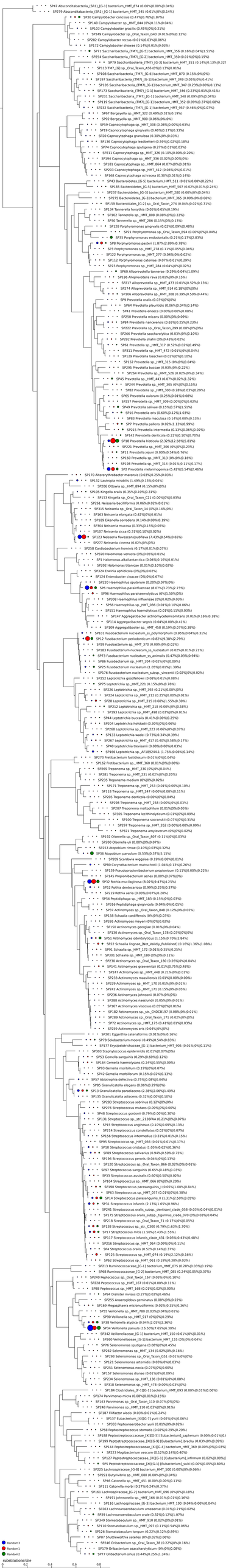
SP47 Abscondibacteria_(SR1)_[G-1]_bacterium_HMT_874 (0.00%|0.00%|0.04%)
SP29 Abscondibacteria_(SR1)_[G-1]_bacterium_HMT_345 (0.01%|0%|0.14%)
SP30 Campylobacter concisus (0.47%|0.76%|1.87%)
SP140 Campylobacter sp._HMT_044 (0%|0.01%|0.04%)
SP103 Campylobacter gracilis (0.45%|0%|0.21%)
SP249 Campylobacter sp._Oral_Taxon_043 (0.01%|0%|0.12%)
SP282 Campylobacter rectus (0.01%|0|0.06%)
SP172 Campylobacter showae (0.14%|0.01%|0.03%)
SP71 Saccharibacteria_(TM7)_[G-1]_bacterium_HMT_356 (0.16%|0.04%|1.51%)
SP254 Saccharibacteria_(TM7)_[G-1]_bacterium_HMT_350 (0.01%|0.01%|0.19%)
SP79 Saccharibacteria_(TM7)_[G-1]_bacterium_HMT_351 (0.14%|0.13%|0.32%)
SP113 TM7_[G-1]_sp._Oral_Taxon_356 (0%|0.13%|0.01%)
SP108 Saccharibacteria_(TM7)_[G-1]_bacterium_HMT_870 (0.15%|0%|0%)
SP197 Saccharibacteria_(TM7)_[G-1]_bacterium_HMT_349 (0.05%|0%|0.41%)
SP105 Saccharibacteria_(TM7)_[G-1]_bacterium_HMT_347 (0.23%|0.09%|0.13%)
SP173 Saccharibacteria_(TM7)_[G-1]_bacterium_HMT_346 (0.23%|0.01%|0.41%)
SP231 Saccharibacteria_(TM7)_[G-1]_bacterium_HMT_348 (0.09%|0%|0.04%)
SP119 Saccharibacteria_(TM7)_[G-1]_bacterium_HMT_352 (0.99%|0.37%|0.68%)
SP232 Saccharibacteria_(TM7)_[G-1]_bacterium_HMT_957 (0.46%|0%|0.07%)
SP67 Bergeyella sp._HMT_322 (0.49%|0.31%|0.19%)
SP97 Bergeyella sp._HMT_900 (0.00%|0%|0%)
SP59 Capnocytophaga sp._HMT_338 (0.08%|0.00%|0.03%)
SP19 Capnocytophaga gingivalis (0.46%|0.17%|0.33%)
SP20 Capnocytophaga granulosa (0.30%|0%|0.03%)
SP136 Capnocytophaga leadbetteri (0.59%|0.02%|0.18%)
SP74 Capnocytophaga sputigena (0.27%|0.01%|0.03%)
SP211 Capnocytophaga sp._HMT_326 (0.10%|0.00%|0.20%)
SP194 Capnocytophaga sp._HMT_336 (0.02%|0.00%|0%)
SP181 Capnocytophaga sp._HMT_864 (0.07%|0%|0.01%)
SP203 Capnocytophaga sp._HMT_412 (0.04%|0%|0.01%)
SP168 Capnocytophaga ochracea (0.30%|0.01%|0.14%)
SP43 Bacteroidetes_[G-5]_bacterium_HMT_511 (0.01%|0.00%|0.22%)
SP185 Bacteroidetes_[G-5]_bacterium_HMT_507 (0.02%|0.01%|0.24%)
SP237 Bacteroidetes_[G-3]_bacterium_HMT_280 (0.00%|0.04%|0.04%)
SP275 Bacteroidetes_[G-3]_bacterium_HMT_365 (0.00%|0%|0.06%)
SP159 Bacteroidetes_[G-2]_sp._Oral_Taxon_274 (0.04%|0.02%|0.31%)
SP134 Tannerella forsythia (0.05%|0.05%|0.19%)
SP102 Tannerella sp._HMT_808 (0.08%|0%|0.33%)
SP50 Tannerella sp._HMT_286 (0.13%|0%|0.13%)
SP128 Porphyromonas gingivalis (0.02%|0.09%|0.48%)
