

Species

- SP1 Streptococcus chosunense
- SP10 Haemophilus parainfluenzae
- SP100 Neisseria cinerea
- SP101 Ruminococcaceae_[G-2] bacterium_HMT_085
- SP102 Prevotella shahii
- SP103 Prevotella sp._HMT_309
- SP104 Campylobacter concisus
- SP105 Streptococcus gordonii
- SP106 Leptotrichia sp._HMT_212
- SP107 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_875
- SP108 Actinomyces johnsonii
- SP109 Streptococcus cristatus_clade_578
- SP11 Streptococcus parasanguinis
- SP110 Capnocytophaga gingivalis
- SP111 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348
- SP112 Megaspheara micronuciformis
- SP113 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957
- SP114 Lachnoanaerobaculum umeaense
- SP115 Stomatobaculum longum
- SP116 Eikenella corrodens
- SP118 Gemella morbillorum
- SP119 Kingella oralis
- SP12 Streptococcus sp._HMT_061
- SP13 Lachnoanaerobaculum gingivalis
- SP14 Streptococcus salivarius
- SP15 Neisseria elongata
- SP16 Streptococcus oralis
- SP17 Streptococcus mitis
- SP18 Fusobacterium periodonticum
- SP2 Granulicatella adiacens
- SP21 Streptococcus peroris
- SP22 Streptococcus sanguinis
- SP23 Streptococcus australis
- SP24 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_352
- SP25 Prevotella pallens
- SP26 Streptococcus oralis_subsp._dentisani_clade_058
- SP27 Streptococcus cristatus
- SP28 Oribacterium sinus
- SP29 Peptostreptococcus stomatis
- SP3 Schaalia odontolytica
- SP30 Streptococcus infantis_clade_431
- SP31 Neisseria mucosa
- SP32 Neisseria perflava
- SP33 Gemella sanguinis
- SP34 Parvimonas sp._HMT_110
- SP35 Capnocytophaga sputigena
- SP36 Prevotella histicola
- SP37 Streptococcus sp._HMT_066
- SP38 Mogibacterium vescum
- SP39 Streptococcus sp._HMT_074
- SP4 Streptococcus rubneri
- SP40 Streptococcus parasanguinis_clade_411
- SP41 Solobacterium moorei
- SP42 Bergeyella sp._HMT_206
- SP43 Neisseria sicca
- SP44 Streptococcus periodonticum
- SP45 Veillonella parvula
- SP46 Schaalia lingnae_[Not_Validly_Published]
- SP47 Schaalia odontolyticus
- SP48 Granulicatella elegans
- SP49 Schaalia sp._HMT_172
- SP5 Porphyromonas pasteri
- SP50 Stomatobaculum sp._HMT_097
- SP51 Gemella haemolysans
- SP52 Schaalia sp._HMT_180
- SP53 Leptotrichia sp._HMT_417
- SP54 Leptotrichia sp._HMT_215
- SP55 Streptococcus sp._HMT_057
- SP56 Lachnospiraceae_[G-2] bacterium_HMT_096
- SP57 Prevotella vespertina
- SP58 Lancefieldella parvula
- SP59 Rothia mucilaginosa
- SP6 Leptotrichia wadei
- SP60 Capnocytophaga leadbetteri
- SP61 Veillonella rogosae
- SP62 Lautropia mirabilis
- SP63 Streptococcus oralis_subsp._tigurinus_clade_070
- SP64 Peptostreptococcaceae_[X1][G-1] [Eubacterium]_sulci
- SP65 Abiotrophia defectiva
- SP66 Rothia dentocariosa
- SP67 Rothia aerea
- SP68 Corynebacterium argentoratense
- SP69 Alloprevotella sp._HMT_308
- SP7 Neisseria flavescens
- SP70 Streptococcus downii
- SP71 Bergeyella sp._HMT_322
- SP72 Klebsiella michiganensis
- SP73 Aggregatibacter sp._HMT_513
- SP74 Arachnia propionica
- SP76 Actinomyces graevenitzi
- SP79 Veillonella atypica
- SP8 Prevotella nanceiensis
- SP80 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870
- SP81 Schaalia meyeri
- SP82 Streptococcus intermedius
- SP84 Streptococcus thermophilus
- SP85 Enterobacter cloacae
- SP87 Fusobacterium nucleatum
- SP88 Prevotella jejuni
- SP89 Prevotella salivae
- SP9 Prevotella melaninogenica
- SP90 Veillonella tobetsuensis
- SP92 Actinomyces naeslundii
- SP93 Lachnoanaerobaculum orale
- SP94 Neisseria subflava
- SP95 Ruminococcaceae_[G-1] bacterium_HMT_075
- SP96 Haemophilus sputorum
- SP97 Streptococcus mutans
- SP98 Parvimonas micra
- SPN1 Streptococcus parasanguinis_nov_96.970%
- SPN10 Streptococcus salivarius_nov_97.561%
- SPN12 Granulicatella adiacens_nov_95.327%
- SPN13 Parvimonas micra_nov_95.914%
- SPN14 Gemella haemolysans_nov_94.981%
- SPN2 Gemella sanguinis_nov_94.595%
- SPN3 Prevotella sp._HMT_305_nov_93.898%
- SPN4 Streptococcus oralis_nov_96.992%
- SPN5 Streptococcus sanguinis_nov_97.727%
- SPN6 Streptococcus mitis_nov_85.393%
- SPN7 Streptobacillus notomitis_nov_93.725%
- SPN8 Schaalia odontolytica_nov_97.104%
- SPN9 Alloprevotella sp._HMT_308_nov_97.436%
- SPP2 Streptococcus sp._HMT_061_sp._HMT_066
- SPP3 Streptococcus infantis_infantis_clade_638
- SPP4 Streptococcus infantis_infantis_clade_431_infantis_clade_638
- SPP7 Veillonella dispar_parvula
- SPP8 Oribacterium asaccharolyticum_parvum
- SPP9 Enterobacter_Leclercia adecarboxylata_cloacae