

cies

SP103 Bifidobacterium scarauvii	SP234 Staphylococcus similaris	SPN178 Gemella haemolysans nov_97.425%
SP104 Bifidobacterium longum	SP235 Anaerococcus obesiensis	SPN18 Enterococcus faecalis nov_96.788%
SP105 Prevotella bivia	SP238 Neisseria flavescens	SPN204 Gemella haemolysans nov_94.030%
SP106 Finnegolia magna	SP239 Peptostreptococcaceae [XII][G-6] [Eubacterium]_nodatum	SPN233 Peptoniphilus coxii nov_97.955%
SP108 Gemella haemolysans	SP240 Limosilactobacillus mucosae	SPN27 Veillonella atypica nov_96.137%
SP109 Collinsella aerofaciens	SP243 Serratia marcescens	SPN301 Clostridium saccharobutylicum nov_96.110%
SP110 Peptoniphilus grossensis	SP244 Actinomyces massiliensis	SPN329 Clostridium chromiireducens nov_97.959%
SP111 Anaerococcus octavius	SP245 Klebsiella variicola	SPN36 Enterococcus faecalis nov_89.507%
SP118 Veillonella atypica	SP250 Eggerthella lenta	SPN45 Mediterraneibacter [Ruminococcus]_gnavus nov_92.601%
SP119 Solobacterium moorei	SP252 Erysipelatoclostridium [Clostridium]_innocuum	SPN496 Megamonas funiformis nov_97.849%
SP120 Serratia nematodiphila	SP257 Gardnerella vaginalis	SPN525 Clostridium saccharobutylicum nov_96.606%
SP121 Bacteroides fragilis	SP265 Prevotella histicola	SPN56 Saccharibacteria (TM7) [G-1] bacterium_HMT_346 nov_97.964%
SP122 Clostridium chromiireducens	SP266 Anaerococcus sp._HMT_294	SPN65 Clostridium tarantellae nov_97.052%
SP123 Phocaecicola vulgatus	SP284 Corynebacterium Kroppenstedtii	SPN86 Clostridium saccharobutylicum nov_97.052%
SP124 Staphylococcus saccharolyticus	SP287 Hymenobacter rigui	SPN9 Rothia mucilaginosa nov_97.991%
SP125 Clostridium paraputrificum	SP288 Bifidobacterium dentium	SPP1 Schaalia odontolyticus sp._HMT_180
SP126 Anaerococcus vaginalis	SP294 Peptostreptococcaceae [XII][G-1] [Eubacterium]_sulci	SPP10 Streptococcus parasanguinis clade_411 sinensis sp._HMT_056
SP127 Clostridium tertium	SP295 Stomatobaculum sp._HMT_097	SPP121 Streptococcus australis rubneri sp._HMT_066
SP130 Gemella sanguinis	SP298 Corynebacterium pilbarensis	SPP127 Klebsiella pneumoniae quasipneumoniae
SP131 Schaalia sp._HMT_172	SP305 Saccharibacteria (TM7) [G-6] bacterium_HMT_870	SPP141 Staphylococcus capitis_caprae
SP134 Rothia aeria	SP307 Dermabacter hominis	SPP142 Staphylococcus capitis_caprae epidermidis
SP135 Granulicatella adiacens	SP309 Peptostreptococcaceae [XII][G-9] [Eubacterium]_brachy	SPP148 Enterobacter asburiae cancerogenus cloacae hormaechei
SP136 Haemophilus parainfluenzae	SP312 Capnocytophaga leadbetteri	SPP149 Streptococcus cristatus cristatus clade_578
SP137 Peptostreptococcus anaerobius	SP325 Actinomyces israelii	SPP155 Streptococcus lactarius_peroris
SP138 Peptoniphilus lacrimalis	SP327 Actinomyces ihuae	SPP16 Streptococcus cristatus cristatus clade_578 downii gwangjuense_i...(18 species)
SP139 Anaerococcus lactolyticus	SP329 Veillonella denticariosi	SPP161 Streptococcus salivarius vestibularis
SP140 Paratractidigestivbacter faecalis	SP341 Bacteroides caccae	SPP165 Escherichia Shigella coli fergusonii flexneri sonnei
SP144 Stutzerimonas [Pseudomonas]_urumqiensis	SP342 Leptotrichia sp._HMT_215	SPP17 Streptococcus cristatus downii gwangjuense_infantis_infantis_cla...(17 species)
SP145 Bacteroides faecis	SP344 Faecalimonas umbilicata	SPP172 Enterococcus casseliflavus_gallinarum
SP146 Mediterraneibacter [Ruminococcus]_gnavus	SP353 Oribacterium parvum	SPP18 Hoefflea Mesorhizobium camelthorni_halophila
SP147 Cutibacterium acnes	SP354 Porphyromonas endodontalis	SPP180 Staphylococcus argenteus aureus Toterodami_simiae
SP149 Staphylococcus lugdunensis	SP357 Winkia neuii	SPP182 Bacillus inaquosorum spizizenii_subtilis_tequilensis
SP150 Corynebacterium amycolatum	SP359 Actinomyces naeslundii	SPP19 Anaerococcus prevotii_tetradius
SP152 Serratia surfactantfaciens	SP365 Ruminococcaceae [G-1] bacterium_HMT_075	SPP21 Anaerococcus mediterraneensis_murdochii
