

Species

SP1 <i>Enterococcus faecalis</i>	SP4 <i>Streptococcus mitis</i>	SP81 <i>Streptococcus chosunense</i>
SP10 <i>Staphylococcus hominis</i>	SP41 <i>Corynebacterium durum</i>	SP82 <i>Enterobacter kobei</i>
SP101 <i>Klebsiella aerogenes</i>	SP42 <i>Bifidobacterium pseudolongum</i>	SP83 <i>Massilia arenae</i>
SP104 <i>Veillonella</i> sp._HMT_780	SP43 <i>Cutibacterium namnetense</i>	SP85 <i>Lawsonella clevelandensis</i>
SP105 <i>Parasutterella excrementihominis</i>	SP44 <i>Microbacterium maritipicum</i>	SP86 <i>Sphingomonas leidyi</i>
SP106 <i>Corynebacterium coyleae</i>	SP45 <i>Corynebacterium ureicelerivorans</i>	SP87 <i>Streptococcus oralis</i>
SP108 <i>Glutamicibacter soli</i>	SP46 <i>Corynebacterium aurimucosum</i>	SP88 <i>Streptococcus mutans</i>
SP109 <i>Tepidiphilus succinatimandens</i>	SP47 <i>Jeotgaliococcus halotolerans</i>	SP89 <i>Acinetobacter septicus</i>
SP110 <i>Cutibacterium avidum</i>	SP48 <i>Corynebacterium jeikeium</i>	SP9 <i>Bacillus halotolerans</i>
SP111 <i>Corynebacterium matruchotii</i>	SP49 <i>Triticum aestivum</i>	SP90 <i>Abiotrophia defectiva</i>
SP114 <i>Neisseria mucosa</i>	SP5 <i>Bifidobacterium longum</i>	SP92 <i>Staphylococcus saprophyticus</i>
SP14 <i>Enterobacter hormaechei</i>	SP50 <i>Faecalibaculum rodentium</i>	SP93 <i>Pseudoramibacter alactolyticus</i>
SP15 <i>Bradyrhizobium valentinum</i>	SP51 <i>Bacillus subtilis</i>	SP95 <i>Corynebacterium pseudodiphtheriticum</i>
SP16 <i>Ligilactobacillus murinus</i>	SP52 <i>Corynebacterium afermentans</i>	SP97 <i>Corynebacterium falsenii</i>
SP17 <i>Janibacter cremeus</i>	SP53 <i>Salmonella enterica</i>	SP98 <i>Lactiplantibacillus plantarum</i>
SP18 <i>Staphylococcus caprae</i>	SP56 <i>Peribacillus frigoritolerans</i>	SP99 <i>Prevotella</i> sp._HMT_314
SP19 <i>Klebsiella pneumoniae</i>	SP57 <i>Rhodococcus qingshengii</i>	SPN30 <i>Kocuria indica</i> _nov_94.872%
SP2 <i>Streptococcus danieliae</i>	SP58 <i>Streptococcus thoraltensis</i>	SPN43 <i>Corynebacterium argentoratense</i> _nov_95.850%
SP20 <i>Varibaculum cambriense</i>	SP59 <i>Streptococcus lactarius</i>	SPN44 <i>Daejeonella oryzae</i> _nov_85.551%
SP21 <i>Staphylococcus capitis</i>	SP6 <i>Acinetobacter pittii</i>	SPN45 <i>Thermodesulfobium acidiphilum</i> _nov_81.729%
SP22 <i>Corynebacterium tuberculostearicum</i>	SP66 <i>Actinidia eriantha</i>	SPN46 <i>Arthrospira platensis</i> _nov_88.987%
SP23 <i>Fusobacterium nucleatum</i>	SP67 <i>Enterococcus casseliflavus</i>	SPN47 <i>Sporosarcina</i> sp._MOT-205_nov_96.360%
SP24 <i>Facklamia languida</i>	SP68 <i>Staphylococcus warneri</i>	SPN48 <i>Brevibacterium paucivorans</i> _nov_97.368%
SP26 <i>Rothia mucilaginosa</i>	SP69 <i>Bradyrhizobium japonicum</i>	SPN49 <i>Streptococcus danieliae</i> _nov_95.594%
SP27 <i>Actinomyces</i> sp._HMT_448	SP7 <i>Staphylococcus pasteurii</i>	SPN50 <i>Clostridium saccharoperbutylacetonicum</i> _nov_95.800%
SP28 <i>Pantoea allii</i>	SP70 <i>Acinetobacter johnsonii</i>	SPN51 <i>Microbacterium saccharophilum</i> _nov_76.981%
SP3 <i>Streptococcus parasanguinis</i> _clade_411	SP71 <i>Enterobacter mori</i>	SPN53 <i>Duncanella freteri</i> _nov_90.075%
SP30 <i>Streptococcus</i> sp._HMT_066	SP72 <i>Tannerella serpentiniformis</i>	SPN54 <i>Ralstonia solanacearum</i> _nov_95.922%
SP31 <i>Methylophilus leisingeri</i>	SP73 <i>Mammaliococcus sciuri</i>	SPN55 <i>Frankia discariae</i> _nov_81.481%
SP32 <i>Neisseriaceae</i> _ [G-1] <i>bacterium</i> _HMT_174	SP74 <i>Shigella flexneri</i>	SPP2 <i>Staphylococcus argenteus_aureus_roterodami</i>
SP33 <i>Cutibacterium acnes</i>	SP75 <i>Corynebacterium kroppenstedtii</i>	SPP4 <i>Enterobacter_Leclercia adecarboxylata_cloacae</i>
SP34 <i>Weizmannia ginsengihumi</i>	SP76 <i>Corynebacterium pilbarensis</i>	SPP5 <i>Streptococcus infantis_infantis</i> _clade_638
SP35 <i>Bradyrhizobium lupini</i>	SP77 <i>Mammaliococcus lentus</i>	SPN6 <i>Bacillus albus_cereus_luti_nitratireducens_paramycoides_tro</i> ...(6 species)
SP36 <i>Enterococcus gallinarum</i>	SP78 <i>Mediterraneibacter faecis</i>	SPPN3 <i>Terrimonas multispecies_sppn3_2_nov_90.822%</i>
SP37 <i>Acinetobacter lwofii</i>	SP8 <i>Escherichia fergusonii</i>	SPPN4 <i>Gloeobacter multispecies_sppn4_2_nov_84.254%</i>
SP38 <i>Enterobacter cloacae</i>	SP80 <i>Bifidobacterium bifidum</i>	