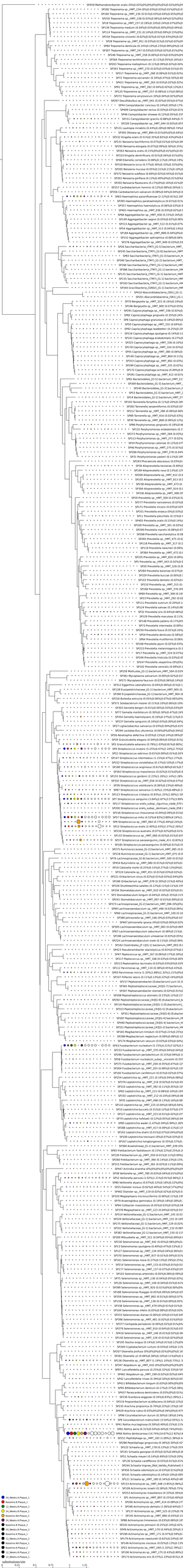
SP459 Methanobrevibacter oralis (0%)
SP182 Treponema sp._HMT_234 (0%)
SP183 Treponema sp._HMT_237 (0%)
SP184 Treponema sp._HMT_238 (0.03%)
SP18 Treponema sp._HMT_237 (0.18%)
SP139 Treponema medium (0.04%)
SP140 Treponema sp._HMT_231 (0.14%)
SP234 Treponema vincentii (0.02%)
SP235 Treponema sp._HMT_351 (0.03%)
SP84 Treponema denticola (0.14%)
SP85 Treponema sp._HMT_242 (0.04%)
SP246 Treponema sp._HMT_258 (0.05%)
SP364 Treponema lecithinolyticum (0.11%)
SP233 Treponema maltophilum (1.41%)
SP343 Treponema sp._HMT_270 (0.02%)
SP117 Treponema sp._HMT_508 (0.06%)
SP72 Treponema socranskii (0.34%)
SP421 Treponema sp._HMT_350 (0.03%)
SP61 Treponema sp._HMT_262 (0.04%)
SP272 Treponema amylovorum (0.04%)
SP267 Desulfobulbus sp._HMT_041 (0.02%)
SP44 Campylobacter concisus (0.14%)
SP49 Campylobacter rectus (0.07%)
SP46 Campylobacter showae (0.12%)
SP45 Campylobacter gracilis (0.49%)
SP128 Campylobacter sp._HMT_044 (0.02%)
SP133 Lautropia mirabilis (0.43%)
SP382 Ottowia sp._HMT_894 (0.01%)
SP232 Kingella oralis (0.31%)
SP322 Neisseria bacilliformis (0.02%)
SP295 Neisseria elongata (0.07%)
SP96 Neisseria oralis (0.18%)
SP357 Haemophilus haemolyticus (0.06%)
SP401 Haemophilus_D_138 (0.03%)
SP68 Aggregatibacter sp._HMT_458 (0.15%)
SP439 Aggregatibacter sp._HMT_513 (0.02%)
SP313 Aggregatibacter sp._HMT_512 (0.01%)
SP54 Aggregatibacter sp._HMT_513 (0.04%)
SP184 Aggregatibacter sp._HMT_898 (0.04%)
SP308 Aggregatibacter sp._HMT_898 (0.08%)
SP278 Aggregatibacter sp._HMT_949 (0.03%)
SP16 Saccharibacteria_(TM7)_[G-1]_sp._HMT_
SP245 Saccharibacteria_(TM7)_[G-4]_bacterium_HMT
SP3 Saccharibacteria_(TM7)_[G-3]_bacterium_H
SP340 Saccharibacteria_(TM7)_[G-2]_bacterium_HMT
SP166 Saccharibacteria_(TM7)_[G-1]_bacterium_HM
SP344 Saccharibacteria_(TM7)_[G-5]_bacterium_HM
SP125 Saccharibacteria_(TM7)_[G-1]_bacterium_HM
SP135 Saccharibacteria_(TM7)_[G-3]_bacterium_HM
SP183 Saccharibacteria_(TM7)_[G-1]_bacterium_HM
SP144 Actinobacillus_(GNB2)_[G-1]_bacterium_HMT
SP410 Abiotrophia_defectiva_(SFA1)_[G-1]
SP21 Abiotrophia_defectiva_(SFA1)_[G-1]
SP70 Bergeyella sp._HMT_322 (0.16%)
SP246 Bergeyella sp._HMT_365 (0.07%)
SP291 Capnocytophaga sp._HMT_338 (0.02%)
SP62 Capnocytophaga gingivalis (0.35%)
SP6 Capnocytophaga sp._HMT_332 (0.04%)
SP50 Capnocytophaga sp._HMT_332 (0.04%)
SP227 Capnocytophaga leadbetteri (0.72%)
SP216 Capnocytophaga ochracea (0.14%)
SP101 Capnocytophaga endodontalis (0.27%)
SP225 Capnocytophaga sp._HMT_336 (0.10%)
SP178 Capnocytophaga sp._HMT_863 (0.02%)
SP45 Capnocytophaga sp._HMT_380 (0.06%)
SP140 Capnocytophaga sp._HMT_412 (0.11%)
SP153 Capnocytophaga sp._HMT_902 (0.03%)
SP299 Capnocytophaga sp._HMT_903 (0.02%)
SP172 Capnocytophaga ochracea (0.09%)
SP311 Capnocytophaga sp._HMT_412 (0.01%)