SP21 Porphyromonas sp._Oral_Taxon_894 (0.00%|0%|0.04%)
SP35 Porphyromonas endodontalis (0.21%|0.11%|2.83%)
SP8 Porphyromonas pasteri (1.87%|2.89%|7.78%)
SP3 Porphyromonas sp._HMT_278 (0.11%|0.05%|0.04%)
SP122 Porphyromonas sp._HMT_277 (0.04%|0%|0.02%)
SP112 Porphyromonas catoniae (0.07%|0.01%|0.29%)
SP23 Porphyromonas sp._HMT_284 (0.04%|0%|0.02%)
SP60 Alloprevotella tannerae (0.29%|0.04%|1.09%)
SP186 Alloprevotella rava (0.01%|0%|0.15%)
SP217 Alloprevotella sp._HMT_473 (0.01%|0.52%|0.13%)
SP274 Alloprevotella sp._HMT_914 (0.18%|0%|0%)
SP106 Alloprevotella sp._HMT_308 (0.39%|0.50%|0.44%)
SP9 Prevotella oralis (0.03%|0%|0%)
SP64 Prevotella oris (0.06%|0.04%|0.14%)
SP41 Prevotella enoeca (0.00%|0.00%|0.08%)
SP250 Prevotella micans (0.00%|0%|0.09%)
SP84 Prevotella nanceiensis (0.06%|0.25%|0.22%)
SP222 Prevotella sp._Oral_Taxon_299 (0.08%|0%|0%)
SP266 Prevotella saccharolytica (0.03%|0%|0.10%)
SP292 Prevotella shahii (0%|0.43%|0.02%)
SP61 Prevotella sp._HMT_317 (0.52%|0.02%|0.49%)
SP311 Prevotella sp._HMT_472 (0.01%|0%|0.04%)
SP129 Prevotella loescheii (0.02%|0%|0.10%)
SP152 Prevotella sp._HMT_315 (0.09%|0%|0%)
SP295 Prevotella buccae (0.03%|0%|0.22%)
SP64 Prevotella sp._HMT_526 (0.02%|0%|0.34%)
SP45 Prevotella sp._HMT_443 (0.07%|0.02%|1.32%)
SP244 Prevotella sp._HMT_305 (0%|0%|0.15%)
SP82 Prevotella sp._HMT_300 (0.28%|0.02%|0.29%)
SP65 Prevotella oulorum (0.25%|0.01%|0.08%)
SP257 Prevotella sp._HMT_309 (0.00%|0%|0.02%)
SP49 Prevotella salivae (0.11%|0.57%|1.51%)
SP16 Prevotella oris (0.60%|0.12%|1.03%)
SP83 Prevotella maculosa (0.14%|0.00%|0.13%)
SP7 Prevotella pallens (0.02%|1.13%|0.99%)
SP215 Prevotella intermedia (0.13%|0.06%|0.92%)
SP142 Prevotella denticola (0.22%|0.10%|0.70%)
SP18 Prevotella histicola (2.32%|12.56%|15.81%)
SP221 Prevotella sp._HMT_306 (0%|0%|0.23%)
SP11 Prevotella jejuni (0.00%|0.54%|0.76%)
SP160 Prevotella sp._HMT_313 (0%|0%|0.16%)
SP198 Prevotella sp._HMT_314 (0.01%|0.11%|0.17%)
SP52 Prevotella melaninogenica (5.42%|5.54%|2.46%)
SP170 Alteriorhabdus marensis (0.02%|0.25%|0.03%)
SP132 Lautropia mirabilis (1.49%|0.13%|0.04%)
SP206 Ottowia sp._HMT_894 (0.15%|0%|0%)
SP195 Kingella oralis (0.35%|0.19%|0.31%)
SP153 Kingella sp._Oral_Taxon_121 (0.00%|0%|0.03%)
SP261 Neisseria bacilliformis (0.06%|0.02%|0.01%)
SP155 Neisseria sp._Oral_Taxon_14 (0%|0.14%|0%)
