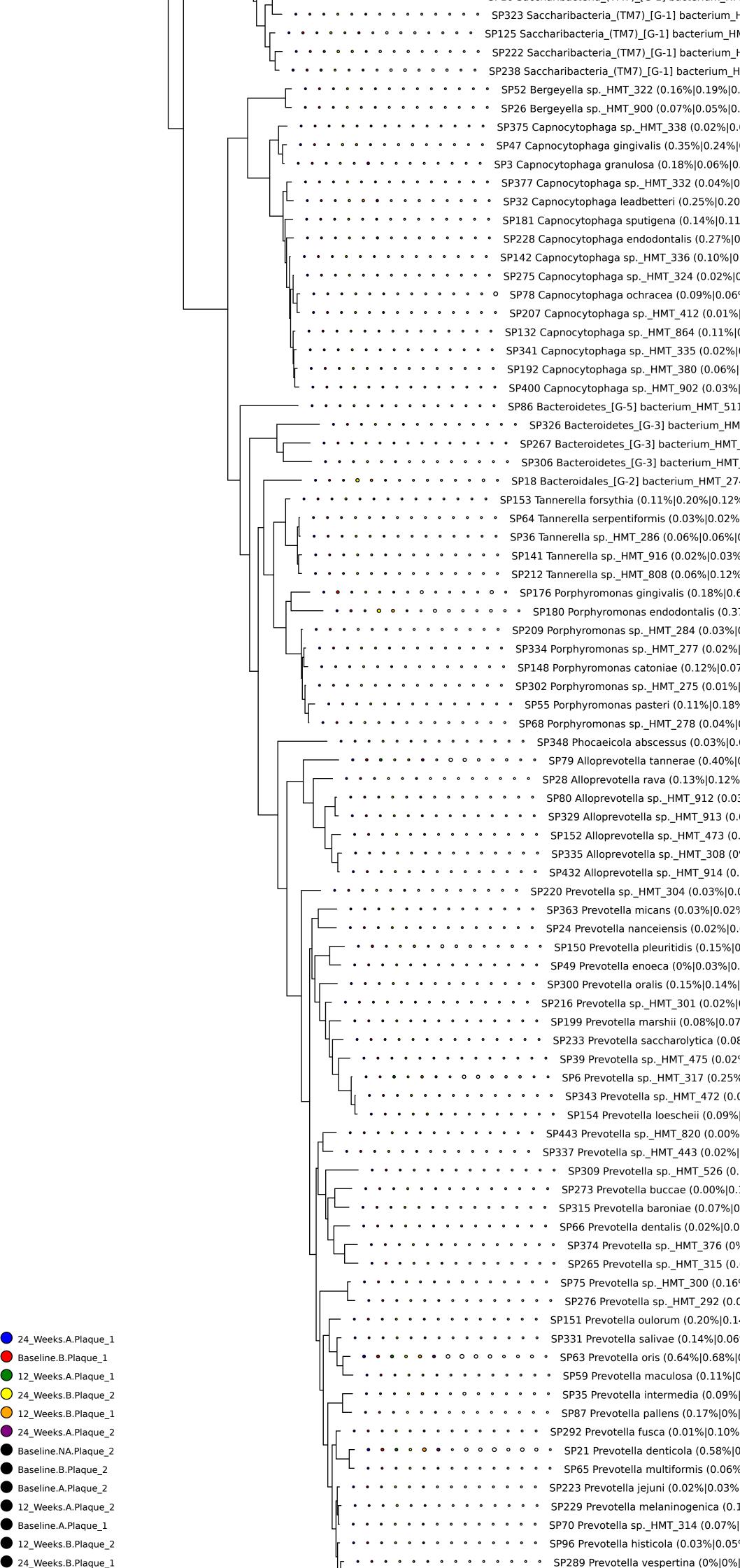
SP467 Methanobrevibacter oralis (0%|0.02%|0%|0.02%|0%|0%|0%|0%|0.06%|0 SP290 Fusobacterium sp._HMT_204 (0.05%|0.07%|0.06%|0.03%|0 SP195 Fusobacterium nucleatum_subsp._vincentii (0.35%|0.32%| 000000 SP20 Fusobacterium nucleatum (5.72%|4.21%|4.04%|4.98%|7.02 SP381 Fusobacterium canifelinum (0.01%|0.02%|0.02%|0.02%|0 SP112 Fusobacterium sp._HMT_203 (0.08%|0.02%|0.03%|0.03% SP124 Fusobacterium sp._HMT_370 (0%|0.04%|0%|0.03%|0.04% SP117 Fusobacterium periodonticum (0.15%|0.09%|0.15%|0.05 SP25 Leptotrichia sp._HMT_221 (0.10%|0.04%|0.03%|0.02%|0.08 SP225 Leptotrichia sp._HMT_219 (0.02%|0.01%|0.01%|0%|0 SP130 Leptotrichia sp. HMT 392 (0.11%|0.05%|0.04%|0.05% SP139 Leptotrichia sp._HMT_215 (0.08%|0.10%|0.06%|0.139 SP104 Leptotrichia sp._HMT_212 (0.10%|0.09%|0.05%|0.08% SP272 Leptotrichia sp. HMT 498 (0.13%|0.16%|0.32%|0.14 SP27 Leptotrichia buccalis (0.31%|0.12%|0.19%|0.35%|0.07% SP158 Leptotrichia sp._HMT_225 (0.04%|0.06%|0.06%|0.00%|0 SP105 Leptotrichia sp._HMT_223 (0.01%|0.02%|0.04%|0.03% SP71 Leptotrichia hofstadii (0.12%|0.05%|0.06%|0.03%|0.06% SP121 Leptotrichia sp._HMT_417 (0.09%|0.11%|0.10%|0.27% SP99 Leptotrichia wadei (1.47% 0.59% 0.82% 1.11% 0.99% SP365 Leptotrichia shahii (0.01%|0.07%|0.07%|0.08%|0.04% SP135 Leptotrichia trevisanii (0%|0.07%|0.03%|0%|0.03%| SP171 Leptotrichia hongkongensis (0.35%|0.37%|0.25%|0.3 SP160 Mycoplasma salivarium (0.00%|0.02%|0.01%|0.03%|0. SP184 Mycoplasma faucium (0.03%|0.06%|0.02%|0.05%|0.14% SP282 Gracilibacteria_(GN02)_[G-1] bacterium_HMT_872 SP324 Absconditabacteria_(SR1)_[G-1] bacteri • • • • SP428 Absconditabacteria_(SR1)_[G-1] bacteriu SP362 Eggerthia catenaformis (0.04%|0.06%|0.00%|0.11%) SP358 Mollicutes [G-1] bacterium HMT_504 (0.03%) • • SP367 Erysipelotrichaceae [G-1] bacterium HMT_904 (0.01 SP97 Erysipelotrichaceae [G-1] bacterium HMT_905 (0.03% SP156 Bulleidia extructa (0.05% 0.06% 0.03% 0.08% 0.07%) SP369 Solobacterium moorei (0.11% | 0.10% | 0.13% | 0.13% | 0.0 SP354 Gemella bergeri (0.01%|0.04%|0.01%|0.02%|0.03%|0. SP15 Gemella morbillorum (0.30%|0.30%|0.34%|0.28%|0.47% ο SP11 Gemella sanguinis (0.16% 0.03% 0.09% 0.04% 0.00% SP190 Gemella haemolysans (0.16%|0.27%|0.27%|0.08%|0.1 SP100 Ligilactobacillus salivarius (0.03%|0.04%|0.04%|0%|0 • SP235 Lactobacillus ultunensis (0.04%|0%|0.02%|0%|0%| SP144 Abiotrophia defectiva (0.05%|0.15%|0.08%|0.06%|0.14 O SP42 Granulicatella adiacens (0.79%|1.03%|1.17%|0.90%|0.8 SP314 Granulicatella elegans (0.04% | 0.06% | 0.02% | 0% | 0.03% ○ ○ ○ ○ • SP128 Streptococcus mutans (5.25%|4.47%|3.35%|2.51%|2 SP242 Streptococcus sobrinus (0.01%|0.09%|0.34%|0.12%| SP494 Streptococcus massiliensis (0.02%|0.01%|0.07%|0.1 SP162 Streptococcus anginosus (0.61%|0.88%|0.66%|0.58 SP45 Streptococcus constellatus (0.17%|0.10%|0.15%|0.2 SP136 Streptococcus intermedius (1.15%|0.47%|0.60%|1.2 SP175 Streptococcus vestibularis (0.06%|0.23%|0.02%|0.1 SP188 Streptococcus salivarius (1.42%|1.23%|1.80%|0.29 SP168 Streptococcus sp._HMT_056 (0.02%|0.03%|0.02%|0. 0 SP123 Streptococcus sanguinis (3.64%|2.87%|3.22%|2.98% SP51 Streptococcus cristatus (0.93%|1.25%|1.01%|0.80% SP14 Streptococcus australis (0.07%|0.02%|0.06%|0.05% SP110 Streptococcus sp._HMT_066 (0.02%|0.01%|0.03%| SP17 Streptococcus parasanguinis (0.00%|0.01%|0.02%| O SP120 Streptococcus parasanguinis_clade_411 (0.83%)1 SP231 Streptococcus oralis_subsp._dentisani_clade_058 (0 0 SP138 Streptococcus oralis_subsp._tigurinus_clade_070 (0 000 SP53 Streptococcus gordonii (2.23%|1.26%|1.97%|2.01% $\bigcirc \circ \circ \circ \circ \circ ()$ SP131 Streptococcus oralis (2.40%|2.43%|1.93%|2.21%|2 • • • • • O SP88 Streptococcus sp._HMT_064 (0.77%|0.49%|0.64%|1. ○ ○ ○ ○ ○ SP12 Streptococcus mitis (4.52%|4.83%|6.01%|4.37%|3.8 SP41 Streptococcus chosunense (0.04%|0.09%|0.03%|0.0 SP76 Actinidia eriantha (0%|0%|0.00%|0%|0%|0%|0%|0%|0 SP245 Dialister sp._HMT_119 (0.02%|0.02%|0.03%|0%|0.0 SP56 Dialister invisus (0.62%|0.40%|0.67%|0.67%|0.54%|0 SP295 Veillonella sp._HMT_780 (0.03%|0.04%|0.02%|0.01 SP7 Veillonella atypica (0.07%|0.12%|0.11%|0.05%|0.18% 0 SP50 Veillonella parvula (1.02%|1.21%|1.44%|0.86%|0.91 SP101 Megasphaera micronuciformis (0.06% 0.11% 0.08%) SP46 Anaeroglobus geminatus (0.19%|0.18%|0.09%|0.17 SP260 Megasphaera sp._HMT_123 (0.04%|0.02%|0.01%| SP113 Colibacter massiliensis (0.05%|0.03%|0.21%|0.04% SP305 Veillonellaceae [G-1] bacterium HMT_145 (0.01%) SP163 Veillonellaceae [G-1] bacterium HMT_155 (0.10% 0. SP237 Veillonellaceae [G-1] bacterium HMT_129 (0.02% 