SP309 Bacteroidales_[G-5]_bacterium_HMT_511
SP240 Bacteroidales_[G-1]_bacterium_HMT_
SP53 Bacteroidales_[G-1]_bacterium_HMT_
SP14 Bacteroidales_[G-2]_bacterium_HMT_
SP333 Tannerella forsythia (0.11%)
SP283 Tannerella sp._HMT_808 (0.01%)
SP312 Tannerella sp._HMT_286 (0.06%)
SP89 Tannerella sp._HMT_916 (0.02%)
SP50 Tannerella sp._HMT_808 (0.04%)
SP86 Porphyromonas endodontalis (0.79%)
SP132 Porphyromonas endodontalis (0.3%)
SP13 Porphyromonas sp._HMT_284 (0.03%)
SP113 Porphyromonas sp._HMT_277 (0.02%)
SP24 Porphyromonas catoniae (2.02%)
SP40 Porphyromonas sp._HMT_275 (0.01%)
SP166 Porphyromonas sp._HMT_278 (0.04%)
SP31 Porphyromonas pasteri (0.13%)
SP83 Phocaeicola abscessus (0.03%)
SP58 Alloprevotella tannerae (0.08%)
SP199 Alloprevotella rava (0.13%)
SP309 Alloprevotella sp._HMT_912 (0.01%)
SP105 Alloprevotella sp._HMT_913 (0.01%)
SP238 Alloprevotella sp._HMT_473 (0.
SP394 Alloprevotella sp._HMT_914 (0.1
SP38 Alloprevotella sp._HMT_308 (0%)
SP59 Prevotella sp._HMT_304 (0.03%)
SP277 Prevotella nanceiensis (0.02%)
SP171 Prevotella micans (0.03%)
SP331 Prevotella enoeca (0%)
SP111 Prevotella pleuritidis (0.15%)
SP405 Prevotella oralis (0.15%)
SP158 Prevotella buccae (0.05%)
SP200 Prevotella marshii (0.48%)
SP258 Prevotella saccharolytica (0.08
SP265 Prevotella sp._HMT_475 (0.
SP418 Prevotella sp._HMT_317 (0.2
SP119 Prevotella loescheii (0.09%)
SP384 Prevotella sp._HMT_472 (0.
SP335 Prevotella sp._HMT_820 (0.06%)
SP5 Prevotella sp._HMT_843 sp._HMT_
SP20 Prevotella sp._HMT_526 (0.0
SP284 Prevotella baroniae (0.17%)
SP329 Prevotella buccalis (0.05%)
SP143 Prevotella dentalis (0.02%)
SP219 Prevotella sp._HMT_313 (0.
SP168 Prevotella sp._HMT_376 (0%
SP63 Prevotella sp._HMT_300 (0.18
SP23 Prevotella sp._HMT_292 (0.02
SP111 Prevotella oulorum (0.29%)
SP124 Prevotella salivae (0.14%)
SP125 Prevotella oris (0.44%)
SP129 Prevotella maculosa (0.1%)
SP148 Prevotella pallens (0.17%)
SP73 Prevotella intermedia (0.09%)
SP159 Prevotella fusca (0.17%)
SP59 Prevotella denticola (0.58%)
SP64 Prevotella multiformis (0.06%
SP108 Prevotella jejuni (0.83%)
SP223 Prevotella melaninogenica (0.1
SP12 Prevotella sp._HMT_314 (0.03%)
SP249 Prevotella histicola (0.03%)
SP167 Prevotella veroralis (0.09%)
SP302 Prevotella sp._HMT_306 (0.03%)
SP381 Myroides odoratimimus (0.05%)
SP371 Myroides odoratus (0.05%)
SP152 Eggerthia catenaformis (0.04%)
SP128 Fusobacterium_[G-1]_bacterium_HMT_
SP386 Erysipelotrichaceae_[G-1]_bacterium_HMT_
SP416 Bulleidia extructa (0.05%)
SP371 Solobacterium moorei (0.52%)
SP203 Gemella bergeri (0.01%)
SP377 Gemella haemolysans (0.01%)
SP154 Gemella haemolysans (0.19%)
SP237 Gemella sanguinis (0.16%)
SP27 Lgalactibacillus salivarius (0.03%)
SP194 Lactobacillus ultunensis (0.04%)
SP85 Absiella dolichum (0.05%)
SP214 Lachnoanaerobaculum orale (0.
