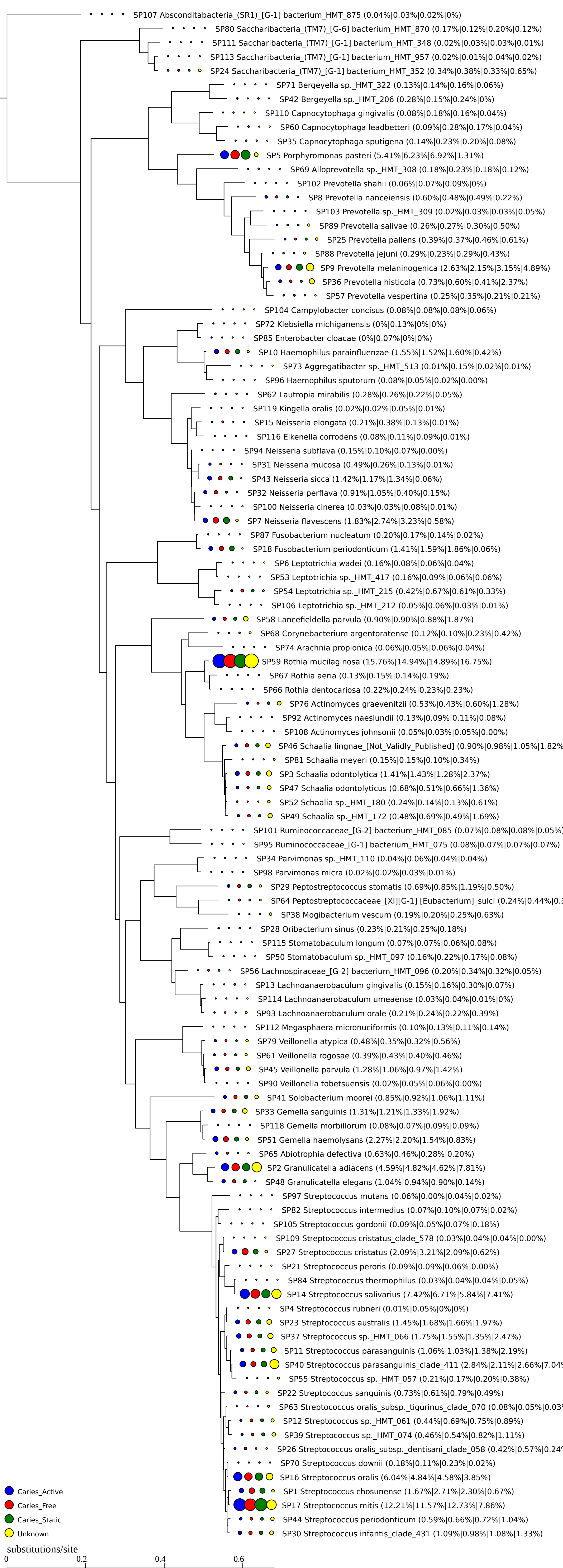
SP107 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_875 (0.04%|0.03%|0.02%|0%)
SP80 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870 (0.17%|0.12%|0.20%|0.12%)
SP111 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348 (0.02%|0.03%|0.03%|0.01%)
SP113 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957 (0.02%|0.01%|0.04%|0.02%)
SP24 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_352 (0.34%|0.38%|0.33%|0.65%)
SP71 Bergeyella sp._HMT_322 (0.13%|0.14%|0.16%|0.06%)
SP42 Bergeyella sp._HMT_206 (0.28%|0.15%|0.24%|0%)
SP110 Capnocytophaga gingivalis (0.08%|0.18%|0.16%|0.04%)
SP60 Capnocytophaga leadbetteri (0.09%|0.28%|0.17%|0.04%)
SP35 Capnocytophaga sputigena (0.14%|0.23%|0.20%|0.08%)
SP5 Porphyromonas pasteri (5.41%|6.23%|6.92%|1.31%)
SP69 Alloprevotella sp._HMT_308 (0.18%|0.23%|0.18%|0.12%)
SP102 Prevotella shahii (0.06%|0.07%|0.09%|0%)
SP8 Prevotella nanceiensis (0.60%|0.48%|0.49%|0.22%)
SP103 Prevotella sp._HMT_309 (0.02%|0.03%|0.03%|0.05%)
SP89 Prevotella salivae (0.26%|0.27%|0.30%|0.50%)
SP25 Prevotella pallens (0.39%|0.37%|0.46%|0.61%)
SP88 Prevotella jejuni (0.29%|0.23%|0.29%|0.43%)
SP9 Prevotella melaninogenica (2.63%|2.15%|3.15%|4.89%)
SP36 Prevotella histicola (0.73%|0.60%|0.41%|2.37%)
SP57 Prevotella vespertina (0.25%|0.35%|0.21%|0.21%)
SP104 Campylobacter concisus (0.08%|0.08%|0.08%|0.06%)
SP72 Klebsiella michiganensis (0%|0.13%|0%|0%)
SP85 Enterobacter cloacae (0%|0.07%|0%|0%)
SP10 Haemophilus parainfluenzae (1.55%|1.52%|1.60%|0.42%)
SP73 Aggregatibacter sp._HMT_513 (0.01%|0.15%|0.02%|0.01%)
SP96 Haemophilus sputorum (0.08%|0.05%|0.02%|0.00%)
SP62 Lautropia mirabilis (0.28%|0.26%|0.22%|0.05%)
SP119 Kingella oralis (0.02%|0.02%|0.05%|0.01%)
SP15 Neisseria elongata (0.21%|0.38%|0.13%|0.01%)
SP116 Eikenella corrodens (0.08%|0.11%|0.09%|0.01%)
SP94 Neisseria subflava (0.15%|0.10%|0.07%|0.00%)
SP31 Neisseria mucosa (0.49%|0.26%|0.13%|0.01%)
SP43 Neisseria sicca (1.42%|1.17%|1.34%|0.06%)
SP32 Neisseria perflava (0.91%|1.05%|0.40%|0.15%)
SP100 Neisseria cinerea (0.03%|0.03%|0.08%|0.01%)
SP7 Neisseria flavescens (1.83%|2.74%|3.23%|0.58%)
