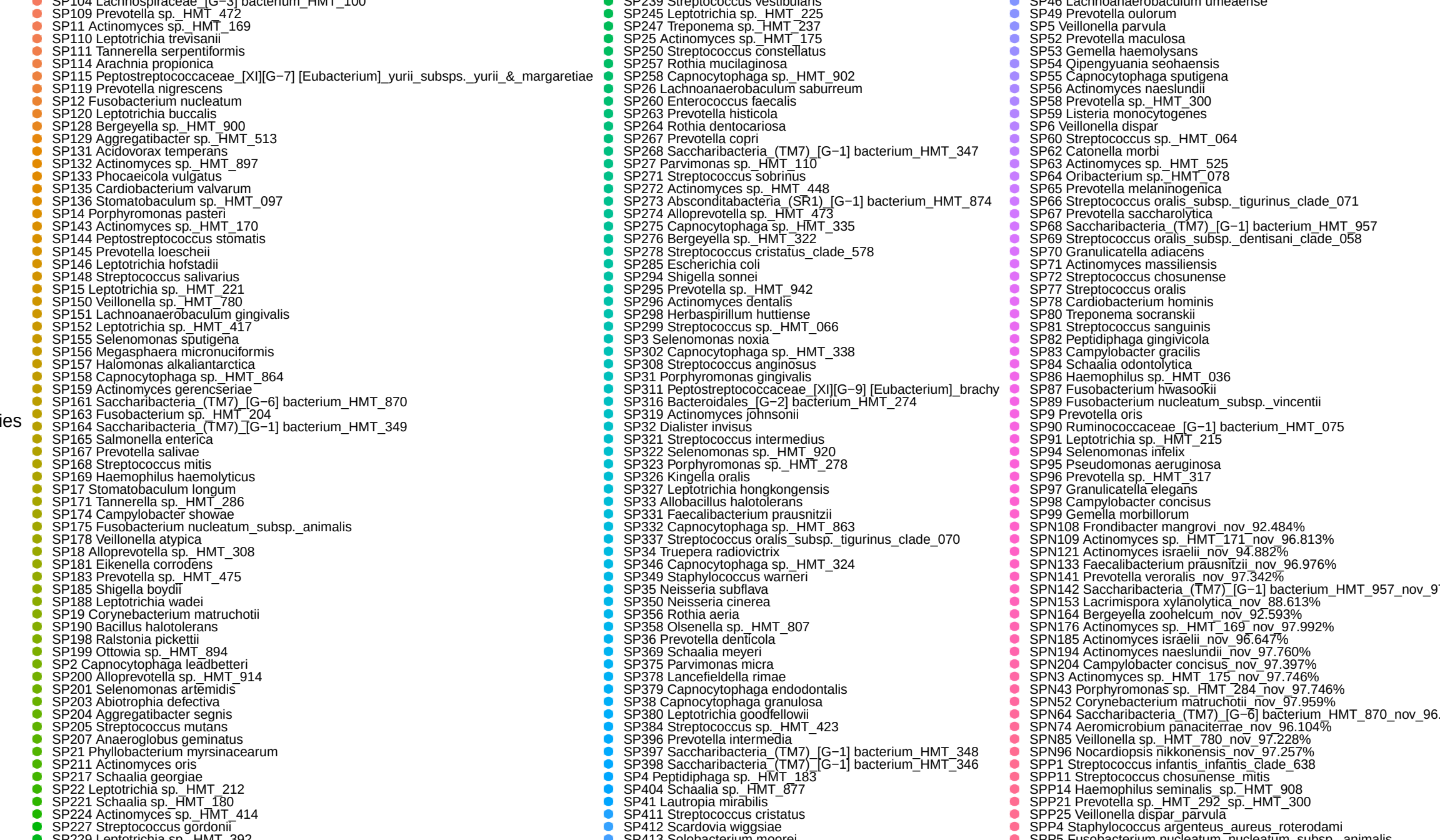
SP104 Lachnospiraceae_[G−3] bacterium_HMT_100
SP109 Prevotella sp._HMT_472
SP11 Actinomyces sp._HMT_169
SP110 Leptotrichia trevisanii
SP111 Tannerella serpentiformis
SP114 Arachnia propionica
SP115 Peptostreptococcaceae_[XI][G−7] [Eubacterium]_yurii_subsps._yurii_&_margaretiae
SP119 Prevotella nigrescens
SP12 Fusobacterium nucleatum
SP120 Leptotrichia buccalis
SP128 Bergeyella sp._HMT_900
SP129 Aggregatibacter sp._HMT_513
SP131 Acidovorax temperans
SP132 Actinomyces sp._HMT_897
SP133 Phocaeicola vulgatus
SP135 Cardiobacterium valvarum
SP136 Stomatobaculum sp._HMT_097
SP14 Porphyromonas pasteri
SP143 Actinomyces sp._HMT_170
SP144 Peptostreptococcus stomatis
SP145 Prevotella loescheii
SP146 Leptotrichia hofstadii
SP148 Streptococcus salivarius
SP15 Leptotrichia sp._HMT_221
SP150 Veillonella sp._HMT_780
SP151 Lachnoanaerobaculum gingivalis
SP152 Leptotrichia sp._HMT_417
SP155 Selenomonas sputigena
SP156 Megasphaera micronuciformis
SP157 Halomonas alkaliantarctica
SP158 Capnocytophaga sp._HMT_864
SP159 Actinomyces gerencseriae
SP161 Saccharibacteria_(TM7)_[G−6] bacterium_HMT_870
SP163 Fusobacterium sp._HMT_204
SP164 Saccharibacteria_(TM7)_[G−1] bacterium_HMT_349
SP165 Salmonella enterica
SP167 Prevotella salivae
SP168 Streptococcus mitis
SP169 Haemophilus haemolyticus
SP17 Stomatobaculum longum
SP171 Tannerella sp._HMT_286
SP174 Campylobacter showae
SP175 Fusobacterium nucleatum_subsp._animalis
SP178 Veillonella atypica
SP18 Alloprevotella sp._HMT_308
SP181 Eikenella corrodens
SP183 Prevotella sp._HMT_475
SP185 Shigella boydii
SP188 Leptotrichia wadei
SP19 Corynebacterium matruchotii
SP190 Bacillus halotolerans
SP198 Ralstonia pickettii
SP199 Ottowia sp._HMT_894
SP2 Capnocytophaga leadbetteri
SP200 Alloprevotella sp._HMT_914
SP201 Selenomonas artemidis
SP203 Abiotrophia defectiva
SP204 Aggregatibacter segnis
SP205 Streptococcus mutans
SP207 Anaeroglobus geminatus
SP21 Phyllobacterium myrsinacearum
SP211 Actinomyces oris
SP217 Schaalia georgiae
SP22 Leptotrichia sp._HMT_212
SP221 Schaalia sp._HMT_180
SP224 Actinomyces sp._HMT_414
SP227 Streptococcus gordonii
SP229 Leptotrichia sp._HMT_392
SP239 Streptococcus vestibularis
SP245 Leptotrichia sp._HMT_225