0.0 SP294 Veillonellaceae [G-1] bacterium HMT_150 (0.11% 0 SP259 Veillonellaceae [G-1] bacterium HMT_132 (0.06% 0. SP339 Mitsuokella sp._HMT_521 (0.04%|0.05%|0.20%|0.08% SP115 Selenomonas sp._HMT_442 (0.05%|0.08%|0%|0.01 SP280 Selenomonas sp. HMT 134 (0%|0.04%|0.06%|0.04 SP1 Selenomonas sputigena (0.40%|0.47%|0.46%|0.31%| SP340 Selenomonas sp._HMT_479 (0%|0.01%|0.01%|0.02% SP501 Selenomonas sp._HMT_126 (0.01%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02%|0.02% SP368 Selenomonas sp._HMT_919 (0.04%|0.01%|0.02%|0. SP291 Selenomonas sp._HMT_481 (0.02%|0.01%|0.02%|0. SP349 Centipeda periodontii (0.00%|0.02%|0.09%|0.02%| SP318 Selenomonas sp._HMT_892 (0.01%|0.00%|0.07%|0.0 SP62 Selenomonas sp._HMT_149 (0.04%|0.01%|0.04%|0% SP240 Selenomonas sp. HMT 136 (0.04%|0.05%|0.03%|0 SP89 Selenomonas sp._HMT_937 (0.01%|0.04%|0.02%|0.0 SP90 Selenomonas noxia (0.27% | 0.13% | 0.31% | 0.21% | 0.2 SP370 Selenomonas sp._HMT_133 (0.03%|0.01%|0.02% SP197 Selenomonas artemidis (0.05%|0.08%|0.01%|0.0 SP215 Selenomonas sp._HMT_137 (0.07%|0.03%|0.05% SP219 Selenomonas sp._HMT_138 (0.01%|0.00%|0.08%|0.0 SP213 Selenomonas infelix (0.05%|0.08%|0.05%|0.08%|0.0 SP330 Selenomonas flueggei (0.05%|0.09%|0.05%|0.07%| SP255 Selenomonas sp._HMT_920 (0.01%|0%|0.04%|0%|0 SP218 Selenomonas sp._HMT_146 (0.01%|0.02%|0.01%|0. SP239 Selenomonas dianae (0.06% 0.10% 0.04% 0.05% 0 SP204 Ruminococcaceae [G-1] bacterium HMT 075 (0.09% SP211 Ruminococcaceae [G-2] bacterium HMT_085 (0.02% SP246 Lachnospiraceae [G-8] bacterium HMT_500 (0.01% 0. SP268 Catonella sp._HMT_451 (0.01%|0.03%|0.01%|0.01 SP103 Catonella morbi (0.25%|0.15%|0.22%|0.22%|0.27%| SP316 Oribacterium sinus (0.02%|0.01%|0.01%|0.09%|0.03% SP107 Oribacterium sp._HMT_078 (0.30%|0.21%|0.27%|0. SP92 Shuttleworthia satelles (0.12%|0.11%|0.10%|0.06%|0 SP327 Stomatobaculum sp._HMT_910 (0.02%|0.02%|0%|0.0 SP133 Stomatobaculum sp._HMT_097 (0.01%|0.09%|0.01% SP106 Stomatobaculum longum (0.04% | 0.16% | 0.36% | 0.11% SP347 Lachnospiraceae [G-2] bacterium HMT_096 (0%|0%|0. SP155 Lachnospiraceae [G-3] bacterium HMT_100 (0.10% SP200 Johnsonella ignava (0%|0.03%|0%|0.03%|0.00 SP402 Johnsonella sp._HMT_166 (0%|0.03%|0.01%|0.06 SP37 Lachnoanaerobaculum sp._HMT_496 (0.02%|0.06%|0 SP296 Lachnoanaerobaculum