

Species

- SP127 Actinomyces sp.\_HMT\_175
- SP128 Prevotella sp.\_HMT\_317
- SP129 Scardovia wiggisiae
- SP13 Stomatobaculum sp.\_HMT\_097
- SP132 Fusobacterium periodonticum
- SP135 Saccharibacteria\_(TM7)\_[G-3] bacterium\_HMT\_351
- SP139 Prevotella buccae
- SP14 Veillonella parvula
- SP140 Megasphaera micronuciformis
- SP142 Capnocytophaga gingivalis
- SP143 Saccharibacteria\_(TM7)\_[G-6] bacterium\_HMT\_870
- SP144 Porphyromonas sp.\_HMT\_275
- SP147 Prevotella loeschii
- SP148 Haemophilus parainfluenzae
- SP15 Veillonella dispar
- SP150 Prevotella pleuritidis
- SP151 Prevotella baroniae
- SP152 Prevotella denticola
- SP154 Schaalia sp.\_HMT\_180
- SP159 Gemella sanguinis
- SP16 Actinomyces dentalis
- SP160 Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_346
- SP163 Corynebacterium matruchotii
- SP165 Leptotrichia sp.\_HMT\_392
- SP166 Prevotella maculosa
- SP168 Veillonella rogosae
- SP17 Streptococcus mitis
- SP170 Stomatobaculum sp.\_HMT\_910
- SP172 Bacteroides zoogloeiformans
- SP178 Haemophilus haemolyticus
- SP179 Saccharibacteria\_(TM7)\_[G-5] bacterium\_HMT\_356
- SP18 Dialister invisus
- SP181 Alloprevotella sp.\_HMT\_473
- SP182 Treponema maltophilum
- SP183 Capnocytophaga leadbetteri
- SP185 Neisseria oralis
- SP188 Lachnospiraceae\_[G-2] bacterium\_HMT\_096
- SP189 Rothia mucilaginosa
- SP19 Streptococcus cristatus
- SP190 Ruminococcaceae\_[G-1] bacterium\_HMT\_075

- SP237 Aggregatibacter segnis
- SP238 Peptostreptococcus stomatis
- SP240 Parvimonas micra
- SP242 Streptococcus oralis\_subsp\_dentisani\_clade\_058
- SP243 Prevotella oulorum
- SP246 Prevotella veroralis
- SP247 Bifidobacterium dentium
- SP248 Peptostreptococcaceae\_[G-1] [Eubacterium]\_sulci
- SP251 Bulleidia extructa
- SP253 Prevotella oralis
- SP258 Peptostreptococcaceae\_[G-5] [Eubacterium]\_saphenum
- SP259 Schaalia cardiffensis
- SP26 Gemella haemolysans
- SP260 Streptococcus oralis\_subsp\_dentisani\_clade\_398
- SP261 Mogibacterium diversum
- SP262 Prevotella histicola
- SP265 Prevotella vespertina
- SP266 Prevotella nigrescens
- SP268 Treponema lecithinolyticum
- SP27 Selenomonas noxia
- SP270 Streptococcus sanguinis
- SP271 Cryptobacterium curtum
- SP275 Fretibacterium sp.\_HMT\_362
- SP277 Bacteroidetes\_[G-3] bacterium\_HMT\_365
- SP279 Pseudoramibacter alactolyticus
- SP28 Actinomyces sp.\_HMT\_414
- SP280 Aggregatibacter aphrophilus
- SP281 Anaeroglobus geminatus
- SP282 Prevotella intermedia
- SP283 Streptococcus oralis
- SP284 Streptococcus oralis\_subsp\_tigurinus\_clade\_070
- SP29 Filifactor alocis
- SP290 Solobacterium moorei
- SP295 Desulfobulbus sp.\_HMT\_041
- SP296 Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_352
- SP298 Streptococcus sp.\_HMT\_066
- SP299 Actinomyces johnsonii
- SP3 Streptococcus australis
- SP300 Schaalia sp.\_HMT\_172
- SP303 Peptostreptococcaceae\_[G-4] bacterium\_HMT\_369

- SP67 Streptococcus infantis\_clade\_431
- SP69 Campylobacter showae
- SP7 Leptotrichia wadei
- SP70 Granulicatella elegans
- SP71 Corynebacterium durum
- SP72 Selenomonas sputigena
- SP73 Streptococcus mutans
- SP74 Rothia dentocariosa
- SP75 Porphyromonas pasteri
- SP77 Finegoldia magna
- SP78 Alloprevotella rava
- SP8 Catonella morbi
- SP80 Fusobacterium nucleatum
- SP82 Streptococcus constellatus
- SP83 Leptotrichia sp.\_HMT\_215
- SP84 Schaalia lingnae
- SP85 Actinomyces timonensis
- SP86 Lachnoanaerobaculum saburreum
- SP87 Streptococcus cristatus\_clade\_578
- SP88 Prevotella jejuni
- SP89 Ruminococcaceae\_[G-2] bacterium\_HMT\_085
- SP9 Gemella morbillorum
- SP90 Prevotella melaninogenica
- SP93 Actinomyces sp.\_HMT\_169
- SP94 Lachnoanaerobaculum umeaense
- SP95 Lachnoanaerobaculum gingivalis
- SP96 Treponema denticola
- SP97 Streptococcus downii
- SP98 Streptococcus sp.\_HMT\_074
- SP99 Porphyromonas endodontalis
- SPN102 Porphyromonas sp.\_HMT\_284\_nov\_97.131%
- SPN103 Selenomonas diana nov\_97.830%
- SPN11 Actinomyces sp.\_HMT\_175\_nov\_97.746%
- SPN112 Lacrimispora xylanolytica\_nov\_88.613%
- SPN119 Porphyromonas uenonis\_nov\_94.456%
- SPN172 Sporanaerobacter acetigenes\_nov\_84.016%
- SPN22 Campylobacter rectus\_nov\_97.826%
- SPN38 Prevotella jejuni\_nov\_97.755%
- SPN58 Bifidobacterium dentium\_nov\_97.240%
- SPN69 Prevotella sp.\_HMT\_305\_nov\_93.865%