SP2 Granulicatella adiacens (0.79%)
SP4 Streptococcus_cristatus_(1.25%)
SP15 Streptococcus sobrinus (0.02%)
SP13 Streptococcus intermedius (1.13%)
SP222 Streptococcus constellatus (0.17%)
SP38 Streptococcus anginosus (0.63%)
SP383 Streptococcus_mitis_oralis_pneumoniae (
SP155 Streptococcus gordonii (2.23%)
SP310 Streptococcus sanguinis (0.18%)
SP35 Streptococcus vestibularis (0.04%)
SP17 Streptococcus salivarius (1.42%)
SP123 Streptococcus cristatus (0.83%)
SP1 Streptococcus_oligofermentans_(0.
SP17 Streptococcus oralis_subsp._tigurinus_(side_770
SP208 Streptococcus oralis_subsp._dentisani_(side_
SP156 Streptococcus_chosuense (0.04%)
SP52 Streptococcus mitis_b_(SK95) (0.
SP9 Streptococcus_sp._HMT_064 (0.77%)
SP10 Streptococcus oralis (2.45%)
SP26 Streptococcus_australis_(SK1059) (0.
SP28 Streptococcus_parasanguinis_(FW213) (0.
SP133 Streptococcus sp._HMT_066 (0.02%)
SP57 Streptococcus parasanguinis_(SK41) (1.
SP185 Streptococcus parasanguinis (0.00%)
SP275 Actinomyces_sp._ID_(2_factors)_HMT_
SP214 Lachnoanaerobaculum orale (0.11%)
SP241 Enterobacter_sp._[G-1]_bacterium_HMT_
SP422 Pseudoramibacter alactolyticus (0.02%)
SP1 Streptococcus_sp._HMT_058 (0.02%)
SP213 Peptostreptococcaceae_[XI][G-1]_bacterium
SP37 Peptostreptococcaceae_[XI][G-4]_bacterium_
SP440 Peptostreptococcaceae_[XI][G-4]_bacterium_
SP21 Peptostreptococcaceae_[XI][G-1]_bacterium_HM
SP165 Mogibacterium timidum (0.07%)
SP268 Mogibacterium neglectum (0.09%)
SP174 Mogibacterium vescum (0.07%)
SP63 Fusobacterium nucleatum_(TS2) (0.72%)
SP233 Fusobacterium sp._TS2_(370) (0.04%)
SP58 Fusobacterium nucleatum_nucleatum_(ATCC
SP359 Fusobacterium nucleatum_subsp._vincentii_(
SP253 Fusobacterium sp._HMT_204 (0.03%)
SP306 Fusobacterium sp._TS1_203 (0.08%)
SP308 Fusobacterium sp._HMT_420 (0.07%)
SP259 Leptotrichia sp._HMT_221 (0.10%)
SP78 Leptotrichia sp._HMT_212 (0.03%)
SP110 Leptotrichia sp._HMT_392 (0.11%)
SP130 Leptotrichia sp._HMT_215 (0.08%)
SP161 Leptotrichia sp._HMT_212 (0.05%)
SP51 Leptotrichia sp._HMT_498 (0.13%)
SP187 Leptotrichia sp._HMT_879 (0.01%)
SP210 Leptotrichia buccalis (0.31%)
SP215 Leptotrichia sp._HMT_223 (0.03%)
SP74 Leptotrichia hofstadii (0.12%)
SP81 Leptotrichia wadei (0.17%)
SP288 Leptotrichia sp._HMT_417 (0.09%)
SP313 Leptotrichia shahii (0.81%)
SP310 Leptotrichia trevisanii (0.07%)
SP317 Leptotrichia hongkongensis (0.