SP247 Treponema sp._HMT_237
SP25 Actinomyces sp._HMT_175
SP250 Streptococcus constellatus
SP257 Rothia mucilaginosa
SP258 Capnocytophaga sp._HMT_902
SP26 Lachnoanaerobaculum saburreum
SP260 Enterococcus faecalis
SP263 Prevotella histicola
SP264 Rothia dentocariosa
SP267 Prevotella copri
SP268 Saccharibacteria_(TM7)_[G−1] bacterium_HMT_347
SP27 Parvimonas sp._HMT_110
SP271 Streptococcus sobrinus
SP272 Actinomyces sp._HMT_448
SP273 Absconditabacteria_(SR1)_[G−1] bacterium_HMT_874
SP274 Alloprevotella sp._HMT_473
SP275 Capnocytophaga sp._HMT_335
SP276 Bergeyella sp._HMT_322
SP278 Streptococcus cristatus_clade_578
SP285 Escherichia coli
SP294 Shigella sonnei
SP295 Prevotella sp._HMT_942
SP296 Actinomyces dentalis
SP298 Herbaspirillum huttiense
SP299 Streptococcus sp._HMT_066
SP3 Selenomonas noxia
SP302 Capnocytophaga sp._HMT_338
SP308 Streptococcus anginosus
SP31 Porphyromonas gingivalis
SP311 Peptostreptococcaceae_[XI][G−9] [Eubacterium]_brachy
SP316 Bacteroidales_[G−2] bacterium_HMT_274
SP319 Actinomyces johnsonii
SP32 Dialister invisus
SP321 Streptococcus intermedius
SP322 Selenomonas sp._HMT_920
SP323 Porphyromonas sp._HMT_278
SP326 Kingella oralis
SP327 Leptotrichia hongkongensis
SP33 Allobacillus halotolerans
SP331 Faecalibacterium prausnitzii
SP332 Capnocytophaga sp._HMT_863
SP337 Streptococcus oralis_subsp._tigurinus_clade_070
SP34 Truepera radiovictrix
SP346 Capnocytophaga sp._HMT_324
SP349 Staphylococcus warneri
SP35 Neisseria subflava
SP350 Neisseria cinerea
SP356 Rothia aeria
SP358 Olsenella sp._HMT_807
SP36 Prevotella denticola
SP369 Schaalia meyeri
SP375 Parvimonas micra
SP378 Lancefieldella rimae
SP379 Capnocytophaga endodontalis
SP38 Capnocytophaga granulosa
SP380 Leptotrichia goodfellowii
SP384 Streptococcus sp._HMT_423
SP396 Prevotella intermedia
SP397 Saccharibacteria_(TM7)_[G−1] bacterium_HMT_348
SP398 Saccharibacteria_(TM7)_[G−1] bacterium_HMT_346
SP4 Peptidiphaga sp._HMT_183
SP404 Schaalia sp._HMT_877
SP41 Lautropia mirabilis
SP411 Streptococcus cristatus
SP412 Scardovia wiggsiae
SP413 Solobacterium moorei
SP46 Lachnoanaerobaculum umeaense
SP49 Prevotella oulorum
SP5 Veillonella parvula
SP52 Prevotella maculosa
SP53 Gemella haemolysans
SP54 Qipengyuania seohaensis
SP55 Capnocytophaga sputigena
