

Species

SP1 Enterococcus faecalis	SP46 Corynebacterium aurimucosum	SPN19 Corynebacterium coyleae_nov_94.908%
SP10 Staphylococcus hominis	SP47 Jeotgallcoccus halotolerans	SPN2 Fusobacterium nucleatum_nov_91.853%
SP100 Fusobacterium canifelinum	SP48 Corynebacterium jeikeium	SPN20 Enterococcus faecalis_nov_95.255%
SP101 Klebsiella aerogenes	SP49 Triticum aestivum	SPN21 Ligilactobacillus murinus_nov_95.911%
SP102 Bifidobacterium breve	SP5 Bifidobacterium longum	SPN22 Faecalibaculum rodentium_nov_91.098%
SP103 Glutamicibacter protophormiae	SP50 Faecalibaculum rodentium	SPN23 Enterobacter mori_nov_89.749%
SP104 Veillonella sp._HMT_780	SP51 Bacillus subtilis	SPN24 Sediminibacterium roseum_nov_92.115%
SP105 Parasutterella excrementihominis	SP52 Corynebacterium afermentans	SPN25 Rheinheimera nanhaiensis_nov_86.051%
SP106 Corynebacterium coyleae	SP53 Salmonella enterica	SPN26 Corynebacterium falsenii_nov_91.304%
SP107 Bradyrhizobium frederickii	SP56 Peribacillus frigitolerans	SPN27 Duncaniella freteri_nov_90.977%
SP108 Glutamicibacter soli	SP57 Rhodococcus qingshengii	SPN28 Bifidobacterium pseudolongum_nov_94.646%
SP109 Tepidiphilus succinatimandens	SP58 Streptococcus thoraltensis	SPN29 Lachnospiraceae_[G-1] bacterium_MOT-176_nov_95.136%
SP11 Staphylococcus aureus	SP59 Streptococcus lactarius	SPN3 Turicibacter sanguinis_nov_95.437%
SP110 Cutibacterium avidum	SP6 Acinetobacter pittii	SPN30 Kocuria indica_nov_94.872%
SP111 Corynebacterium matruchotii	SP66 Actinidia eriantha	SPN31 Phycocomes zhengii_nov_88.764%
SP112 Hathewayia histolytica	SP67 Enterococcus casseliflavus	SPN32 Photobacterium frigidophilum_nov_83.878%
SP113 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870	SP68 Staphylococcus warneri	SPN33 Streptococcus thoraltensis_nov_95.652%
SP114 Neisseria mucosa	SP69 Bradyrhizobium japonicum	SPN34 Bradyrhizobium valentinum_nov_90.985%
SP115 Capnocytophaga sputigena	SP7 Staphylococcus pasteuri	SPN35 Adlercreutzia caecimuris_nov_92.843%
SP116 Shigella sonnei	SP70 Acinetobacter johnsonii	SPN36 Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.427%
SP117 Escherichia coli	SP71 Enterobacter mori	SPN37 Cutibacterium avidum_nov_95.072%
SP118 Corynebacterium mastitidis	SP72 Tannerella serpentiniformis	SPN38 Enterococcus gallinarum_nov_91.392%
SP121 Leptothrix sp._HMT_025	SP73 Mammaliicoccus sciuri	SPN39 Granulicatella elegans_nov_84.142%
SP122 Faecalicatena fissicatena	SP74 Shigella flexneri	SPN4 Weizmannia ginsengihumi_nov_96.346%
SP123 Anaerostipes caccae	SP75 Corynebacterium kroppenstedtii	SPN40 Eggerthella timonensis_nov_88.867%
SP124 Bacteroides thetaiotaomicron	SP76 Corynebacterium pilbarensis	SPN41 Corynebacterium glyciniphilum_nov_88.528%
SP125 Corynebacterium propinquum	SP77 Mammaliicoccus lentus	SPN42 Enterococcus faecalis_nov_96.175%
SP14 Enterobacter hormaechei	SP78 Mediterraneibacter faecis	SPN43 Corynebacterium argentoratensis_nov_95.850%
SP15 Bradyrhizobium valentinum	SP79 Klebsiella michiganensis	SPN44 Daejeonella oryzae_nov_85.551%
SP16 Ligilactobacillus murinus	SP8 Escherichia fergusonii	SPN45 Thermodesulfobium acidiphilum_nov_81.729%
SP17 Janibacter cremeus	SP80 Bifidobacterium bifidum	SPN46 Arthrosira platensis_nov_88.987%