SP163 Neisseria elongata (0.42%|0%|0.01%)
SP189 Eikenella corrodens (0.14%|0.00%|0.19%)
SP304 Neisseria mucosa (0.33%|0.15%|0.05%)
SP107 Neisseria sicca (0.31%|0.10%|0.02%)
SP123 Neisseria flavescens|subflava (7.43%|8.54%|0.65%)
SP277 Neisseria cinerea (0.02%|0%|0%)
SP258 Cardiobacterium hominis (0.17%|0.01%|0.07%)
SP320 Halomonas venusta (0%|0.05%|0.01%)
SP91 Halomonas alkalilacrifica (0.04%|0.16%|0.01%)
SP202 Halomonas titanicae (0.01%|0.10%|0.02%)
SP324 Erwinia aphidicola (0%|0%|0.02%)
SP124 Enterobacter cloacae (0%|0%|0.67%)
SP220 Haemophilus sputorum (0.20%|0.07%|0%)
SP6 Haemophilus parainfluenae (8.07%|3.73%|2.73%)
SP96 Haemophilus parahaemolyticus (0%|1.50%|0%)
SP308 Haemophilus influenzae (0%|0.02%|0.03%)
SP56 Haemophilus sp._HMT_036 (0.01%|0.10%|0.06%)
SP151 Haemophilus haemolyticus (0.01%|0.11%|0.03%)
SP147 Aggregatibacter actinomycetemcomitans (0.01%|0.16%|0.18%)
SP114 Aggregatibacter segnis (0.04%|0.00%|0.41%)
SP109 Aggregatibacter sp._HMT_458 (0.10%|0.07%|0.38%)
SP101 Fusobacterium nucleatum_ss_polymorphum (0.95%|0.04%|0.31%)
SP12 Fusobacterium periodonticum (0.82%|3.38%|2.79%)
SP29 Fusobacterium sp._HMT_370 (0.00%|0%|0.02%)
SP183 Fusobacterium nucleatum_ss_nucleatum (0.02%|0.01%|0.21%)
SP73 Fusobacterium nucleatum_ss_animalis (0.47%|0.03%|0.94%)
SP86 Fusobacterium sp._HMT_204 (0.02%|0%|0.09%)
SP25 Fusobacterium nucleatum (1.05%|0.01%|1.39%)
SP176 Fusobacterium nucleatum_subsp._vincentii (0.02%|0%|0.02%)
SP252 Leptotrichia goodfellowii (0.08%|0.01%|0.08%)
SP75 Leptotrichia sp._HMT_221 (0.15%|0%|0.76%)
SP226 Leptotrichia sp._HMT_392 (0.01%|0%|0%)
SP224 Leptotrichia sp._HMT_212 (0.25%|0.00%|0.01%)
SP28 Leptotrichia sp._HMT_215 (0.60%|1.55%|0.30%)
SP212 Leptotrichia sp._HMT_218 (0.00%|0%|0.58%)
SP193 Leptotrichia sp._HMT_498 (0.03%|0%|0.01%)
SP44 Leptotrichia buccalis (0.41%|0.00%|0.25%)
SP204 Leptotrichia hofstadii (0.39%|0%|0.06%)
SP268 Leptotrichia sp._HMT_223 (0.06%|0%|0.07%)
SP133 Leptotrichia wadei (0.73%|0.34%|0.39%)
SP267 Leptotrichia sp._HMT_417 (0.84%|0.58%|0.17%)
SP40 Leptotrichia trevisanii (0.08%|0.00%|0.03%)
SP166 Leptotrichia sp._AF189244.1 (1.75%|0.06%|0.14%)
SP233 Filifactor fastidiosum (0.01%|0%|0.04%)
SP162 Fretibacterium sp._HMT_360 (0.01%|0%|0.08%)
SP269 Treponema sp._HMT_230 (0%|0%|0.04%)
SP281 Treponema sp._HMT_231 (0.02%|0%|0.20%)
