

## Species

SP1 Enterococcus faecalis	SP4 Streptococcus mitis	SP81 Streptococcus chosunense
SP10 Staphylococcus hominis	SP41 Corynebacterium durum	SP82 Enterobacter kobei
SP101 Klebsiella aerogenes	SP42 Bifidobacterium pseudolongum	SP83 Massilia arenae
SP104 Veillonella sp._HMT_780	SP43 Cutibacterium namnetense	SP85 Lawsonella clevelandensis
SP105 Parasutterella excrementihominis	SP44 Microbacterium maritipicum	SP86 Sphingomonas leidyi
SP106 Corynebacterium coyleae	SP45 Corynebacterium ureicelerivorans	SP87 Streptococcus oralis
SP108 Glutamicibacter soli	SP46 Corynebacterium aurimucosum	SP88 Streptococcus mutans
SP109 Tepidiphilus succinatimandens	SP47 Jeotgaliococcus halotolerans	SP89 Acinetobacter septicus
SP110 Cutibacterium avidum	SP48 Corynebacterium jeikeium	SP9 Bacillus halotolerans
SP111 Corynebacterium matruchotii	SP49 Triticum aestivum	SP90 Abiotrophia defectiva
SP114 Neisseria mucosa	SP5 Bifidobacterium longum	SP92 Staphylococcus saprophyticus
SP14 Enterobacter hormaechei	SP50 Faecalibaculum rodentium	SP93 Pseudoramibacter alactolyticus
SP15 Bradyrhizobium valentinum	SP51 Bacillus subtilis	SP95 Corynebacterium pseudodiphtheriticum
SP16 Ligilactobacillus murinus	SP52 Corynebacterium afermentans	SP97 Corynebacterium falsenii
SP17 Janibacter cremeus	SP53 Salmonella enterica	SP98 Lactiplantibacillus plantarum
SP18 Staphylococcus caprae	SP56 Peribacillus frigoritolerans	SP99 Prevotella sp._HMT_314
SP19 Klebsiella pneumoniae	SP57 Rhodococcus qingshengii	SPN30 Kocuria indica_nov_94.872%
SP2 Streptococcus danieliae	SP58 Streptococcus thoraltensis	SPN43 Corynebacterium argentoratense_nov_95.850%
SP20 Varibaculum cambriense	SP59 Streptococcus lactarius	SPN44 Daejeonella oryzae_nov_85.551%
SP21 Staphylococcus capitis	SP6 Acinetobacter pittii	SPN45 Thermodesulfobium acidiphilum_nov_81.729%
SP22 Corynebacterium tuberculoostearicum	SP66 Actinidia eriantha	SPN46 Arthrospira platensis_nov_88.987%
SP23 Fusobacterium nucleatum	SP67 Enterococcus casseliflavus	SPN47 Sporosarcina sp._MOT-205_nov_96.360%
SP24 Facklamia languida	SP68 Staphylococcus warneri	SPN48 Brevibacterium paucivorans_nov_97.368%
SP26 Rothia mucilaginosa	SP69 Bradyrhizobium japonicum	SPN49 Streptococcus danieliae_nov_95.594%
SP27 Actinomyces sp._HMT_448	SP7 Staphylococcus pasteurii	SPN50 Clostridium saccharoperbutylacetonicum_nov_95.800%
SP28 Pantoea allii	SP70 Acinetobacter johnsonii	SPN51 Microbacterium saccharophilum_nov_76.981%
SP3 Streptococcus parasanguinis_clade_411	SP71 Enterobacter mori	SPN53 Duncaniella freteri_nov_90.075%
SP30 Streptococcus sp._HMT_066	SP72 Tannerella serpentiniformis	SPN54 Ralstonia solanacearum_nov_95.922%
SP31 Methylophilus leisingeri	SP73 Mammaliococcus sciuri	SPN55 Frankia discariae_nov_81.481%
SP32 Neisseriaceae_[G-1] bacterium_HMT_174	SP74 Shigella flexneri	SPP2 Staphylococcus argenteus_aureus_roterodami
SP33 Cutibacterium acnes	SP75 Corynebacterium kroppenstedtii	SPP4 Enterobacter_Leclercia adecarboxylata_cloacae
SP34 Weizmannia ginsengihumi	SP76 Corynebacterium pilbarensis	SPP5 Streptococcus infantis_infantis_clade_638
SP35 Bradyrhizobium lupini	SP77 Mammaliococcus lentus	SPN6 Bacillus albus_cereus_luti_nitratireducens_paramycoides_tro...(6 species)
SP36 Enterococcus gallinarum	SP78 Mediterraneibacter faecis	SPPN3 Terrimonas multispecies_sppn3_2_nov_90.822%
SP37 Acinetobacter lwofii	SP8 Escherichia fergusonii	SPPN4 Gloeobacter multispecies_sppn4_2_nov_84.254%
SP38 Enterobacter cloacae	SP80 Bifidobacterium bifidum	