

Species

SP1 Triticum aestivum	SP56 Actinidia eriantha	SPN179 Peptococcus sp._HMT_168_nov_84.866%
SP10 Delftia acidovorans	SP57 Atopostipes sp._MOT-201	SPN183 Duncaniella freteri_nov_89.718%
SP11 Enterococcus gallinarum	SP59 Mollicutes_[G-1] bacterium_MOT-186	SPN190 Oscillospiraceae_[G-6] bacterium_MOT-153_nov_91.631%
SP13 Staphylococcus saprophyticus	SP6 Ralstonia sp._HMT_406	SPN194 Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.000%
SP15 Ligilactobacillus murinus	SP63 Anaerococcus sp._HMT_290	SPN208 Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.719%
SP17 Pseudomonas helleri	SP65 Streptomyces aculeolatus	SPN218 Fusobacterium perfoetens_nov_91.126%
SP18 Eubacteriales_[G-4] bacterium_MOT-164	SP66 Lactobacillus johnsonii	SPN220 Actinidia eriantha_nov_97.011%
SP19 Stenotrophomonas maltophilia	SP67 Bifidobacterium pseudolongum	SPN28 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.423%
SP2 Psychrobacter alimentarius	SP7 Staphylococcus ureilyticus	SPN32 Yaniella halotolerans_nov_97.040%
SP24 Eubacteriales_[G-2] bacterium_MOT-162	SP70 Lactocaseibacillus rhamnosus	SPN33 Duncaniella freteri_nov_88.934%
SP26 Clostridium disporicum	SP72 Akkermansia muciniphila	SPN39 Oscillospiraceae_[G-1] bacterium_MOT-147_nov_96.674%
SP27 Stenotrophomonas [Pseudomonas] hibiscicola	SP75 Leptothrix sp._HMT_025	SPN45 Lachnoclostridium [Clostridium] aminophilum_nov_89.792%
SP29 Cutibacterium acnes	SP8 Limosilactobacillus reuteri	SPN49 Duncaniella freteri_nov_93.699%
SP33 Lachnospiraceae_[G-11] bacterium_MOT-177	SP80 Moraxella osloensis	SPN50 Oscillospiraceae_[G-3] bacterium_MOT-150_nov_92.917%
SP34 Cutibacterium granulosum	SP81 Secundilactobacillus paracollinoides	SPN55 Muribaculaceae_[G-1] bacterium_MOT-129_nov_88.105%
SP35 Corynebacterium stationis	SP88 Pelomonas saccharophila	SPN60 Muribaculaceae_[G-1] bacterium_MOT-129_nov_85.887%
SP36 Streptococcus thermophilus	SP9 Ligilactobacillus animalis	SPN66 Parafannyhessea umbonata_nov_92.161%
SP39 Staphylococcus equorum	SP94 Bradyrhizobium pachyrhizi	SPN81 Turicibacter sanguinis_nov_95.923%
SP4 Massilia aurea	SPN102 Fusobacterium varium_nov_96.696%	SPN82 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.198%
SP41 Jeotgalicoccus halotolerans	SPN107 Duncaniella freteri_nov_93.293%	SPN9 Oribacterium parvum_nov_89.770%
SP42 Mammaliicoccus lentus	SPN116 Muribaculaceae_[G-2] bacterium_MOT-104_nov_88.867%	SPP1 Staphylococcus saprophyticus_xylosus
SP44 Lachnospiraceae_[G-14] bacterium_MOT-185	SPN117 Enterococcus faecalis_nov_95.825%	SPP3 Sphingomonas aquatilis_melonis
SP50 Dubosiella newyorkensis	SPN15 Fusicatenibacter saccharivorans_nov_90.526%	SPPN3 Faecalicatena multispecies_sppn3_2_nov_92.067%
SP51 Corynebacterium ammoniagenes	SPN150 Duncaniella freteri_nov_89.775%	
SP53 Enterococcus faecalis	SPN173 Alistipes senegalensis_nov_93.443%	