sp._HMT_083 (0.02%|0.06%| SP254 Lachnoanaerobaculum umeaense (0.02%|0.05%|0.0 SP34 Lachnoanaerobaculum saburreum (0.48%|0.21%|0.3 SP198 Lachnoanaerobaculum orale (0.11%|0.10%|0.10%|0 SP191 Clostridiales [F-1][G-1] bacterium HMT_093 (0.02 SP396 Peptoniphilus lacrimalis (0.03%|0.03%|0.01%|0.01 SP166 Parvimonas micra (1.35% 1.68% 1.02% 1.37% 1.32% SP119 Parvimonas sp._HMT_110 (0.38%|0.05%|0.13%|0.06 SP217 Pseudoramibacter alactolyticus (0.02%|0.07%|0.02 SP57 Peptococcus sp._HMT_167 (0.06%|0.17%|0.09%|0. SP95 Peptococcus sp._HMT_168 (0.03%|0.03%|0.04%|0 SP356 Filifactor alocis (0.11%|0.13%|0.09%|0.19%|0.22%| SP164 Peptoanaerobacter [Eubacterium] yurii (0.10% SP423 Peptoanaerobacter stomatis (0.02%|0.01%|0.0 SP398 Peptostreptococcaceae_[XI][G-7] bacterium_H SP140 Peptostreptococcus stomatis (0.23%|0.12%|0.17% SP177 Peptostreptococcaceae [XI][G-9] [Eubacterium]_bra SP250 Peptostreptococcaceae [XI][G-1] [Eubacterium]_in SP247 Peptostreptococcaceae_[XI][G-5] [Eubacterium]_ SP413 Peptostreptococcaceae [XI][G-6] [Eubacteriu SP461 Peptostreptococcaceae_[XI][G-4] bacterium_HM SP376 Peptostreptococcaceae_[XI][G-4] bacterium_HM SP415 Peptostreptococcaceae [XI][G-2] bacterium HMT SP22 Mogibacterium timidum (0.07%|0.21%|0.13%|0.2 SP193 Mogibacterium vescum (0.03%|0.03%|0.04%|0 SP263 Mogibacterium neglectum (0.09%|0.08%|0.11% SP287 Treponema sp._HMT_234 (0%|0.03%|0.02%|0.05%| SP72 Treponema sp._HMT_230 (0.01%|0.03%|0.01%|0.04% SP165 Treponema sp._HMT_238 (0.03%|0.06%|0.02%|0.0 SP4 Treponema sp._HMT_237 (0.18%|0.10%|0.15%|0.11% SP73 Treponema medium (0.04%|0.03%|0.04%|0.02%|0% SP203 Treponema vincentii (0.02%|0.02%|0.05%|0.03%|0 SP74 Treponema sp._HMT_231 (0.14%|0.05%|0.16%|0.06 SP30 Treponema sp._HMT_951 (0.03%|0.01%|0.03%|0 SP270 Treponema sp. HMT 247 (0.03%|0.03%|0.06%|0.0 SP2 Treponema denticola (0.14%|0.10%|0.15%|0.09%|0 SP258 Treponema sp._HMT_258 (0.00%|0.01%|0%|0 SP122 Treponema maltophilum (0.11%|0.09%|0.09% SP320 Treponema lecithinolyticum (0.11%|0.05%|0 SP352 Treponema sp._HMT_270 (0.02%|0.03%|0% SP143 Treponema sp. HMT 268 (0.06% | 0.01% SP60 Treponema socranskii (0.34%|0.47%|0.3 SP313 Treponema sp._HMT_263 (0.03%|0.02% SP167 Treponema sp._HMT_262 (0.04%|0.02%| SP48 Treponema sp._HMT_257 (0.08%|0.11%|0 SP94 