

Species

SP1 Staphylococcus saprophyticus	SP48 Atopostipes sp._MOT-201	SPN179 Peptococcus sp._HMT_168_nov_84.866%
SP11 Lachnospiraceae_[G-14] bacterium_MOT-185	SP54 Massilia aurea	SPN183 Duncaniella freteri_nov_89.718%
SP13 Enterococcus faecalis	SP59 Akkermansia muciniphila	SPN189 Oscillospiraceae_[G-6] bacterium_MOT-153_nov_91.631%
SP14 Corynebacterium ammoniagenes	SP6 Pelomonas saccharophila	SPN194 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.000%
SP16 Staphylococcus equorum	SP61 Ralstonia sp._HMT_406	SPN210 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.719%
SP18 Mammaliicoccus lentus	SP65 Pseudomonas helleri	SPN218 Fusobacterium perfoetens_nov_91.126%
SP20 Bradyrhizobium pachyrrhizi	SP68 Cutibacterium granulosum	SPN220 Actinidia eriantha_nov_97.011%
SP22 Triticum aestivum	SP69 Delftia acidovorans	SPN28 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.423%
SP24 Moraxella osloensis	SP7 Ligilactobacillus animalis	SPN33 Duncaniella freteri_nov_88.934%
SP25 Eubacteriales_[G-4] bacterium_MOT-164	SP72 Lactobacillus johnsonii	SPN36 Yaniella halotolerans_nov_97.040%
SP27 Jeotgaliococcus halotolerans	SP77 Stenotrophomonas [Pseudomonas] hibiscicola	SPN38 Oscillospiraceae_[G-1] bacterium_MOT-147_nov_96.674%
SP3 Lachnospiraceae_[G-11] bacterium_MOT-177	SP78 Eubacteriales_[G-2] bacterium_MOT-162	SPN43 Lachnoclostridium [Clostridium] aminophilum_nov_89.792%
SP30 Actinidia eriantha	SP8 Leptothrix sp._HMT_025	SPN50 Duncaniella freteri_nov_93.699%
SP31 Limosilactobacillus reuteri	SP81 Dubosiella newyorkensis	SPN51 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_92.917%
SP32 Clostridium disporicum	SP85 Streptomyces aculeolatus	SPN61 Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.887%
SP33 Enterococcus gallinarum	SP9 Secundilactobacillus paracollinoides	SPN64 Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.105%
SP34 Cutibacterium acnes	SP92 Bifidobacterium pseudolongum	SPN65 Parafannyhessea umbonata_nov_92.161%
SP35 Mollicutes_[G-1] bacterium_MOT-186	SP94 Anaerococcus sp._HMT_290	SPN8 Oribacterium parvum_nov_89.770%
SP36 Corynebacterium stationis	SPN102 Fusobacterium varium_nov_96.696%	SPN81 Turicibacter sanguinis_nov_95.923%
SP39 Staphylococcus ureilyticus	SPN107 Duncaniella freteri_nov_93.293%	SPN82 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.198%
SP4 Psychrobacter alimentarius	SPN118 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.867%	SPP1 Staphylococcus saprophyticus_xylosus
SP40 Ligilactobacillus murinus	SPN127 Enterococcus faecalis_nov_95.825%	SPP3 Sphingomonas aquatilis_melonis
SP41 Streptococcus thermophilus	SPN15 Fusicatenibacter saccharivorans_nov_90.526%	SPPN3 Faecalicaltena multispecies_sppn3_2_nov_92.067%
SP45 Stenotrophomonas maltophilia	SPN150 Duncaniella freteri_nov_89.775%	
SP47 Lactocaseibacillus rhamnosus	SPN173 Alistipes senegalensis_nov_93.443%	