526%	
SP53 Clostridium disporicum	SPN173 Alistipes senegalensis_nov_93.443%	