

## Species

- SP1 Streptococcus salivarius
- SP10 Leptotrichia wadei
- SP100 Prevotella buccae
- SP101 Leptotrichia hongkongensis
- SP103 Aggregatibacter sp.\_HMT\_458
- SP104 Capnocytophaga leadbetteri
- SP105 Eikenella corrodens
- SP107 Lachnoanaerobaculum gingivalis
- SP108 Prevotella sp.\_HMT\_317
- SP109 Anaeroglobus geminatus
- SP11 Peptidiphaga sp.\_HMT\_183
- SP110 Selenomonas sp.\_HMT\_137
- SP111 Prevotella sp.\_HMT\_314
- SP112 Tannerella sp.\_HMT\_286
- SP113 Leptotrichia sp.\_HMT\_392
- SP114 Alloprevotella sp.\_HMT\_473
- SP115 Actinomyces naeslundii
- SP116 Lancefieldella rimae
- SP117 Cardiobacterium hominis
- SP119 Bergeyella sp.\_HMT\_900
- SP12 Sphingomonas oligophenolica
- SP120 Arachnia propionica
- SP122 Lachnospiraceae\_[G-3] bacterium\_HMT\_100
- SP123 Tannerella serpentiniformis
- SP124 Streptococcus chosunense
- SP125 Absconditabacteria\_(SR1)\_[G-1] bacterium\_HMT\_345
- SP127 Prevotella oulorum
- SP128 Capnocytophaga sp.\_HMT\_380
- SP129 Streptococcus oralis\_subsp.\_dentisani\_clade\_058
- SP13 Streptococcus intermedius
- SP130 Lautropia mirabilis
- SP131 Treponema maltophilum
- SP132 Arachnia rubra
- SP133 Selenomonas flueggei
- SP134 Fusobacterium nucleatum\_subsp.\_vincentii
- SP136 Streptococcus sp.\_HMT\_056
- SP137 Capnocytophaga sp.\_HMT\_864
- SP138 Catonella morbi
- SP139 Rothia aeria
- SP14 Gemella morbillorum
- SP140 Lachnospiraceae\_[G-8] bacterium\_HMT\_500
- SP141 Campylobacter gracilis
- SP143 Rothia dentocariosa
- SP144 Campylobacter sp.\_HMT\_044
- SP146 Scardovia wiggisiae
- SP147 Prevotella marshii
- SP15 Bergeyella sp.\_HMT\_322
- SP150 Saccharibacteria\_(TM7)\_[G-2] bacterium\_HMT\_350
- SP153 Porphyromonas gingivalis
- SP154 Prevotella loeschei
- SP155 Fretibacterium sp.\_HMT\_360
- SP157 Prevotella veroralis
- SP159 Prevotella sp.\_HMT\_292
- SP16 Veillonella parvula
- SP160 Streptococcus cristatus\_clade\_578
- SP161 Actinomyces sp.\_HMT\_175
- SP163 Veillonella rogosae
- SP164 Schaalia sp.\_HMT\_180
- SP165 Selenomonas sp.\_HMT\_138
- SP168 Lachnoanaerobaculum saburreum
- SP17 Dialister invisus
- SP171 Streptococcus oralis\_subsp.\_dentisani\_clade\_398
- SP172 Capnocytophaga sp.\_HMT\_336
- SP180 Prevotella salivae
- SP182 Peptostreptococcaceae\_[XII][G-4] bacterium\_HMT\_103
- SP184 Shuttleworthia satelles
- SP19 Peptostreptococcus stomatis
- SP192 Schaalia sp.\_HMT\_172
- SP194 Leptotrichia sp.\_HMT\_498
- SP2 Solobacterium moorei
- SP20 Prevotella intermedia
- SP21 Veillonella atypica
- SP22 Leptotrichia sp.\_HMT\_223
- SP23 Porphyromonas sp.\_HMT\_275
- SP24 Prevotella saccharolytica
- SP25 Veillonella dispar
- SP26 Stomatobaculum longum
- SP27 Neisseria elongata
- SP28 Fusobacterium canifelinum
- SP29 Neisseria sicca
- SP3 Streptococcus sp.\_HMT\_064
- SP30 Streptococcus mitis
- SP31 Neisseria flavescens
- SP32 Selenomonas noxia
- SP33 Porphyromonas endodontalis
- SP34 Porphyromonas catoniae
- SP35 Ottowia sp.\_HMT\_894
- SP36 Neisseria subflava
- SP37 Prevotella oris
- SP38 Leptotrichia sp.\_HMT\_215
- SP39 Selenomonas sputigena
- SP4 Bacteroidales\_[G-2] bacterium\_HMT\_274
- SP40 Prevotella sp.\_HMT\_300
- SP41 Streptococcus sanguinis
- SP42 Fusobacterium periodonticum
- SP43 Streptococcus vestibularis
- SP44 Capnocytophaga endodontalis
- SP45 Capnocytophaga ochracea
