

## Species

SP150 Aggregatibacter sp.\_HMT\_313  
SP151 Peptidiphaga gingivicola  
SP152 Streptococcus australis  
SP153 Leptotrichia trevisanii  
SP154 Leptotrichia shahii  
SP155 Peptoanaerobacter yunii  
SP156 Lactobacillus salivarius  
SP157 Treponema sp.\_HMT\_268  
SP158 Treponema lecitininolyticum  
SP159 Streptococcus sinensis  
SP16 Actinomyces viscosus  
SP160 Peptidiphaga sp.\_HMT\_183  
SP161 Porphyromonas gingivalis  
SP162 Actinomyces sp.\_HMT\_170  
SP163 Gemella moribillum  
SP164 Gemella haemolysans  
SP165 Actinomyces israelii  
SP166 Saccharibacteria\_(TM7)\_[G-2] bacterium\_HMT\_350  
SP167 Actinomyces sp.\_HMT\_896  
SP168 Lachnoanaerobaculum umeaense  
SP169 Actinomyces cardiffensis  
SP17 Streptococcus anginosus  
SP170 Streptococcus oralis\_subsp.\_tigurinus\_clade\_071  
SP171 Actinomyces odontolyticus  
SP172 Prevotella sp.\_HMT\_475  
SP173 Actinomyces sp.\_Oral\_Taxon\_848  
SP174 Streptococcus thermophilus  
SP175 Porphyromonas sp.\_HMT\_277  
SP176 Johnsonella ignava  
SP177 Selenomonas flueggei  
SP178 Peptostreptococcaceae\_[XII][G-9] [Eubacterium]\_brachy  
SP179 Ottowia sp.\_HMT\_894  
SP18 Fusobacterium periodonticum  
SP180 Selenomonas artemidis  
SP181 Centipeda periodontii  
SP182 Actinomyces timonensis  
SP183 Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_347  
SP184 Gracilibacteria\_(GN02)\_[G-2] bacterium\_HMT\_873  
SP185 Parvimonas sp.\_HMT\_110  
SP186 Veillonella atypica  
SP187 Saccharibacteria\_(TM7)\_[G-8] bacterium\_HMT\_955  
SP188 Kingella sp.\_HMT\_012  
SP189 Capnocytophaga sp.\_HMT\_326  
SP19 Streptococcus infantis  
SP190 Megaspheera micronuciformis  
SP191 Atopobium rimae  
SP193 Veillonellaceae\_[G-1] bacterium\_HMT\_135  
SP194 Alloprevotella sp.\_HMT\_912  
SP195 Capnocytophaga granulosa  
SP196 Streptococcus sobrinus  
SP197 Prevotella sp.\_HMT\_300  
SP198 Leptotrichia sp.\_HMT\_221  
SP199 Veillonellaceae\_[G-1] sp.\_Oral\_Taxon\_155  
SP2 Lautropia mirabilis  
SP20 Haemophilus parainfluenzae  
SP200 Lachnospiraceae\_[G-8] bacterium\_HMT\_500  
SP201 Treponema socranskii  
SP202 Filifactor alocis  
SP203 Stomatobaculum longum  
SP204 Prevotella baroniae  
SP205 Freitbacterium fastidiosum  
SP206 Campylobacter gracilis  
SP207 Porphyromonas sp.\_Oral\_Taxon\_B43  
SP208 Fusobacterium sp.\_HMT\_203  
SP209 Porphyromonas endodontalis  
SP21 Selenomonas sp.\_HMT\_137  
SP210 Gemella sanguinis  
SP211 Bergeyella sp.\_HMT\_900  
SP212 Bacteroidales\_[G-2] bacterium\_HMT\_274  
SP213 Eikenella corrodens  
SP214 Eubacterium sensillum

SP316 Leptotrichia sp.\_HMT\_2417  
SP319 Capnocytophaga sp.\_HMT\_323  
SP32 Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_957  
SP320 Fusobacterium sp.\_HMT\_204  
SP321 Prevotella pleuritidis  
SP324 Treponema sp.\_HMT\_951  
SP325 Treponema sp.\_HMT\_270  
SP326 Desulfobulbus sp.\_HMT\_041  
SP327 Lachnospiraceae\_[G-7] bacterium\_HMT\_086  
