

Species

- SP2 Haemophilus parainfluenzae
- SP28 Rothia mucilaginosa
- SP29 Prevotella melaninogenica
- SP3 Veillonella atypica
- SP31 Prevotella hispicola
- SP32 Streptococcus peroris
- SP36 Leptotrichia sp._HMT_215
- SP39 Prevotella vespertina
- SP4 Streptococcus gordonii
- SP40 Schaalia lingnae_[Not_Validly_Published]
- SP41 Veillonella dispar
- SP45 Granulicatella adiacens
- SP48 Megasphaera micronuciformis
- SP49 Prevotella sp._HMT_313
- SP51 Streptococcus sp._HMT_074
- SP52 Abiotrophia defectiva
- SP59 Gemella sanguinis
- SP61 Selenomonas sp._HMT_136
- SP63 Actinomyces graevenitzi
- SP65 Gemella haemolysans
- SP67 Actinomyces oris
- SP68 Bifidobacterium longum
- SP69 Schaalia sp._HMT_180
- SP7 Capnocytophaga sputigena
- SP71 Capnocytophaga leadbetteri
- SP73 Alloprevotella sp._HMT_308
- SP74 Fusobacterium periodonticum
- SP76 Prevotella salivae
- SP77 Oribacterium sinus
- SP78 Campylobacter concisus
- SP8 Streptococcus parasanguinis_clade_411
- SP81 Schaalia sp._HMT_172
- SP83 Prevotella nanceiensis
- SP84 Schaalia odontolytica
- SP85 Stomatobaculum longum
- SPN1 Rothia mucilaginosa_nov_90.443%
- SPN10 Rothia mucilaginosa_nov_93.627%
- SPN11 Abiotrophia defectiva_nov_97.669%
- SPN12 Megasphaera micronuciformis_nov_89.744%
- SPN13 Megasphaera micronuciformis_nov_96.262%
- SPN14 Schaalia lingnae_[Not_Validly_Published]_nov_94.299%
- SPN15 Anaerobacillus arseniciselenatis_nov_88.605%
- SPN16 Actinomyces graevenitzi_nov_93.381%
- SPN17 Alloprevotella sp._HMT_308_nov_87.470%
- SPN18 Bifidobacterium longum_nov_97.789%
- SPN19 Prevotella salivae_nov_96.919%
- SPN2 Prevotella sp._HMT_942_nov_83.059%
- SPN20 Prevotella melaninogenica_nov_97.630%
- SPN21 Rothia mucilaginosa_nov_92.402%
- SPN22 Haemophilus parainfluenzae_nov_97.424%
- SPN24 Veillonella dispar_nov_90.164%
- SPN3 Veillonella atypica_nov_92.774%
- SPN33 Veillonella dispar_nov_92.541%
- SPN4 Streptococcus gordonii_nov_91.183%
- SPN44 Abiotrophia defectiva_nov_88.345%
- SPN5 Bifidobacterium cebidarum_nov_95.332%
- SPN6 Anaerobacillus arseniciselenatis_nov_88.372%
- SPN61 Schaalia sp._HMT_180_nov_95.735%
- SPN7 Blautia faecicola_nov_90.000%
- SPN74 Oxalophagus oxalicus_nov_88.152%
- SPN8 Rothia mucilaginosa_nov_95.599%
- SPN81 Actinomyces oris_nov_96.927%
- SPN9 Prevotella nanceiensis_nov_94.340%
- SPP12 Streptococcus australis_rubneri_sp._HMT_066
- SPP14 Streptococcus salivarius_vestibularis
- SPP15 Schaalia odontolyticus_sp._HMT_180
- SPP16 Atopobium_Lancefieldella parvula_parvulum
- SPP2 Streptococcus parasanguinis_clade_411_sinensis_sp._HMT_056
- SPP3 Streptococcus cristatus_downii_gwangjuense_infantis_infantis_cla...(17 species)
- SPP4 Streptococcus cristatus_cristatus_clade_578_downii_gwangjuense_i...(18 species)
- SPP5 Streptococcus cristatus_downii_gordonii_gwangjuense_infantis_inf...(18 species)
- SPP6 Streptococcus cristatus_cristatus_clade_578
- SPP7 Streptococcus lactarius_peroris
- SPP9 Streptococcus parasanguinis_parasanguinis_clade_721_sp._HMT_066
- SPPN1 Schaalia multispecies_sppn1_3_nov_93.365%
- SPPN2 Arthrobacter multispecies_sppn2_2_nov_88.605%