SP246 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_875 (0.09%|0%|0%|0.06%|0%|0.08%)
SP281 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345 (0.03%|0%|0.01%|0.01%|0.04%|0.07%)
SP279 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_874 (0.05%|0.12%|0.08%|0%|0.10%|0%)
SP183 Gracilibacteria_(GN02)_[G-1] bacterium_HMT_872 (0.00%|0.15%|0.35%|0%|0.18%|0.
SP175 Bergeyella sp._HMT_900 (0%|0.14%|0.01%|0%|0.11%|0%)
SP43 Bergeyella sp._HMT_322 (0.14%|0.30%|0.46%|0.07%|0.43%|0.09%)
SP149 Capnocytophaga gingivalis (0.11%|0.39%|0.57%|0.14%|0.38%|0.05
SP117 Capnocytophaga granulosa (0.01%|0.19%|0.23%|0.03%|0.14%|0.0
SP67 Capnocytophaga leadbetteri (0.10%|0.34%|0.46%|0.02%|0.32%|0.1
SP100 Capnocytophaga sputigena (0.05%|0.36%|0.29%|0.04%|0.22%|0.
SP127 Capnocytophaga endodontalis (0%|0.11%|0.27%|0%|0.18%|0.01%
SP305 Capnocytophaga sp._HMT_335 (0%|0.05%|0.04%|0%|0.04%|0%)
SP92 Capnocytophaga ochracea (0%|0.21%|0.19%|0%|0.16%|0%)
SP52 Capnocytophaga sp._HMT_380 (0%|0.07%|0.13%|0.02%|0.01%|0.
SP282 Capnocytophaga sp._HMT_336 (0%|0.20%|0.17%|0%|0.12%|0.0
SP155 Capnocytophaga sp._HMT_864 (0%|0.10%|0.09%|0%|0.11%|0%)
SP125 Bacteroidales_[G-2] sp._Oral_Taxon_274 (0.02%|0.04%|0.21%|0.04%|
SP203 Tannerella forsythia (0.01%|0%|0.04%|0%|0%|0.01%)
SP56 Tannerella sp._HMT_286 (0.04%|0.19%|0.25%|0.02%|0.23%|0.06%
SP239 Tannerella sp._HMT_808 (0%|0%|0.01%|0%|0.07%|0%)
SP179 Porphyromonas endodontalis (0.10%|0.14%|0.20%|0.10%|0.2
SP251 Porphyromonas gingivalis (0.04%|0%|0%|0.09%|0%|0.05%)
SP253 Porphyromonas sp._HMT_284 (0%|0.02%|0.08%|0%|0.10%|0%)
SP40 Porphyromonas pasteri (1.00%|0.58%|0.46%|0.68%|0.40%|0.9
SP46 Porphyromonas catoniae (0.05%|0.26%|0.48%|0.01%|0.28%|0.0
SP173 Porphyromonas sp._HMT_277 (0%|0%|0%|0%|0.05%|0.02%)
SP288 Porphyromonas sp._Oral_Taxon_B43 (0.02%|0.04%|0%|0.01%
SP84 Alloprevotella tannerae (0.10%|0.33%|0.55%|0.12%|0.
SP180 Alloprevotella rava (0.05%|0%|0%|0.04%|0%|0.04%)
SP204 Alloprevotella sp._HMT_473 (0.17%|0.04%|0.03%|0.0
SP222 Alloprevotella sp._HMT_914 (0.01%|0%|0%|0.09%|0%|
SP171 Alloprevotella sp._HMT_308 (0.33%|0%|0.01%|0.33%|
SP295 Prevotella pleuritidis (0.02%|0%|0.19%|0%|0.04%|0
SP287 Prevotella sp._HMT_304 (0.03%|0%|0%|0.04%|0%|0
SP41 Prevotella nanceiensis (0.27%|0%|0.03%|0.17%|0%|0.2
SP238 Prevotella saccharolytica (0%|0.06%|0.06%|0%|0.06
SP290 Prevotella shahii (0.08%|0%|0%|0.10%|0%|0.02%)
SP174 Prevotella sp._HMT_317 (0.02%|0.23%|0.37%|0%|0.