SP235 Treponema medium (0%|0%|0.02%)
SP171 Treponema sp._HMT_253 (0.01%|0.00%|0.10%)
SP118 Treponema sp._HMT_247 (0.00%|0.00%|0.11%)
SP205 Treponema denticola (0.00%|0%|0.04%)
SP268 Treponema sp._HMT_258 (0.00%|0%|0.03%)
SP297 Treponema maltophilum (0.01%|0%|0.05%)
SP305 Treponema lecithinolyticum (0.01%|0%|0.09%)
SP100 Treponema socranskii (0.07%|0%|0.31%)
SP297 Treponema sp._HMT_262 (0.00%|0.00%|0.09%)
SP321 Treponema amylovorum (0%|0%|0.02%)
SP192 Olsenella sp._Oral_Taxon_807 (0.11%|0%|0.03%)
SP200 Olsenella uli (0.00%|0%|0.07%)
SP253 Atopobium rimae (0.10%|0.03%|0.32%)
SP36 Atopobium parvulum (0.53%|0.37%|5.15%)
SP209 Scardovia wiggsiae (0.19%|0%|0.01%)
SP80 Corynebacterium matruchotii (1.04%|0.13%|0.26%)
SP139 Propionibacterium propionicum (0.11%|0.00%|0.22%)
SP145 Propionibacterium acnes (0.00%|0.07%|0%)
SP32 Rothia mucilaginosa (8.02%|9.47%|4.25%)
SP52 Rothia dentocariosa (0.99%|0.25%|0.37%)
SP219 Rothia aeria (0.03%|0.07%|0.20%)
SP54 Peptidhpga sp._HMT_183 (0.15%|0%|0.03%)
SP316 Peptidhpga gnavicola (0.04%|0%|0.05%)
SP37 Actinomyces sp._Oral_Taxon_848 (0.12%|0%|0.02%)
SP158 Schaalia cardiffensis (0%|0%|0.03%)
SP326 Actinomyces meyeri (0%|0%|0.02%)
SP150 Actinomyces georgiae (0.01%|0%|0.04%)
SP130 Actinomyces sp._Oral_Taxon_178 (0.03%|0%|0%)
SP51 Actinomyces odontolyticus (1.15%|0.70%|2.84%)
SP22 Schaalia longae_[Not_Validly_Published] (0.16%|1.36%|1.08%)
SP91 Schaalia sp._HMT_172 (0.01%|0.35%|0.25%)
SP301 Schaalia sp._HMT_180 (0%|0%|0.11%)
SP230 Actinomyces sp._Oral_Taxon_180 (0.26%|0.04%|0%)
SP141 Actinomyces graevenitzii (0.01%|0.75%|0.48%)
SP247 Actinomyces sp._HMT_448 (0.21%|0%|0.01%)
SP233 Actinomyces massiliensis (0.01%|0.00%|0.00%)
SP229 Actinomyces sp._HMT_170 (0.01%|0%|0.01%)
SP242 Actinomyces sp._HMT_171 (0.15%|0%|0.05%)
SP236 Actinomyces johnsonii (0%|0.02%|0%)
SP288 Actinomyces naeslundii (0.05%|0%|0.01%)
SP167 Actinomyces viscosus (0.05%|0%|0.01%)
SP182 Actinomyces sp._str._CHOC83? (0.08%|0%|0.01%)
SP289 Actinomyces sp._Oral_Taxon_171 (0.02%|0%|0%)
SP72 Actinomyces sp._HMT_175 (0.41%|0.01%|0.03%)
SP259 Actinomyces oris (0.04%|0%|0%)
SP201 Eggerthia catenaformis (0.01%|0%|0.16%)
SP78 Solobacterium moorei (0.49%|0.54%|0.83%)
SP177 Erysipelothrichaceae_[G-1]_bacterium_HMT_905 (0.01%|0%|0.01%)
SP303 Staphylococcus epidermidis (0.01%|0.07%|0%)
SP53 Gemella sanguinis (0.29%|0.60%|0.12%)