SP328 Veillonellaceae\_[G-1] bacterium\_HMT\_155  
SP329 Prevotella sp.\_HMT\_942  
SP33 Rothia aerea  
SP330 Citrobacter koseri  
SP331 Peptococcus sp.\_HMT\_168  
SP332 Bulleidia extructa  
SP333 Mogibacterium neglectum  
SP334 Veillonella sp.\_HMT\_780  
SP335 Selenomonas sp.\_HMT\_479  
SP336 Treponema sp.\_ovinefootrot  
SP337 Treponema amylovorum  
SP338 Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_352  
SP339 Prevotella sp.\_Oral\_Taxon\_B62  
SP34 Alloprevotella sp.\_HMT\_473  
SP340 Streptococcus parasanguinis\_I  
SP341 NA nodatum  
SP342 Selenomonas sp.\_Oral\_Taxon\_G67  
SP343 Selenomonas sp.\_HMT\_138  
SP344 Shuttleworthia satelles  
SP345 Treponema sp.\_HMT\_257  
SP346 Stomatobaculum sp.\_HMT\_373  
SP347 Aggregatibacter sp.\_HMT\_949  
SP348 Streptococcus cristatus\_clade\_578  
SP349 unclassified\_Peptostreptococcaceae oral\_sp.\_str\_CM2  
SP35 Treponema sp.\_HMT\_237  
SP350 Selenomonas sp.\_HMT\_937  
SP351 Pseudomonas fluorescens  
SP352 Tannerella forsythia  
SP353 Anaerolineae\_[G-1] bacterium\_HMT\_439  
SP354 Mycoplasma faucium  
SP355 Prevotella jejuni  
SP356 Treponema sp.\_Oral\_Taxon\_G85  
SP357 Leptotrichia sp.\_HMT\_463  
SP358 Neisseria sp.\_HMT\_523  
SP359 Freitbacterium sp.\_HMT\_359  
SP36 Capnocytophaga gingivalis  
SP360 Aggregatibacter sp.\_HMT\_512  
SP361 Selenomonas sp.\_HMT\_136  
SP362 Capnocytophaga sp.\_HMT\_335  
SP363 Leptotrichia goodfellowii  
SP364 Bifidobacterium dentium  
SP365 Selenomonas sp.\_HMT\_919  
SP366 Collibacter massiliensis  
SP367 Selenomonas sp.\_Oral\_Taxon\_E20  
SP368 unclassified\_Peptostreptococcaceae Oral\_sp.\_str\_CM5  
SP369 Selenomonas sp.\_Oral\_Taxon\_G55  
SP37 Parvimonas sp.\_Oral\_Taxon\_110  
SP370 Alloprevotella sp.\_HMT\_308  
SP371 Capnocytophaga sp.\_HMT\_878  
SP372 Veillonellaceae\_[G-1] bacterium\_HMT\_132  
SP373 Schaalia sp.\_HMT\_178  
SP374 Neisseria sp.\_HMT\_499  
SP375 Streptococcus sp.\_Oral\_Taxon\_B59  
SP376 Campylobacter sp.\_HMT\_044  
SP377 Phocaeicola abscessus  
SP378 Bacteroidetes\_[G-3] bacterium\_HMT\_503  
SP379 Haemophilus sp.\_HMT\_036  
SP38 Aggregatibacter segnis  
SP380 Saccharibacteria\_(TM7)\_[G-4] bacterium\_HMT\_355  
SP387 Schaalia cardiffensis  
SP388 Selenomonas sp.\_Oral\_Taxon\_F21  
SP389 Selenomonas sp.\_HMT\_446

SPN126 Selenomonas sp.\_Oral\_Taxon\_137\_nov\_96.324%  
SPN127 Selenomonas infelix\_nov\_97.059%  
SPN128 Selenomonas sp.\_HMT\_149\_nov\_96.947%  
SPN129 Corynebacterium sp.\_Oral\_Taxon\_A46\_nov\_95.898%  
SPN13 Selenomonas artemidis\_nov\_97.606%  
SPN130 Actinomyces viscosus\_nov\_96.234%  
SPN131 Corynebacterium durum\_nov\_96.032%  
SPN14 Actinomyces sp.\_Oral\_Taxon\_178\_nov\_97.514%  
SPN15 Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_957\_nov\_97.639%  
SPN16 Streptococcus gordonii\_nov\_97.782%  
SPN17 Treponema denticola\_nov\_97.170%  
SPN18 Corynebacterium sp.\_Oral\_Taxon\_A46\_nov\_97.065%  
