

SP149 Peptostreptococcaceae_[XI][G-7] [Eubacterium]_yurii_subsp._yurii_&_margaretiae	SP311 Ruminococcaceae_[G-2] bacterium_HMT_085	SP83 Veillonella atypica
SP151 Prevotella sp._HMT_317	SP312 Treponema sp._HMT_234	SP84 Fusobacterium hwasookii
SP150 Streptococcus cristatus	SP315 Selenomonas sp._HMT_919	SP85 Alloprevotella sp._HMT_473
SP151 Treponema sp._HMT_237	SP316 Solobacterium moorei	SP87 Streptococcus gordonii
SP152 Capnocytophaga sp._HMT_864	SP317 Prevotella sp._HMT_301	SP88 Streptococcus sanguinis
SP153 Lancefieldella parvula	SP318 Prevotella buccae	SP9 Leptotrichia sp._HMT_212
SP154 Actinomyces dentalis	SP32 Abiotrophia defectiva	SP90 Fusobacterium periodonticum
SP155 Schaalia sp._HMT_180	SP320 Fusobacterium sp._HMT_204	SP91 Leptotrichia sp._HMT_215
SP157 Bacteroidetes_[G-3] bacterium_HMT_365	SP323 Treponema sp._HMT_253	SP92 Treponema sp._HMT_231
SP159 Actinomyces israelii	SP324 Schaalia odontolytica	SP93 Haemophilus haemolyticus
SP16 Selenomonas sputigena	SP325 Kingella denitrificans	SP94 Aggregatibacter sp._HMT_949
SP160 Campylobacter rectus	SP326 Actinomyces massiliensis	SP95 Campylobacter concisus
SP161 Peptostreptococcaceae_[XI][G-6] [Eubacterium]_nodatum	SP328 Capnocytophaga sp._HMT_338	SP97 Lachnoanaerobaculum sp._HMT_083
SP162 Ruminococcaceae_[G-1] bacterium_HMT_075	SP33 Selenomonas noxia	SP98 Selenomonas diana
SP165 Corynebacterium durum	SP330 Prevotella sp._HMT_315	SP99 Porphyromonas sp._HMT_275
SP166 Veillonellaceae_[G-1] bacterium_HMT_129	SP331 Prevotella jejuni	SPN103 Actinomyces israelii_nov_96.838%
SP167 Lautropia mirabilis	SP333 Treponema sp._HMT_238	SPN108 Cardiobacterium hominis_nov_97.510%
SP168 Streptococcus australis	SP336 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957	SPN111 Prevotella veroralis_nov_97.959%
SP169 Veillonellaceae_[G-1] bacterium_HMT_145	SP338 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348	SPN121 Porphyromonas catoniae_nov_97.951%
SP17 Selenomonas sp._HMT_937	SP339 Fretibacterium sp._HMT_361	SPN129 Actinomyces israelii_nov_94.882%
SP170 Lachnoanaerobaculum saburreum	SP340 Bifidobacterium dentium	SPN137 Corynebacterium matruchotii_nov_97.769%
SP171 Selenomonas flueggei	SP341 Stomatobaculum sp._HMT_910	SPN145 Selenomonas sp._HMT_481_nov_97.822%
SP172 Actinomyces sp._HMT_896	SP342 Capnocytophaga sp._HMT_324	SPN153 Hoppeia youngheungensis_nov_80.457%
SP173 Veillonellaceae_[G-1] bacterium_HMT_132	SP344 Olsenella uli	SPN162 Actinomyces sp._HMT_171_nov_96.813%
SP174 Capnocytophaga granulosa	SP347 Actinomyces oricola	SPN163 Prevotella sp._HMT_300_nov_97.955%
SP175 Leptotrichia sp._HMT_221	SP35 Capnocytophaga sp._HMT_336	SPN173 Actinomyces sp._HMT_897_nov_97.769%
SP176 Neisseria flavescens	SP350 Tannerella forsythia	SPN178 Desulfovibrio legallii_nov_94.343%
SP177 Gemella haemolysans	SP351 Streptococcus intermedius	SPN184 Tannerella forsythia_nov_97.308%
SP178 Campylobacter showae	SP353 Treponema lecithinolyticum	SPN193 Sneathia sanguinegens_nov_90.578%
SP179 Aggregatibacter sp._HMT_513	SP355 Prevotella marshii	SPN200 Lacrimispora xylanolytica_nov_88.613%
SP18 Streptococcus parasanguinis_clade_411	SP356 Gracilibacteria_(GN02)_[G-1] bacterium_HMT_872	SPN207 Corynebacterium durum_nov_97.484%
SP180 Treponema sp._HMT_257	SP358 Streptococcus sinensis	SPN210 Actinomyces sp._HMT_175_nov_97.751%
SP181 Bacteroidales_[G-2] bacterium_HMT_274	SP36 Porphyromonas pasteri	SPN214 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348_nov_95.0
SP182 Selenomonas sp._HMT_479	SP361 Neisseria bacilliformis	SPN216 Corynebacterium matruchotii_nov_97.955%
SP183 Granulicatella elegans	SP362 Saccharibacteria_(TM7)_[G-8] bacterium_HMT_955	SPN224 Kingella oralis_nov_97.536%
SP184 Prevotella maculosa	SP364 Peptostreptococcaceae_[XI][G-3] bacterium_HMT_495	SPN233 Tepidisphaera mucosa_nov_92.461%
SP186 Capnocytophaga sp._HMT_412	SP365 Peptoniphilaceae_[G-1] bacterium_HMT_113	SPN234 Selenomonas infelix_nov_97.822%
SP187 Capnocytophaga sp._HMT_335	SP366 Pseudoleptotrichia goodfellowii	SPN242 Actinomyces naeslundii_nov_97.963%
SP188 Streptococcus chosunense	SP367 Lachnospiraceae_[G-7] bacterium_HMT_086	SPN251 Prevotella oris_nov_97.131%