SP360 Anaerolineae_[G-1]_bacterium_HMT_439
SP65 Prebacterium helcogenes (0.13%)
SP239 Prebacterium sp._HMT_473 (0.01%)
SP350 Finegoldia magna (0.01%)
SP447 Actinomyces israelii (0.02%)
SP180 Actinomyces_sp._HMT_848 (0.01%)
SP252 Veillonella parvula (1.02%)
SP80 Veillonella atypica (0.07%)
SP53 Dialister invisus (0.82%)
SP442 Dialister_sp._HMT_119 (0.02%)
SP220 Megasphaera micronuciformis (0.06%)
SP329 Megasphaera genomosp._type_1_(phylotype_
SP66 Selenomonas sputigena (0.05%)
SP137 Megasphaera sp._HMT_123 (0.04%)
SP214 Selenomonas_[G-1]_bacterium_HMT_145
SP229 Veillonellaceae_[G-1]_bacterium_HMT_155
SP175 Veillonellaceae_[G-1]_bacterium_HMT_129
SP252 Veillonellaceae_[G-1]_bacterium_HMT_132
SP380 Veillonellaceae_[G-1]_bacterium_HMT_156
SP266 Mitsuokella sp._HMT_521
SP436 Selenomonas sp._HMT_442
SP311 Selenomonas sputigena (0.02%)
SP127 Selenomonas sp._HMT_134
SP339 Selenomonas sp._HMT_937
SP136 Selenomonas noxia (0.27%)
SP214 Selenomonas sp._HMT_133
SP167 Selenomonas sp._HMT_137
SP103 Selenomonas artemidis (0.05%)
SP15 Selenomonas sp._HMT_134 (0.04%)
SP235 Selenomonas sp._HMT_149 (0.04%)
SP266 Selenomonas sp._HMT_922 (0.01%)
SP188 Selenomonas flueggei (0.05%)
SP150 Selenomonas sp._HMT_892 (0.01%)
SP158 Selenomonas sp._HMT_478 (0.01%)
SP138 Selenomonas sp._HMT_479 (0.01%)
SP279 Selenomonas infelix (0.05%)
SP255 Selenomonas dianae (0.06%)
SP219 Selenomonas sp._HMT_150 (0.01%)
SP377 Centipeda periodontii (0.05%)
SP379 Selenomonas noxia (0.01%)
SP416 Selenomonas sp._HMT_146 (0.01%)
SP158 Selenomonas sp._HMT_149 (0.01%)
SP195 Slackia exigua (0.14%)
SP349 Cryptobacterium curtum (0.03%)
SP261 Olsenella profusa (0.01%)
SP281 Olsenella sp. (0.06%)
SP194 Olsenella sp._HMT_811 (1.19%)
SP347 Atopobium sp._HMT_416 (0.02%)
SP107 Lancefieldella parvula (0.25%)
SP481 Atopobium sp._HMT_199 (0.02%)
SP422 Lancefieldella rimosa (0.59%)
SP411 Bifidobacterium dentium (0.18%)
SP373 Parascardovia denticolens (0.03%)
SP236 Scardovia wiggsiae (0.34%)
SP219 Propionibacterium acidifaciens (0.04%)
SP225 Arachnia propionica (0.79%)
SP429 Arachnia rubra (0.03%)
SP66 Corynebacterium durum (0.30%)
SP4 Corynebacterium matruchotii (1.08%)
SP41 Rothia mucilaginosa (0.59%)
SP61 Rothia aeria (0.13%)
SP68 Rothia dentocariosa (14.74%)
SP415 Peptidiphaga_HMT_183_(PM2_HMT_183)
SP286 Peptidiphaga gingivicola (0.36%)
SP29 Actinomyces sp._HMT_178 (0.13%)
SP331 Schaalia georgiae (0.02%)
SP191 Schaalia meyeri (0.14%)
SP130 Schaalia cardiffensis (0.03%)
SP414 Schaalia_(Integr._HMT_180) (0.
SP456 Schaalia odontolytica (0.03%)
SP243 Schaalia odontolytica (0.14%)
SP251 Schaalia sp._HMT_180 (0.03%)
SP153 Actinomyces sp._HMT_448 (3.85%)
SP169 Actinomyces israelii (0.58%)
SP253 Actinomyces massiliensis (0.37%)
SP1 Actinomyces sp._HMT_897
SP206 Actinomyces sp._HMT_414 (0.08%)
SP366 Actinomyces dentalis (1.05%)
SP137 Actinomyces sp._HMT_525 (0.23%)
SP181 Actinomyces sp._HMT_896 (0.02%)
SP388 Actinomyces timonensis (0.03%)
SP242 Actinomyces johnsonii (0.03%)
SP69 Actinomyces sp._HMT_170 (0.44%)
SP139 Actinomyces sp._HMT_171 (0.97%)
SP146 Actinomyces viscosus (0.02%)
SP212 Actinomyces sp._HMT_169 (0.03%)
SP190 Actinomyces sp._HMT_175 (0.45%)
SP173 Actinomyces oris (0.57%)
Contributions/site
0.25
0.5
0.75
1.25
1K Weeks_A Plaque_1
Baseline A Plaque_1
1K Weeks_B Plaque_1
Baseline B Plaque_1
1K Weeks_A Plaque_2
Baseline A Plaque_2
1K Weeks_B Plaque_2
Baseline B Plaque_2
1K Weeks_A Plaque_3
Baseline A Plaque_3
1K Weeks_B Plaque_3
Baseline B Plaque_3