SP184 Gemella haemolysans (0.24%|0.55%|0.09%)
SP93 Gemella morbillorum (0.19%|0%|0.07%)
SP42 Gemella morbillorum (0.15%|0.02%|0.13%)
SP57 Abiotrophia defectiva (0.75%|0.08%|0.04%)
SP85 Granulicatella elegans (0.06%|0.29%|0%)
SP13 Granulicatella paradiacens (2.38%|2.06%|1.49%)
SP135 Granulicatella adiacens (0.32%|0.00%|0.10%)
SP23 Streptococcus sobrinus (0.12%|0%|0%)
SP276 Streptococcus mutans (0.09%|0%|0.00%)
SP48 Streptococcus gordonii (0.79%|0.00%|0.30%)
SP131 Streptococcus sp._str._2136FAA (0.21%|0%|0.07%)
SP15 Streptococcus anginosus (0.10%|0.09%|0.13%)
SP214 Streptococcus constellatus (0.02%|0%|0.07%)
SP156 Streptococcus intermedius (0.31%|0.01%|0.15%)
SP95 Streptococcus sp._HMT_056 (0.01%|0.01%|0.17%)
SP10 Streptococcus cristatus (1.05%|0.62%|0.36%)
SP89 Streptococcus salivarius (0.94%|0.59%|0.75%)
SP196 Streptococcus peroris (0.04%|0%|0.13%)
SP100 Streptococcus sp._Oral_Taxon_866 (0.02%|0%|0.01%)
SP97 Streptococcus sanguinis (0.65%|0.18%|0.33%)
SP33 Streptococcus australis (0.60%|0.59%|0.02%)
SP108 Streptococcus sp._HMT_666 (0%|0%|0.20%)
SP190 Streptococcus parasanguinis (1.05%|1.00%|0.84%)
SP63 Streptococcus sp._HMT_057 (0.01%|0%|0.38%)
SP14 Streptococcus sp._HMT_058 (0.01%|0%|0.01%)
SP31 Streptococcus infantis (2.13%|1.65%|0.96%)
SP241 Streptococcus oralis_subsp._dentisani_058 (0.03%|0.04%|0.01%)
SP175 Streptococcus oralis_subsp._tigurinus_clade_070 (0.03%|0.03%|0.04%)
SP218 Streptococcus sp._Oral_Taxon_71 (0.17%|0%|0.05%)
SP138 Streptococcus sp._str._HGH2 (0.79%|1.43%|1.70%)
SP17 Streptococcus mitis (1.50%|2.43%|1.55%)
SP117 Streptococcus infantis_clade_431 (0.03%|0.43%|0.48%)
SP214 Streptococcus sp._HMT_071 (0.01%|0%|0%)
SP4 Streptococcus oralis (0.52%|0.14%|0.37%)
SP125 Streptococcus sp._HMT_074 (0.19%|2.12%|0.16%)
SP62 Streptococcus sp._HMT_061 (0.13%|0.30%|0.03%)
SP213 Ruminococcaceae_[G-1]_bacterium_HMT_075 (0.28%|0.03%|0.19%)
SP68 Ruminococcaceae_[G-2]_bacterium_HMT_085 (0.24%|0.05%|0.37%)
SP240 Peptococcus sp._Oral_Taxon_167 (0.03%|0%|0.10%)
SP228 Peptococcus sp._HMT_167 (0.01%|0.00%|0.11%)
SP88 Peptococcus sp._HMT_168 (0.01%|0.02%|0.00%)
SP84 Dialister invisus (0.27%|0.02%|0.46%)
SP253 Anaerroglobus geminatus (0.08%|0%|0.22%)
SP169 Megasphaera micronuciformis (0.02%|0.35%|0.36%)
SP55 Veillonella sp._HMT_780 (0.03%|0.04%|0.01%)