SP257 Prevotella loescheii (0%|0.06%|0.12%|0%|0.12%|0
SP190 Prevotella sp._HMT_472 (0%|0.12%|0.19%|0%|0%|
SP247 Prevotella sp._HMT_300 (0%|0.18%|0%|0%|0.19%
SP154 Prevotella oulorum (0.03%|0.34%|0.15%|0.05%|0.21%|
SP126 Prevotella sp._HMT_309 (0.07%|0.02%|0%|0.01%|0%|0
SP57 Prevotella salivae (0.45%|0.10%|0%|0.50%|0.10%|0.75%
SP273 Prevotella maculosa (0%|0.08%|0.05%|0%|0.03%|0
SP33 Prevotella oris (0.13%|0.80%|0.64%|0.12%|0.75%|0.1
SP98 Prevotella pallens (0.39%|0%|0.03%|0.36%|0%|0.69%|
SP201 Prevotella aurantiaca (0.09%|0%|0%|0.05%|0%|0.05%
SP90 Prevotella intermedia (0.14%|0.06%|0.10%|0.09%|0.
SP42 Prevotella denticola (0.04%|1.04%|0.31%|0.11%|0.50%
SP7 Prevotella jejuni (0.28%|0%|0%|0.24%|0%|0.55%)
SP99 Prevotella sp._HMT_314 (0.02%|0%|0.04%|0.01%|0.02%|
SP29 Prevotella melaninogenica (5.10%|0.39%|0.41%|4.52%|0
SP129 Prevotella histicola (1.06%|0.23%|0.04%|2.39%|0.12%|
SP30 Prevotella sp._HMT_306 (0.24%|0%|0%|0.35%|0%|0.31%
SP256 Prevotella veroralis (0.01%|0.01%|0.06%|0%|0.12%|0%
SP197 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356 (0.01%|0.42%|0.21%
SP264 Saccharibacteria_(TM7)_[G-2] bacterium_HMT_350 (0%|0%|0%|0%|0.08%|0
SP71 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351 (0.16%|0.20%
SP192 TM7_[G] sp._Oral_Taxon_A56 (0.06%|0%|0%|0.01%|0%|0.12%)
SP172 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870 (0.07%|0.06%|0.08%
SP109 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348 (0.03%|1.02%|0.51
SP73 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957 (0.02%|0.18%|0.05%|
SP118 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_352 (0.38%|0.00%|0.01%
SP119 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347 (0.01%|0.32%|0.05%
SP79 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349 (0%|0.12%|0.27%|0.