- SP46 Stenotrophomonas maltophilia
- SP47 Streptococcus oralis\_subsp.\_tigurinus\_clade\_070
- SP48 Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_348
- SP49 Leptotrichia sp.\_HMT\_225
- SP50 Absconditabacteria\_(SR1)\_[G-1] bacterium\_HMT\_874
- SP51 Neisseria sp.\_HMT\_020
- SP52 Aggregatibacter segnis
- SP53 Prevotella sp.\_HMT\_820
- SP54 Campylobacter rectus
- SP55 Porphyromonas pasteri
- SP56 Saccharibacteria\_(TM7)\_[G-5] bacterium\_HMT\_356
- SP57 Selenomonas infelix
- SP58 Gracilibacteria\_(GN02)\_[G-2] bacterium\_HMT\_873
- SP59 Saccharibacteria\_(TM7)\_[G-3] bacterium\_HMT\_351
- SP6 Streptococcus gordonii
- SP60 Peptostreptococcaceae\_[XIII][G-4] bacterium\_HMT\_369
- SP61 Kingella oralis
- SP62 Campylobacter concisus
- SP63 Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_869
- SP64 Leptotrichia buccalis
- SP65 Aggregatibacter sp.\_HMT\_512
- SP66 Prevotella melaninogenica
- SP67 Streptococcus cristatus
- SP68 Capnocytophaga gingivalis
- SP69 Haemophilus parainfluenzae
- SP7 Leptotrichia hofstadii
- SP70 Granulicatella adiacens
- SP71 Corynebacterium matruchotii
- SP72 Campylobacter showae
- SP73 Filifactor alocis
- SP75 Capnocytophaga granulosa
- SP76 Leptotrichia sp.\_HMT\_212
- SP77 Actinomyces sp.\_HMT\_171
- SP78 Klebsiella michiganensis
- SP79 Peptoanaerobacter [Eubacterium] yurii
- SP8 Fusobacterium nucleatum
- SP80 Treponema socranskii
- SP81 Parvimonas micra
- SP82 Prevotella maculosa
- SP83 Alloprevotella tannerae
- SP84 Streptococcus anginosus
- SP85 Prevotella denticola
- SP86 Phocaeicola abscessus
- SP87 Prevotella nanceiensis
- SP88 Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_957
- SP89 Cardiobacterium valvarum
- SP9 Streptococcus oralis
- SP91 Actinomyces sp.\_HMT\_169
- SP92 Saccharibacteria\_(TM7)\_[G-4] bacterium\_HMT\_355
- SP93 Tannerella forsythia
- SP94 Peptostreptococcaceae\_[XII][G-1] [Eubacterium]\_infirmum
- SP95 Ruminococcaceae\_[G-2] bacterium\_HMT\_085
- SP96 Capnocytophaga sputigena
- SP97 Bacteroidetes\_[G-3] bacterium\_HMT\_280
- SP98 Lachnoanaerobaculum orale
- SP99 Saccharibacteria\_(TM7)\_[G-1] bacterium\_HMT\_346
- SPN1 Porphyromonas sp.\_HMT\_284\_nov\_97.436%
- SPN11 Bergeyella zoohelcum\_nov\_92.673%
- SPN15 Bacteroides zoogloeiformans\_nov\_96.578%
- SPN2 Aggregatibacter sp.\_HMT\_458\_nov\_97.826%
- SPN22 Actinomyces sp.\_HMT\_448\_nov\_96.718%
- SPN27 Prevotella sp.\_HMT\_305\_nov\_93.688%
- SPN3 Sphingomonas jeddahensis\_nov\_97.447%
- SPN33 Selenomonas sp.\_HMT\_137\_nov\_96.947%
- SPN39 Saccharibacteria\_(TM7)\_[G-3] bacterium\_HMT\_351\_nov\_92.871%
- SPN4 Neisseria elongata\_nov\_96.374%
- SPN44 Selenomonas sp.\_HMT\_137\_nov\_97.126%
- SPN5 Neisseria sp.\_HMT\_020\_nov\_97.619%
- SPN51 Leptotrichia sp.\_HMT\_215\_nov\_97.292%
- SPN55 Actinomyces naeslundii\_nov\_97.593%
- SPN6 Prevotella melaninogenica\_nov\_97.170%
- SPN63 Streptococcus sanguinis\_nov\_97.538%
- SPN66 Selenomonas noxia\_nov\_96.337%
- SPN71 Selenomonas sp.\_HMT\_137\_nov\_97.901%
- SPN72 Corynebacterium durum\_nov\_97.384%
- SPN73 Abiotrophia defectiva\_nov\_96.481%
- SPN74 Selenomonas sputigena\_nov\_97.026%
- SPN75 Selenomonas sp.\_HMT\_479\_nov\_96.756%
- SPP2 Lachnoanaerobaculum gingivalis\_umeaense
- SPP3 Selenomonas flueggei\_noxia
- SPP6 Veillonella dispar\_parvula