SPN19 Selenomonas sp.\_Oral\_Taxon\_137\_nov\_97.059%  
SPN2 Actinomyces sp.\_HMT\_897\_nov\_97.461%  
SPN20 Streptococcus mitis\_nov\_97.538%  
SPN21 Selenomonas artemidis\_nov\_97.710%  
SPN22 Actinomyces viscosus\_nov\_97.393%  
SPN23 Bacteroidetes\_[G-5] bacterium\_HMT\_511\_nov\_96.393%  
SPN24 Prevotella enoea\_nov\_96.680%  
SPN25 Peptostreptococcaceae\_[XII][G-2] bacterium\_HMT\_091\_nov\_97.194%  
SPN26 Kingella oralis\_nov\_97.529%  
SPN27 Capnocytophaga sp.\_Oral\_Taxon\_A48\_nov\_94.556%  
SPN28 Peptococcus sp.\_Oral\_Taxon\_167\_nov\_97.798%  
SPN29 Streptococcus gordonii\_nov\_92.818%  
SPN3 Actinomyces sp.\_HMT\_448\_nov\_95.969%  
SPN30 Actinomyces massiliensis\_nov\_97.790%  
SPN31 Neisseria sp.\_Oral\_Taxon\_14\_nov\_97.154%  
SPN32 Capnocytophaga ochracea\_nov\_97.093%  
SPN33 Neisseria elongata\_nov\_97.533%  
SPN34 Veillonella sp.\_HMT\_780\_nov\_96.958%  
SPN35 Actinomyces massiliensis\_nov\_89.496%  
SPN36 Actinomyces israelii\_nov\_97.393%  
SPN37 Granulicatella adiacens\_nov\_97.575%  
SPN38 Anaeroglobus geminatus\_nov\_95.104%  
SPN39 Treponema sp.\_HMT\_257\_nov\_97.826%  
SPN4 Leptotrichia sp.\_HMT\_215\_nov\_97.083%  
SPN40 Renibacterium salmoninarum\_nov\_90.979%  
SPN41 Prevotella pleuritidis\_nov\_97.087%  
SPN42 Actinomyces oris\_nov\_92.435%  
SPN43 Actinomyces massiliensis\_nov\_93.566%  
SPN44 Veillonellaceae\_[G-1] sp.\_Oral\_Taxon\_155\_nov\_96.241%  
SPN45 Actinomyces sp.\_HMT\_175\_nov\_97.830%  
SPN46 Actinomyces sp.\_Oral\_Taxon\_171\_nov\_91.728%  
SPN47 Actinomyces israelii\_nov\_96.685%  
SPN48 Brachymonas chironomi\_nov\_91.939%  
SPN49 Pleomorphochaeta multiformis\_nov\_85.882%  
SPN5 Selenomonas flueggei\_nov\_96.750%  
SPN50 Aggregatibacter sp.\_HMT\_898\_nov\_88.521%  
SPN51 Treponema sp.\_HMT\_258\_nov\_97.843%  
SPN52 Streptococcus anginosus\_nov\_97.974%  
SPN53 Leptotrichia sp.\_HMT\_223\_nov\_93.347%  
SPN54 Neisseria sp.\_Oral\_Taxon\_14\_nov\_96.584%  
SPN55 Streptococcus oralis\_nov\_95.113%  
SPN56 Capnocytophaga gingivalis\_nov\_95.720%  
SPN57 Capnocytophaga gingivalis\_nov\_97.471%  
SPN58 Prevotella sp.\_HMT\_443\_nov\_97.047%  
SPN59 Vagococcus penaei\_nov\_79.167%  
SPN6 Porphyromonas catoniae\_nov\_97.328%  
SPN60 Heribiconiux ginsengi\_nov\_83.270%  
SPN61 Neisseria sp.\_Oral\_Taxon\_14\_nov\_96.970%  
SPN62 Actinomyces johnsonii\_nov\_95.194%  
SPN63 Megaspheera micronuciformis\_nov\_94.918%  
SPN64 Fusobacterium nucleatum\_ss\_animalis\_nov\_97.605%  
SPN65 Saccharibacillus kuereensis\_nov\_78.378%  
SPN66 Streptococcus intermedius\_nov\_92.421%  
SPN67 Veillonellaceae\_[G-1] bacterium\_HMT\_150\_nov\_97.456%  
SPN68 Actinomyces massiliensis\_nov\_90.842%  
SPN69 Firmicutes\_[G] sp.\_Oral\_Taxon\_C68\_nov\_86.481%  
SPN7 Actinomyces dentalis\_nov\_97.885%  
SPN70 Actinomyces sp.\_HMT\_414\_nov\_92.952%  
SPN71 Leptotrichia mirabilis\_nov\_94.050%