Treponema amylovorum (0%|0.04%|0.019 SP385 Anaerolineae [G-1] bacterium HMT 439 (0%) SP33 Fretibacterium fastidiosum (0.15% | 0.22% | 0.17% | 0.0 SP170 Fretibacterium sp._HMT_359 (0.01%|0.11%|0.05%| SP102 Fretibacterium sp._HMT_362 (0.02%|0.11%|0%|0.0 SP58 Fretibacterium sp._HMT_360 (0.14%|0.23%|0.13%|0 SP10 Slackia exigua (0.14% 0.14% 0.05% 0.11% 0. SP338 Cryptobacterium curtum (0.03%|0.10%|0.06 SP29 Olsenella profusa (0%|0%|0%|0%|0%|0%|0%|0.1 SP118 Olsenella uli (0.06% 0.36% 0.08% 0.11% 0.10% SP108 Olsenella sp._HMT_807 (1.19%|1.10%|0.53%|1 SP8 Atopobium sp._HMT_416 (0%|0%|0%|0%|0%|0% SP353 Lancefieldella parvula (0.25%|0.32%|0.62%|0 SP269 Atopobium sp. HMT 199 (0.02%|0.02%|0%| SP227 Lancefieldella rimae (0.39%|0.50%|0.50%|0 SP401 Bifidobacterium dentium (0.17%|0.37%|1.15 SP475 Bifidobacterium longum (0.03%|0.00%|0.03% SP277 Scardovia wiggsiae (0.34%|0.63%|0.45%|0 SP436 Parascardovia denticolens (0.03%|0%|0%|0. SP159 Propionibacterium acidifaciens (0.04% | 0.12% | SP497 Arachnia rubra (0.03%|0%|0.05%|0%|0%|0.07 SP182 Arachnia propionica (0.70%|0.22%|0.27%|0.17 SP31 Corynebacterium matruchotii (3.54%|2.05%|1 SP241 Corynebacterium durum (0.30%|0.18%|0.27 SP161 Rothia mucilaginosa (0.59%|0.49%|0.42%|0.4 SP147 Rothia dentocariosa (14.74%|14.67%|15.73% SP311 Rothia aeria (0.31%|0.54%|0.31%|0.52%|0.4 SP146 Peptidiphaga sp._HMT_183 (1.09%|1.38%|1.2 SP264 Peptidiphaga gingivicola (0.36%|0.30%|0.16 SP232 Schaalia sp._HMT_178 (0.13%|0.17%|0.17 SP350 Schaalia georgiae (0.05% 0.02% 0.08% 0.0 SP173 Schaalia meyeri (0.14%|0.49%|0.11%|0.14 SP178 Schaalia cardiffensis (0.03%|0.01%|0%|0 SP226 Schaalia lingnae_[Not_Validly_Published] SP257 Schaalia odontolyticus (0.03%|0.01%|0.0 SP266 Schaalia sp._HMT_180 (0.34%|0.40%|0.4 SP13 Schaalia odontolytica (0.14%|0.19%|0.26% 0 00000(SP230 Actinomyces sp. HMT 448 (3.85%|5.96% SP126 Actinomyces israelii (0.58%|0.79%|0.20% SP19 Actinomyces massiliensis (0.33%|0.30%|0 SP5 Actinomyces sp._HMT_897 (0.25%|0.40%|0 SP186 Actinomyces sp._HMT_414 (0.08%|0.27 SP69 Actinomyces dentalis (1.00%|0.64%|0.8 SP361 Actinomyces sp._HMT_896 (0.03%|0.1. SP91 Actinomyces sp._HMT_525 (0.23%|0.76 SP319 Actinomyces timonensis (0.03%|0.08%|0.0 SP351 Actinomyces johnsonii (0.25%|0.10%|0.10% SP129 Actinomyces