SP28 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_869 (0.01%|0.56%|0.10%|
SP58 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346 (0.01%|1.09%|0.71%
SP93 Campylobacter showae (0.05%|0.14%|0.24%|0.03%|0.30%|0%)
SP34 Campylobacter gracilis (0%|0.40%|0.31%|0.03%|0.24%|0.01%)
SP16 Campylobacter concisus (0.41%|0.26%|0.23%|0.56%|0.32%|0.6
SP128 Cardiobacterium valvarum (0%|0.08%|0.07%|0%|0.07%|0%)
SP55 Cardiobacterium hominis (0.01%|0.42%|0.51%|0.00%|0.38%|0.01%)
SP228 Haemophilus parahaemolyticus (0.18%|0%|0.04%|0.12%|0%|0.0
SP31 Haemophilus parainfluenzae (4.07%|0.61%|2.09%|3.91%|1.37%
SP133 Haemophilus sputorum (0.03%|0%|0.02%|0.04%|0%|0.03%)
SP233 Haemophilus sp._HMT_036 (0.01%|0%|0%|0.02%|0.01%|0.05%)
SP112 Aggregatibacter sp._HMT_458 (0.07%|0.34%|0.31%|0.05%|0.13%
SP72 Aggregatibacter aphrophilus (0%|0%|0.08%|0%|0.05%|0%)
SP187 Aggregatibacter segnis (0%|0.05%|0%|0%|0.04%|0%)
SP60 Lautropia mirabilis (0.18%|1.62%|2.38%|0.32%|1.65%|0.12%)
SP150 Ottowia sp._HMT_894 (0%|0.07%|0.11%|0%|0.15%|0.01%)
SP182 Kingella oralis (0.03%|0.77%|0.57%|0.01%|0.33%|0.01%)
SP162 Neisseria sp._HMT_018 (0%|0.19%|0%|0%|0%|0%)
SP45 Neisseria bacilliformis (0.00%|0.14%|0.33%|0%|0.29%|0.01%)
SP75 Neisseria flavescens|subflava (3.58%|0.23%|0.25%|2.39%|0.16%|2.86%
SP11 Neisseria subflava (0.23%|0.05%|0.48%|0.26%|0.16%|0.26%)
SP36 Neisseria sicca (0.23%|0.65%|0.27%|0.11%|1.17%|0.01%)
SP53 Neisseria mucosa (0.21%|0.45%|0.48%|0.13%|0.49%|0.08%)
SP214 Neisseria sp._Oral_Taxon_14 (0%|0.09%|0.17%|0%|0.13%|0%)
SP209 Eikenella corrodens (0%|0.17%|0.32%|0%|0.17%|0%)
SP339 Kingella denitrificans (0%|0%|0.10%|0%|0.00%|0%)
SP68 Neisseria elongata (0.11%|0.83%|0.81%|0.02%|0.82%|0.07%)
SP51 Fusobacterium periodonticum (1.80%|0%|0.14%|1.32%|0%|1.91%)
SP158 Fusobacterium nucleatum_ss_polymorphum (0%|0.48%|0.35%|0.01%
SP147 Fusobacterium sp._HMT_204 (0%|0.10%|0.13%|0.02%|0.01%|0.02%
SP185 Fusobacterium nucleatum_subsp._vincentii (0%|0.05%|0.16%|0.01%
SP37 Fusobacterium nucleatum (0.14%|0.94%|1.53%|0.12%|0.92%|0.13%
SP219 Fusobacterium nucleatum_ss_animalis (0%|0.09%|0.22%|0%|0.02%
SP3 Leptotrichia sp._HMT_221 (0.55%|0.06%|0.02%|0.61%|0%|0.70%)
SP235 Pseudoleptotrichia goodfellowii (0%|0.03%|0.04%|0%|0.04%|0%)
SP268 Leptotrichia goodfellowii (0%|0.03%|0.07%|0%|0.02%|0%)
SP184 Leptotrichia sp._HMT_392 (0.01%|0.06%|0.25%|0%|0.12%|0.02
SP168 Leptotrichia sp._HMT_215 (0.28%|0.03%|0.25%|0.25%|0.09%|
SP194 Leptotrichia sp._HMT_212 (0.03%|0.25%|0.37%|0%|0.30%|0%