SP90 Veillonella sp._HMT_917 (0%|0%|0.29%)
SP38 Veillonella atypica (0.94%|2.05%|2.36%)
SP54 Veillonella parvula (16.50%|7.65%|6.30%)
SP342 Veillonellaceae_[G-1]_bacterium_HMT_150 (0.01%|0%|0.01%)
SP260 Veillonellaceae_[G-1]_bacterium_HMT_155 (0.09%|0%|0.04%)
SP76 Selenomonas sputigena (0.08%|0%|0.45%)
SP262 Selenomonas sp._HMT_134 (0.02%|0%|0.16%)
SP293 Selenomonas sp._Oral_Taxon_151 (0.01%|0%|0%)
SP122 Selenomonas artemidis (0%|0%|0.03%)
SP251 Selenomonas noxia (0.07%|0%|0.00%)
SP157 Selenomonas dianae (0.01%|0%|0.09%)
SP324 Selenomonas sp._HMT_136 (0.01%|0%|0.08%)
SP318 Selenomonas sp._HMT_478 (0.00%|0.03%|0%)
SP184 Clostridiales_[F-1][G-1]_bacterium_HMT_093 (0.00%|0.01%|0.06%)
SP74 Parvimonas micra (0.08%|0.01%|0.15%)
SP143 Parvimonas sp._Oral_Taxon_110 (0.07%|0%|0%)
SP248 Parvimonas sp._HMT_110 (0.03%|0%|0.01%)
SP187 Filifactor alocis (0.07%|0.01%|0.24%)
SP137 Eubacterium_[X][G-7] yurii (0.02%|0%|0.06%)
SP333 Peptoanaerobacter yurii (0.01%|0%|0.02%)
SP58 Peptostreptococcus stomatis (0.02%|0.29%|0.29%)
SP189 Peptostreptococcaceae_[X][G-5] [Eubacterium]_saphenum (0.00%|0.01%|0.0
SP348 Peptostreptococcaceae_[X][G-9] [Eubacterium]_brachy (0.03%|0%|0.09%)
SP144 Peptostreptococcaceae_[X][G-4] [Eubacterium]_nodatum_HMT_369 (0.00%|0%|0.03%)
SP223 Mogibacterium vescum (0.12%|0.14%|0.40%)
SP127 Peptostreptococcaceae_[X][G-1] [Eubacterium]_infirmum (0.02%|0.00%|
SP5 Peptostreptococcaceae_[X][G-1] [Eubacterium]_sulci (0.00%|0.05%|0.89%)
SP225 Lachnospiraceae_[G-8] bacterium_HMT_500 (0.00%|0%|0.06%)
SP291 Butyrivibrio sp._HMT_080 (0.00%|0%|0.04%)
SP46 Catonella sp._HMT_451 (0.00%|0%|0.11%)
SP111 Catonella morbi (0.27%|0%|0.37%)
SP161 Lachnospiraceae_[G-2] bacterium_HMT_096 (0%|0%|0.18%)
SP191 Johnsonella sp._HMT_146 (0.01%|0.01%|0.16%)
SP116 Lachnospiraceae_[G-3] bacterium_HMT_100 (0.04%|0.00%|0.04%)
SP263 Lachnoanaerobaculum umeaense (0.01%|0.21%|0.02%)
SP139 Lachnoanaerobaculum orale (0.32%|0.12%|1.07%)
SP349 Stomatobaculum sp._HMT_910 (0.02%|0%|0.01%)
SP110 Stomatobaculum sp._HMT_097 (0.11%|0.54%|0.06%)
SP126 Stomatobaculum longum (0.22%|0.12%|0.89%)
SP87 Shuttleworthia satelles (0%|0.02%|0.06%)
SP146 Oribacterium sp._Oral_Taxon_78 (0.22%|0%|0.16%)
SP126 Oribacterium asaccharolyticum (0%|0%|0.08%)
SP77 Oribacterium sinus (0.44%|0.25%|1.34%)
Random3
Random1
Random2
Constitutions/site
0.2
0.4
0.6
0.8
1