SP153 Capnocytophaga sputigena	SP367 Cutibacterium avidum	SPP25 Bifidobacterium adolescentis_faecale
SP154 Prevotella corporis	SP370 Leptotrichia sp._HMT_392	SPP28 Peptoniphilus gorbachii_lacydonensis sp._HMT_187
SP155 Acinetobacter baumannii	SP400 Bacteroides stercoris	SPP33 Streptococcus parasanguinis clade_411 sp._HMT_057
SP156 Veillonella dispar	SP41 Veillonella rogosae	SPP37 Fusobacterium nucleatum_nucleatum_subsp._vincentii
SP157 Gemella morbillorum	SP42 Anaerococcus hydrogenalis	SPP39 Streptococcus parasanguinis parasanguinis clade_721 sp._HMT_066
SP159 Porphyromonas pasteri	SP43 Anaerostipes caccae	SPP41 Atopobium Lancefieldella parvula_parvulum
SP160 Citrobacter koseri	SP430 Corynebacterium durum	SPP42 Colibacter_Megasphaera massiliensis sp._HMT_123
SP162 Lactobacillus gasseri	SP44 Schaalia lingnae [Not_Validly_Published]	SPP44 Atopobium_Fannyhessea vaginae
SP173 Faecalibacterium prausnitzii	SP45 Proteus mirabilis	SPP46 Pseudomonas oryzaehabitans_psychrotolerans
SP176 Pinus sylvestris	SP451 Ruminococcaceae [G-2] bacterium_HMT_085	SPP47 Clostridium disporicum_saudiense
SP179 Prevotella timonensis	SP458 Bilophila wadsworthia	SPP50 Lactococcus cremoris_lactis
SP182 Bifidobacterium bifidum	SP49 Klebsiella pneumoniae	SPP52 Fusobacterium canifelinum_nucleatum_nucleatum_subsp._polymorphum
SP183 Rothia dentocariosa	SP51 Rothia mucilaginosa	SPP53 Hungatella effluvii_hathewayi
SP184 Bergeyella sp._HMT_931	SP52 Enterobacter bugandensis	SPP54 Bifidobacterium catenulatum_pseudocatenulatum
SP186 Parabacteroides distansoni	SP53 Enterococcus faecalis	SPP55 Olsenella sp._HMT_939_urininfantis
SP187 Pantoea conspicua	SP54 Fusobacterium nucleatum	SPP56 Streptococcus australis_rubneri sp._HMT_066 sp._HMT_074
SP189 Actinomyces oris	SP55 Streptococcus parasanguinis clade_411	SPP61 Staphylococcus argensis_pettenkoferi
SP192 Prevotella melaninogenica	SP56 Staphylococcus capitis	SPP63 Afipia_Bradyrhizobium archetypum_australiense_broomeae_elkanii_embraepens...(18 species)
SP193 Bacteroides thetaiotaomicron	SP58 Clostridium perfringens	SPP64 Haemophilus seminalis sp._HMT_908
SP194 Peptoniphilus coxii	SP63 Staphylococcus hominis	SPP68 Neisseria flava_macacae_mucosa_sicca
SP195 Actinomyces urogenitalis	SP64 Streptococcus lactarius	SPP76 Acinetobacter_Prolinoborus fasciculus_lwoffii
SP196 Schaalia odontolyticus	SP65 Streptococcus peroris	SPP79 Haemophilus haemolyticus sp._HMT_036
SP197 Varibaculum anthropi	SP66 Megamonas funiformis	SPP8 Streptococcus parasanguinis clade_411 sinensis
SP198 Parvimonas micra	SP67 Enterobacter mori	SPP83 Lactobacillus crispatus_gallinarum
SP199 Phocaecicola dorei	SP68 Streptococcus sp._HMT_056	SPP85 Veillonella parvula_tobetsuensis
SP2 Veillonella parvula	SP75 Staphylococcus haemolyticus	SPP87 Veillonella parvula_rogosae_tobetsuensis
SP203 Streptococcus sanguinis	SP79 Abiotrophia defectiva	SPP90 Lachnoanaerobaculum gingivalis_umeaense
SP204 Stenotrophomonas maltophilia	SP80 Veillonella sp._HMT_780	SPP98 Parvimonas_Peptostreptococcus Candidatus massiliensis sp._HMT_110 sp._HMT_393
SP205 Varibaculum cambriense	SP81 Blautia wexlerae	SPPN104 Enterococcus multispecies_sppn104_2 nov_96.360%
SP206 Oribacterium sinus	SP82 Stenotrophomonas geniculata	SPPN109 Schaalia multispecies_sppn109_2 nov_87.419%
SP211 Clostridium butyricum	SP87 Lautropia mirabilis	SPPN13 Streptococcus multispecies_sppn13_18 nov_97.859%
SP212 Peptostreptococcus stomatis	SP88 Raoultella planticola	SPPN143 Streptococcus multispecies_sppn143_2 nov_97.849%
SP217 Granulicatella elegans	SP89 Lawsonella clevelandensis	SPPN151 Streptococcus multispecies_sppn151_17 nov_92.505%
SP219 Saccharibacteria (TM7) [G-3] bacterium_HMT_351	SP90 Bifidobacterium breve	SPPN171 Streptococcus multispecies_sppn171_17 nov_94.658%