SP250 Leptotrichia sp._HMT_218 (0.14%|0%|0%|0.08%|0%|0.23%)
SP170 Leptotrichia sp._HMT_498 (0%|0.26%|0.19%|0%|0.15%|0%)
SP138 Leptotrichia sp._HMT_225 (0%|0%|0.10%|0.27%|0%)
SP156 Leptotrichia buccalis (0.11%|0.35%|0.19%|0%|0.52%|0.04%)
SP22 Leptotrichia hofstadii (0.02%|0.40%|0.53%|0.04%|0.35%|0%)
SP151 Leptotrichia sp._HMT_223 (0%|0.05%|0%|0%|0.06%|0%)
SP139 Leptotrichia sp._HMT_417 (0.12%|0.02%|0.08%|0.13%|0.07%|
SP69 Leptotrichia wadei (0.26%|1.71%|1.06%|0.16%|1.83%|0.16%)
SP306 Leptotrichia shahii (0%|0.09%|0%|0%|0%|0%)
SP134 Leptotrichia sp._AF189244.1 (0.08%|1.39%|0.81%|0.12%|0.5
SP252 Leptotrichia trevisanii (0%|0.05%|0.09%|0%|0%|0%)
SP148 Fretibacterium sp._HMT_360 (0.01%|0%|0.05%|0%|0.03%|0%)
SP120 Fretibacterium sp._HMT_359 (0%|0%|0.08%|0%|0%|0%)
SP334 Cryptobacterium curtum (0%|0%|0%|0%|0.10%|0%)
SP200 Olsenella sp._Oral_Taxon_807 (0%|0.26%|0.03%|0%|0.04%|0%)
SP27 Lancefieldella rimae (0%|0.09%|0.16%|0%|0.29%|0%)
SP177 Bifidobacterium dentium (0.04%|0.05%|0%|0%|0.17%|0%)
SP121 Scardovia wiggsiae (0.01%|1.00%|0.65%|0.05%|0.54%|0.0
SP63 Parascardovia denticolens (0%|0.02%|0.17%|0.06%|0.11%|0.0
SP303 Propionibacterium acidifaciens (0%|0.08%|0%|0.01%|0%|0%)
SP15 Arachnia propionica (0.02%|1.05%|1.42%|0%|0.86%|0%)
SP193 Arachnia rubra (0.07%|1.32%|1.99%|0.03%|1.59%|0.04%)
SP12 Corynebacterium matruchotii (0.09%|3.91%|2.36%|0.15%|2.77
SP191 Corynebacterium accolens (0%|0%|0%|0.11%|0%|0%)
SP206 Corynebacterium sp._Oral_Taxon_A16 (0.04%|0.19%|0.22%|0
SP66 Corynebacterium durum (0.05%|0.55%|0.41%|0.08%|0.40%|0
SP50 Rothia mucilaginosa (11.85%|0%|0.43%|9.88%|0.22%|9.80%)
SP163 Rothia aeria (0.13%|0.92%|1.17%|0.13%|0.70%|0.21%)
SP54 Rothia dentocariosa (1.31%|3.01%|1.84%|1.09%|2.67%|0.69%)
SP122 Peptidiphaga sp._HMT_183 (0%|2.42%|1.76%|0.10%|0.84%|0%
SP265 Schaalia lingnae_[Not_Validly_Published] (0.15%|0%|0%|0%|0
SP39 Schaalia odontolytica (1.61%|0.21%|0.15%|1.39%|0.08%|2.07
SP181 Schaalia sp._HMT_180 (0.32%|0.05%|0%|0.13%|0.09%|0.30%
SP130 Actinomyces sp._Oral_Taxon_180 (0.02%|0.94%|0.44%|0.11%
SP20 Actinomyces odontolyticus (2.44%|0.19%|0.12%|2.93%|0.22%
SP124 Schaalia sp._HMT_172 (0.67%|0.05%|0%|0.74%|0%|0.99%)
SP234 Schaalia meyeri (0%|0.09%|0.05%|0%|0.03%|0%)
SP196 Schaalia georgiae (0%|0.14%|0.11%|0.01%|0.15%|0%)
SP89 Actinomyces sp._Oral_Taxon_178 (0.01%|0.26%|0.15%|0%
SP166 Schaalia sp._HMT_877 (0.01%|0.25%|0.18%|0.02%|0.29%
SP8 Actinomyces graevenitzii (1.54%|0%|0%|1.11%|0%|1.37%)
SP161 Actinomyces sp._HMT_448 (0%|5.55%|6.30%|0.06%|7.80
SP157 Actinomyces massiliensis (0%|1.01%|1.17%|0%|0.94%|0