sp._HMT_170 (0.44%|0.20%|0 SP93 Actinomyces sp._HMT_171 (0.97%|0.59%|0. SP372 Actinomyces viscosus (0.02%|0.13%|0.09 SP244 Actinomyces naeslundii (0.62%|0.24%|0.4 0 SP9 Actinomyces sp._HMT_169 (1.32%|2.78%|2.8 SP85 Actinomyces oris (0.57%|1.23%|1.27%|0.99 0 SP77 Actinomyces sp._HMT_175 (0.43%|0.89%|0 SP189 Desulfobulbus sp._HMT_041 (0.02%|0.05%|0.05% SP84 Campylobacter concisus (0.14%|0.20%|0.18% SP134 Campylobacter showae (0.12%|0.05%|0.1 SP390 Campylobacter rectus (0.03%|0.02%|0.02% SP185 Campylobacter gracilis (0.88% | 0.44% | 0.5 SP23 Campylobacter sp._HMT_044 (0.02%|0.05% SP145 Lautropia mirabilis (0.43%|0.28%|0.49%|0.75%| SP279 Ottowia sp._HMT_894 (0.01%|0%|0%|0.04% SP299 Kingella oralis (0.51%|0.35%|0.33%|0.50%|0.82% SP38 Neisseria bacilliformis (0.07%|0.01%|0.02%|0.07% 0 SP359 Neisseria oralis (0.15%|0%|0.11%|0%|0%|0.05 SP205 Neisseria elongata (0.07%|0.09%|0.06%|0.27% SP109 Eikenella corrodens (0.08%|0.11%|0.13%|0.14 SP297 Kingella denitrificans (0.01%|0.04%|0%|0.01% SP210 Neisseria flavescens (0.17%|0%|0.05%|0.02%|0 SP243 Neisseria perflava (0.13%|0.00%|0.22%|0.01%| SP278 Neisseria subflava (0.00%|0.02%|0.00%|0.00%| SP67 Neisseria sicca (0.77% | 0.30% | 0.28% | 0.42% | 0.33 SP98 Neisseria mucosa (0.05% 0.21% 0.16% 0.15% 0 SP54 Cardiobacterium hominis (0.12%|0.08%|0.14%|0.15 SP261 Cardiobacterium valvarum (0.08%|0.04%|0.03%|0. SP286 Haemophilus parahaemolyticus (0.01%|0.01% 0 SP81 Haemophilus parainfluenzae (2.12%|0.91%|1. О SP274 Haemophilus sp._HMT_036 (0.03%|0.02%|0.0 SP174 Haemophilus haemolyticus (0.06% | 0.03% | 0.0 SP234 Aggregatibacter sp._HMT_458 (0.15%|0.16%|0 SP382 Aggregatibacter sp._HMT_512 (0.01%|0.07%| SP322 Aggregatibacter segnis (0.03%|0.02%|0.30% SP43 Aggregatibacter sp._HMT_513 (0.04%|0.14%|0 SP284 Aggregatibacter sp._HMT_898 (0.04%|0%|0. SP317 Aggregatibacter sp._HMT_949 (0.03%|0.03% SP61 Aggregatibacter aphrophilus (0.08%|0.06%|0 SP40 Saccharibacteria_(TM7)_[G-5] bacterium_HM⁻ SP298 Saccharibacteria_(TM7)_[G-6] bacterium_HM SP201 Saccharibacteria_(TM7)_[G-3] bacteriur SP116 Saccharibacteria_(TM7)_[G-2] bacterium_HM SP16 Saccharibacteria_(TM7)_[G-1] bacterium_HM



SP111 Prevotella veroralis (0.09%|0.1

Baseline.NA.Plaque 1

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