ecies

- SP139 Lautropia mirabilis
- SP14 Fusobacterium sp._HMT_203
- SP140 Porphyromonas pasteri
- SP141 Kingella oralis
- SP142 Tannerella sp._HMT_286
- SP143 Capnocytophaga haemolytica
- SP145 Veillonella parvula
- SP146 Veillonella dispar
- SP147 Prevotella loescheii
- SP148 Schaalia odontolytica
- SP149 Treponema denticola
- SP15 Actinomyces timonensis
- SP150 Pseudoleptotrichia sp._HMT_221
- SP151 Saccharibacteria_(TM7)_[G−5] bacterium_HMT_356
- SP152 Pseudoleptotrichia sp._HMT_219
- SP153 Selenomonas sp._HMT_920
- SP154 Butyrivibrio sp._HMT_455
- SP156 Solobacterium moorei
- SP157 Streptococcus sp._HMT_066
- SP158 Phocaeicola abscessus
- SP159 Aggregatibacter actinomycetemcomitans
- SP16 Actinomyces sp._HMT_170
- SP160 Capnocytophaga sp._HMT_864
- SP161 Prevotella nigrescens
- SP162 Peptostreptococcaceae_[G−9] [Eubacterium]_brachy
- SP163 Parvimonas sp._HMT_110
- SP164 Selenomonas sputigena
- SP165 Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
- SP166 Rothia mucilaginosa
- SP168 Actinomyces johnsonii
- SP169 Leptotrichia sp._HMT_212
- SP17 Actinomyces graevenitzii
- SP170 Leptotrichia shahii
- SP172 Treponema maltophilum
- SP173 Schaalia georgiae
- SP174 Pseudoleptotrichia goodfellowii
- SP175 Kingella sp._HMT_932
- SP176 Actinomyces naeslundii
- SP177 Prevotella dentalis
- SP272 Haemophilus sp._HMT_036
- SP273 Veillonellaceae_[G−1] bacterium_HMT_132
- SP276 Porphyromonas sp._HMT_284
- SP277 Selenomonas dianae
- SP278 Streptococcus oralis_subsp._tigurinus_clade_071
- SP279 Leptotrichia sp._HMT_225
- SP28 Actinomyces sp._HMT_169
- SP280 Aggregatibacter aphrophilus
- SP281 Leptotrichia wadei
- SP284 Fusobacterium canifelinum
- SP285 Streptococcus australis
- SP287 Prevotella micans
- SP289 Neisseria sp._HMT_018
- SP29 Lachnoanaerobaculum saburreum
- SP290 Streptococcus anginosus
- SP292 Capnocytophaga sp._HMT_338
- SP293 Streptococcus cristatus
- SP297 Neisseria oralis
- SP299 Capnocytophaga ochracea
- SP30 Granulicatella adiacens
- SP301 Parvimonas micra
- SP302 Streptococcus intermedius
- SP308 Scardovia wiggsiae
- SP309 Fusobacterium hwasookii
- SP310 Absconditabacteria_(SR1)_[G−1] bacterium_HMT_874
- SP314 Prevotella koreensis
- SP315 Leptotrichia hongkongensis
- SP317 Saccharibacteria_(TM7)_[G−1] bacterium_HMT_869
- SP318 Schaalia sp._HMT_172
- SP319 Peptidiphaga sp._HMT_183
- SP320 Treponema sp._HMT_258
- SP321 Haemophilus sputorum
- SP322 Porphyromonas sp._HMT_930
- SP325 Olsenella profusa
- SP328 Cardiobacterium hominis
- SP33 Haemophilus pittmaniae
- SP330 Propionibacterium acidifaciens
- SP339 Selenomonas sp._HMT_479
- SP34 Fretibacterium sp._HMT_360
- SP69 Prevotella oulorum
- SP7 Prevotella sp._HMT_472
- SP70 Lachnospiraceae_[G−2] bacterium_HMT_096
- SP71 Prevotella pallens
- SP72 Filifactor alocis
- SP73 Alloprevotella tannerae
- SP74 Treponema socranskii
- SP75 Streptococcus cristatus_clade_578
- SP76 Corynebacterium matruchotii
- SP77 Prevotella fusca
- SP79 Stomatobaculum sp._HMT_097
- SP8 Fusobacterium nucleatum
- SP80 Dialister invisus
- SP81 Prevotella jejuni
- SP82 Haemophilus paraphrohaemolyticus
- SP83 Haemophilus parainfluenzae
- SP84 Streptococcus parasanguinis_clade_411
- SP85 Campylobacter concisus
- SP86 Prevotella melaninogenica
- SP87 Bacteroidales_[G−2] bacterium_HMT_274
- SP88 Streptococcus koreensis
- SP89 Streptococcus sp._HMT_057
- SP9 Peptococcus sp._HMT_167
- SP90 Neisseria elongata
- SP92 Neisseria subflava
- SP93 Prevotella sp._HMT_315
- SP95 Johnsonella sp._HMT_166
- SP97 Saccharibacteria_(TM7)_[G−6] bacterium_HMT_870
- SP98 Streptococcus periodonticum
- SP99 Streptococcus infantis_clade_431
- SPN104 Corynebacterium matruchotii_nov_97.951%
- SPN106 Treponema sp._HMT_927_nov_91.411%
- SPN115 Peptostreptococcaceae_[G−5] bacterium_HMT_493_nov_96.4
- SPN127 Prevotella sp._HMT_305_nov_93.865%
- SPN139 Leptotrichia sp._HMT_225_nov_96.304%
- SPN148 Kingella oralis_nov_97.536%
- SPN15 Peptidiphaga gingivicola_nov_94.057%
- SPN156 Actinomyces sp._HMT_175_nov_97.746%
- SPN166 Lacrimispora xylanolytica_nov_88.866%