SP152 Actinomyces israelii (0%|0.10%|0.11%|0%|0.10%|0%)
SP212 Actinomyces sp._HMT_897 (0%|0%|0.05%|0.15%|0%|0
SP259 Actinomyces sp._HMT_414 (0%|0.25%|0.03%|0.05%
SP64 Actinomyces dentalis (0%|1.85%|0.48%|0.05%|1.66
SP210 Actinomyces sp._HMT_896 (0%|0.55%|0%|0.02%|
SP217 Actinomyces sp._HMT_525 (0%|0.18%|0.54%|0%|
SP24 Actinomyces sp._HMT_170 (0.12%|0.66%|0.80%|0.06%|0.86
SP94 Actinomyces johnsonii (0.00%|1.04%|0.79%|0.05%|0.70%|
SP10 Actinomyces sp._HMT_171 (0.03%|0.41%|0.30%|0.01%|0.35
SP2 Actinomyces sp._Oral_Taxon_171 (0%|0.11%|0.17%|0%|0.11
SP142 Actinomyces sp._HMT_175 (0.05%|1.11%|1.09%|0.27%|0
SP96 Actinomyces oris (0%|0.59%|0.44%|0%|0.30%|0.06%)
SP6 Actinomyces viscosus (0.14%|1.17%|0.88%|0.15%|0.86%|0.
SP17 Actinomyces naeslundii (0%|2.30%|2.76%|0.09%|2.04%|0
SP86 Actinomyces sp._str._ChDCB197 (0.21%|1.53%|1.36%|0.3
SP80 Actinomyces sp._HMT_169 (0%|0.12%|0.26%|0.01%|0.09%
SP26 Parvimonas micra (0.17%|0.74%|0.61%|0.14%|0.58%|0.17%)
SP25 Parvimonas sp._Oral_Taxon_110 (0.01%|0.05%|0.22%|0.03%|0.06%
SP293 Filifactor alocis (0.04%|0%|0.02%|0.03%|0.04%|0.02%)
SP240 Eubacterium_[XI][G-7] yurii (0%|0.02%|0.06%|0%|0.03%|0%
SP309 Peptoanaerobacter [Eubacterium] yurii (0%|0%|0.02%|0%|0
SP47 Peptostreptococcus stomatis (0.10%|0.01%|0.07%|0.07%|0.15%|0.
SP114 Peptostreptococcaceae_[XI][G-9] [Eubacterium]_brachy (0.03%|0.
SP245 Peptostreptococcaceae_[XI][G-6] [Eubacterium]_nodatu
SP211 Peptostreptococcaceae_[XI][G-1] [Eubacterium]_sulci (0.17%|0
SP249 Peptostreptococcaceae_[XI][G-1] [Eubacterium]_infirmum (0%
SP301 Mogibacterium timidum (0.01%|0.02%|0.01%|0%|0.07%|0.0
SP216 Mogibacterium neglectum (0%|0.16%|0.08%|0%|0.12%|0%)
SP108 Mogibacterium vescum (0.35%|0%|0.06%|0.30%|0.05%|0.44
SP160 Lachnospiraceae_[G-2] bacterium_HMT_096 (0.23%|0%|0%|0.09%|0%|
SP328 Lachnospiraceae_[G] sp._Oral_Taxon_B32 (0%|0%|0%|0.06%|0%|0.0
SP164 Lachnospiraceae_[G-3] bacterium_HMT_100 (0.04%|0.03%|0%|
SP202 Lachnoanaerobaculum orale (0.07%|0.01%|0%|0.25%|0%|0.12%)
SP215 Lachnoanaerobaculum gingivalis (0.03%|0%|0%|0.06%|0%|0%|0.03%)
SP165 Lachnoanaerobaculum umeaense (0%|0.04%|0.07%|0%|0.05%|0.0
SP141 Shuttleworthia satelles (0%|0.00%|0.04%|0%|0.05%|0%)
SP267 Oribacterium sp._Oral_Taxon_78 (0%|0.06%|0.07%|0.01%|0.03%
SP59 Oribacterium sinus (0.82%|0.09%|0.03%|0.59%|0.09%|0.90%)
SP143 Stomatobaculum longum (0.19%|0.05%|0.05%|0.28%|0.13%|0.16%
SP115 Stomatobaculum sp._HMT_097 (0.23%|0.02%|0.05%|0.21%|0.01%
SP274 Butyrivibrio sp._HMT_455 (0.05%|0%|0%|0%|0%|0.06%)