SP18 Staphylococcus caprae	SP81 Streptococcus chosunense	SPN47 Sporosarcina sp._MOT-205_nov_96.360%
SP19 Klebsiella pneumoniae	SP82 Enterobacter kobei	SPN48 Brevibacterium paucivorans_nov_97.368%
SP2 Streptococcus danieliae	SP83 Massilia arenae	SPN49 Streptococcus danieliae_nov_95.594%
SP20 Varibaculum cambriense	SP84 Actinomyces sp._HMT_525	SPN5 Mediterraneibacter glycyrrhizinilyticus_nov_95.367%
SP21 Staphylococcus capitis	SP85 Lawsonella clevelandensis	SPN50 Clostridium saccharoperbutylacetonicum_nov_95.800%
SP22 Corynebacterium tuberculoostearicum	SP86 Sphingomonas leidyi	SPN51 Microbacterium saccharophilum_nov_76.981%
SP23 Fusobacterium nucleatum	SP87 Streptococcus oralis	SPN52 Ligilactobacillus murinus_nov_97.020%
SP24 Facklamia languida	SP88 Streptococcus mutans	SPN53 Duncaniella freteri_nov_90.075%
SP25 Citrobacter murlinae	SP89 Acinetobacter septicus	SPN54 Ralstonia solanacearum_nov_95.922%
SP26 Rothia mucilaginosa	SP9 Bacillus halotolerans	SPN55 Frankia discariae_nov_81.481%
SP27 Actinomyces sp._HMT_448	SP90 Abiotrophia defectiva	SPN56 Ligilactobacillus murinus_nov_96.667%
SP28 Pantoea allii	SP91 Acinetobacter radioresistens	SPN57 Faecalibaculum rodentium_nov_96.571%
SP29 Neisseria sicca	SP92 Staphylococcus saprophyticus	SPN58 Lactobacillus gasseri_nov_94.075%
SP3 Streptococcus parasanguinis_clade_411	SP93 Pseudoramibacter alactolyticus	SPN59 Enterococcus faecalis_nov_94.118%
SP30 Streptococcus sp._HMT_066	SP94 Corynebacterium appendicis	SPN6 Streptococcus thoraltensis_nov_94.361%
SP31 Methylophilus leisingeri	SP95 Corynebacterium pseudodiphtheriticum	SPN7 Cutibacterium acnes_nov_94.643%
SP32 Neisseriaceae_[G-1] bacterium_HMT_174	SP96 Robinsoniella peoriensis	SPN8 Prauserella oleivorans_nov_80.113%
SP33 Cutibacterium acnes	SP97 Corynebacterium falsenii	SPN9 Enterococcus faecalis_nov_96.685%
SP34 Weizmannia ginsengihumi	SP98 Lactiplantibacillus plantarum	SPP1 Bradyrhizobium mercantei_viridifuturi
SP35 Bradyrhizobium lupini	SP99 Prevotella sp._HMT_314	SPP2 Staphylococcus argenteus_aureus_roterodami
SP36 Enterococcus gallinarum	SPN1 Mammaliicoccus sciuri_nov_96.743%	SPP3 Staphylococcus equorum_ureilyticus
SP37 Acinetobacter lwoffii	SPN10 Faecalibaculum rodentium_nov_93.103%	SPP4 Enterobacter_Leclercia_adeccarboxylata_cloacae
SP38 Enterobacter cloacae	SPN11 Olsenella phocaeensis_nov_91.977%	SPP5 Streptococcus infantis_infantis_clade_638
SP4 Streptococcus mitis	SPN12 Anaerostipes caccae_nov_96.266%	SPP6 Bacillus albus_cereus_luti_nitratireducens_paramycoides_tro...(6 species)
SP40 Prevotella oralis	SPN13 Longibaculum muris_nov_90.607%	SPPN1 Staphylococcus multispecies_sppn1_3_nov_95.362%
SP41 Corynebacterium durum	SPN14 Corynebacterium matruchotii_nov_86.497%	SPPN2 Staphylococcus multispecies_sppn2_3_nov_96.987%
SP42 Bifidobacterium pseudolongum	SPN15 Limosilactobacillus reuteri_nov_93.571%	SPPN3 Terrimonas multispecies_sppn3_2_nov_90.822%
SP43 Cutibacterium namnetense	SPN16 Klebsiella pneumoniae_nov_91.098%	SPPN4 Gloebacter multispecies_sppn4_2_nov_84.254%
SP44 Microbacterium maritipicum	SPN17 Zavartzinia aquatilis_nov_84.906%	
SP45 Corynebacterium ureicelivorans	SPN18 Robinsoniella peoriensis_nov_94.563%	