SP56 Actinomyces naeslundii
SP58 Prevotella sp._HMT_300
SP59 Listeria monocytogenes
SP6 Veillonella dispar
SP60 Streptococcus sp._HMT_064
SP62 Catonella morbi
SP63 Actinomyces sp._HMT_525
SP64 Oribacterium sp._HMT_078
SP65 Prevotella melaninogenica
SP66 Streptococcus oralis_subsp._tigurinus_clade_071
SP67 Prevotella saccharolytica
SP68 Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957
SP69 Streptococcus oralis_subsp._dentisani_clade_058
SP70 Granulicatella adiacens
SP71 Actinomyces massiliensis
SP72 Streptococcus chosunense
SP77 Streptococcus oralis
SP78 Cardiobacterium hominis
SP80 Treponema socranskii
SP81 Streptococcus sanguinis
SP82 Peptidiphaga gingivicola
SP83 Campylobacter gracilis
SP84 Schaalia odontolytica
SP86 Haemophilus sp._HMT_036
SP87 Fusobacterium hwasookii
SP89 Fusobacterium nucleatum_subsp._vincentii
SP9 Prevotella oris
SP90 Ruminococcaceae_[G−1] bacterium_HMT_075
SP91 Leptotrichia sp._HMT_215
SP94 Selenomonas infelix
SP95 Pseudomonas aeruginosa
SP96 Prevotella sp._HMT_317
SP97 Granulicatella elegans
SP98 Campylobacter concisus
SP99 Gemella morbillorum
SPN108 Frondibacter mangrovi_nov_92.484%
SPN109 Actinomyces sp._HMT_171_nov_96.813%
SPN121 Actinomyces israelii_nov_94.882%
SPN133 Faecalibacterium prausnitzii_nov_96.976%
SPN141 Prevotella veroralis_nov_97.342%
SPN142 Saccharibacteria_(TM7)_[G−1] bacterium_HMT_957_nov_9
SPN153 Lacrimispora xylanolytica_nov_88.613%
SPN164 Bergeyella zoohelcum_nov_92.593%
SPN176 Actinomyces sp._HMT_169_nov_97.992%
SPN185 Actinomyces israelii_nov_96.647%
SPN194 Actinomyces naeslundii_nov_97.760%
SPN204 Campylobacter concisus_nov_97.397%
SPN3 Actinomyces sp._HMT_175_nov_97.746%
SPN43 Porphyromonas sp._HMT_284_nov_97.746%
SPN52 Corynebacterium matruchotii_nov_97.959%
SPN64 Saccharibacteria_(TM7)_[G−6] bacterium_HMT_870_nov_96.
SPN74 Aeromicrobium panaciterrae_nov_96.104%
SPN85 Veillonella sp._HMT_780_nov_97.228%
SPN96 Nocardiopsis nikkonensis_nov_97.257%
SPP1 Streptococcus infantis_infantis_clade_638
SPP11 Streptococcus chosunense_mitis
SPP14 Haemophilus seminalis_sp._HMT_908
SPP21 Prevotella sp._HMT_292_sp._HMT_300
SPP25 Veillonella dispar_parvula
SPP4 Staphylococcus argenteus_aureus_roterodami
SPP5 Fusobacterium nucleatum_nucleatum_subsp._animalis