SP178 Catonella morbi (0.06%|0.04%|0.10%|0.11%|0.14%|0.16%)
SP207 Ruminococcaceae_[G-2] bacterium_HMT_085 (0.07%|0.01%|0.0
SP81 Ruminococcaceae_[G-1] bacterium_HMT_075 (0.11%|0.20%|0.07
SP140 Dialister invisus (0%|0.42%|0.37%|0.04%|0.33%|0%)
SP342 Veillonella sp._HMT_917 (0%|0%|0%|0.06%|0%|0.03%)
SP91 Veillonella atypica (2.13%|0.17%|0.21%|2.54%|0.10%|2.46%)
SP1 Veillonella parvula (3.52%|4.68%|5.58%|3.52%|5.16%|4.13%)
SP237 Megasphaera micronuciformis (0.36%|0.03%|0.02%|0.40%|0.02
SP284 Colibacter massiliensis (0%|0%|0.04%|0%|0.05%|0%)
SP248 Anaeroglobus geminatus (0%|0.06%|0%|0%|0.03%|0%)
SP65 Selenomonas sputigena (0%|0.51%|0.25%|0.05%|0.51%|0.03%
SP205 Veillonellaceae_[G-1] bacterium_HMT_155 (0%|0.10%|0%
SP199 Veillonellaceae_[G-1] bacterium_HMT_150 (0%|0.02%|0.03
SP220 Selenomonas sp._Oral_Taxon_E83 (0%|0.07%|0.10%|0%|0.08%
SP102 Selenomonas artemidis (0%|0.11%|0.15%|0%|0.15%|0%)
SP229 Selenomonas sp._HMT_137 (0.00%|0.07%|0.01%|0%|0.04%|0
SP14 Selenomonas sp._Oral_Taxon_137 (0%|0.14%|0.10%|0%|0.07%
SP35 Selenomonas noxia (0.02%|0.74%|0.35%|0.03%|0.59%|0.03%)
SP307 Selenomonas infelix (0%|0.06%|0.04%|0%|0.02%|0%)
SP242 Selenomonas flueggei (0%|0.03%|0.03%|0%|0.08%|0%)
SP153 Selenomonas sp._HMT_149 (0.04%|0%|0.01%|0.03%|0%|0.07
SP271 Selenomonas sp._HMT_136 (0.06%|0.01%|0%|0.07%|0.06%
SP70 Solobacterium moorei (0.27%|0.19%|0.19%|0.25%|0.13%|0.46%
SP74 Gemella morbillorum (0.11%|0.59%|0.75%|0.05%|0.60%|0.06%)
SP101 Gemella haemolysans (0.84%|0.05%|0.07%|0.81%|0.03%|0.54%)
SP83 Gemella sanguinis (1.94%|0.11%|0.12%|1.55%|0.16%|1.71%)
SP44 Lactobacillus salivarius (0.13%|0%|0%|0%|0.14%|0.33%)
SP231 Lactobacillus ultunensis (0.03%|0.04%|0.04%|0.00%|0.01
SP188 Abiotrophia defectiva (0.09%|0.46%|0.46%|0.12%|0.31%|0.11%)
SP283 Granulicatella elegans (0.11%|0.03%|0%|0.10%|0%|0.02%)
SP13 Granulicatella paradiacens (3.04%|0.68%|0.73%|2.54%|0.68%|2.5
SP4 Granulicatella adiacens (0.02%|0.91%|1.07%|0.07%|1.10%|0.05%)
SP88 Streptococcus mutans (0.03%|0.91%|0.54%|0.08%|2.47%|0.0
SP176 Streptococcus sobrinus (0.01%|0.05%|2.03%|0%|0.05%|0.04
SP116 Streptococcus sp._str._2136FAA (0.05%|0.93%|0.22%|0.10%|0.4
SP97 Streptococcus gordonii (0.32%|1.81%|1.31%|0.34%|1.62%|0.07
SP48 Streptococcus sinensis (0.02%|0%|0.03%|0.00%|0.07%|0.01%
SP85 Streptococcus anginosus (0.02%|0.89%|0.75%|0.08%|0.56%
SP169 Streptococcus constellatus (0.01%|0.03%|0.08%|0.01%|0.0
SP105 Streptococcus intermedius (0.03%|1.26%|0.60%|0.08%|0.39
SP82 Streptococcus sp._HMT_056 (0.01%|0.21%|0.07%|0%|0.13%|0.
SP107 Streptococcus sp._str._C150 (1.38%|0.01%|0.30%|0.88%|0.
SP123 Streptococcus vestibularis (0.04%|0.05%|0.00%|0.08%|0.01
SP76 Streptococcus salivarius (16.39%|0.53%|1.05%|25.60%|0.25%
SP78 Streptococcus sp._Oral_Taxon_B66 (0.05%|0.47%|0.60%|0.0
SP49 Streptococcus sanguinis (1.16%|6.08%|6.29%|0.72%|8.15%|0.
SP23 Streptococcus cristatus (1.14%|1.82%|1.86%|0.80%|1.80%|0.
SP77 Streptococcus australis (1.46%|0.03%|0.40%|0.78%|0.10%|0.
SP19 Streptococcus sp._HMT_066 (0.57%|0%|0.11%|0.57%|0.10%|
SP21 Streptococcus koreensis (0.30%|0.01%|0.07%|0.19%|0.01%
SP110 Streptococcus parasanguinis_I (0.62%|0.08%|0.04%|0.67%
SP159 Streptococcus sp._HMT_057 (0.29%|0%|0%|0.32%|0%|0.4
SP18 Streptococcus parasanguinis_II (3.60%|0.34%|0.32%|3.46%
SP103 Streptococcus sp._Oral_Taxon_71 (0.04%|0.60%|0.28%|0.04
SP263 Streptococcus oralis_subsp._tigurinus_clade_071 (0%|0.01%|
SP132 Streptococcus oralis_subsp._tigurinus_clade_070 (0.19%|0.5
SP213 Streptococcus sp._HMT_074 (0.45%|0.01%|0.03%|0.29%|0.
SP113 Streptococcus sp._HMT_061 (0.63%|0.01%|0.04%|0.47%|0.
SP104 Streptococcus oralis_subsp._dentisani_clade_058 (0.10%|0.0
SP5 Streptococcus sp._str._C300 (2.07%|4.96%|6.66%|1.38%|4.96%
SP9 Streptococcus chosunense (0.47%|0.13%|0.21%|0.38%|0.14%|
SP106 Streptococcus sp._HMT_064 (0.03%|0.07%|0.26%|0.02%|0.1
SP38 Streptococcus infantis (2.34%|0.01%|0.31%|1.54%|0.14%|1.9
SP131 Streptococcus infantis_clade_431 (0.47%|0%|0%|0.37%|0%|
SP32 Streptococcus oralis (1.45%|1.95%|1.51%|0.88%|1.98%|0.54%
SP87 Streptococcus mitis (2.97%|2.13%|1.13%|2.70%|1.43%|1.49%
Saliva_Visit_1
Plaque_Visit_2
Plaque_Visit_3
Saliva_Visit_2
Plaque_Visit_1
Saliva_Visit_3
substitutions/site
0
0.2
0.4
0.6
0.8