SP87 Fusobacterium nucleatum (0.20%|0.17%|0.14%|0.02%)
SP18 Fusobacterium periodonticum (1.41%|1.59%|1.86%|0.06%)
SP6 Leptotrichia wadei (0.16%|0.08%|0.06%|0.04%)
SP53 Leptotrichia sp._HMT_417 (0.16%|0.09%|0.06%|0.06%)
SP54 Leptotrichia sp._HMT_215 (0.42%|0.67%|0.61%|0.33%)
SP106 Leptotrichia sp._HMT_212 (0.05%|0.06%|0.03%|0.01%)
SP58 Lancefieldella parvula (0.90%|0.90%|0.88%|1.87%)
SP68 Corynebacterium argentoratense (0.12%|0.10%|0.23%|0.42%)
SP74 Arachnia propionica (0.06%|0.05%|0.06%|0.04%)
SP59 Rothia mucilaginosa (15.76%|14.94%|14.89%|16.75%)
SP67 Rothia aeria (0.13%|0.15%|0.14%|0.19%)
SP66 Rothia dentocariosa (0.22%|0.24%|0.23%|0.23%)
SP76 Actinomyces graevenitzii (0.53%|0.43%|0.60%|1.28%)
SP92 Actinomyces naeslundii (0.13%|0.09%|0.11%|0.08%)
SP108 Actinomyces johnsonii (0.05%|0.03%|0.05%|0.00%)
SP46 Schaalia lingnae_[Not_Validly_Published] (0.90%|0.98%|1.05%|1.82%
SP81 Schaalia meyeri (0.15%|0.15%|0.10%|0.34%)
SP3 Schaalia odontolytica (1.41%|1.43%|1.28%|2.37%)
SP47 Schaalia odontolyticus (0.68%|0.51%|0.66%|1.36%)
SP52 Schaalia sp._HMT_180 (0.24%|0.14%|0.13%|0.61%)
SP49 Schaalia sp._HMT_172 (0.48%|0.69%|0.49%|1.69%)
SP101 Ruminococcaceae_[G-2] bacterium_HMT_085 (0.07%|0.08%|0.08%|0.05%
SP95 Ruminococcaceae_[G-1] bacterium_HMT_075 (0.08%|0.07%|0.07%|0.07%)
SP34 Parvimonas sp._HMT_110 (0.04%|0.06%|0.04%|0.04%)
SP98 Parvimonas micra (0.02%|0.02%|0.03%|0.01%)
SP29 Peptostreptococcus stomatis (0.69%|0.85%|1.19%|0.50%)
SP64 Peptostreptococcaceae_[XI][G-1] [Eubacterium]_sulci (0.24%|0.44%|0.
SP38 Mogibacterium vescum (0.19%|0.20%|0.25%|0.63%)
SP28 Oribacterium sinus (0.23%|0.21%|0.25%|0.18%)
SP115 Stomatobaculum longum (0.07%|0.07%|0.06%|0.08%)
SP50 Stomatobaculum sp._HMT_097 (0.16%|0.22%|0.17%|0.08%)
SP56 Lachnospiraceae_[G-2] bacterium_HMT_096 (0.20%|0.34%|0.32%|0.05%)
SP13 Lachnoanaerobaculum gingivalis (0.15%|0.16%|0.30%|0.07%)
SP114 Lachnoanaerobaculum umeaense (0.03%|0.04%|0.01%|0%)
SP93 Lachnoanaerobaculum orale (0.21%|0.24%|0.22%|0.39%)
SP112 Megasphaera micronuciformis (0.10%|0.13%|0.11%|0.14%)
SP79 Veillonella atypica (0.48%|0.35%|0.32%|0.56%)
SP61 Veillonella rogosae (0.39%|0.43%|0.40%|0.46%)
SP45 Veillonella parvula (1.28%|1.06%|0.97%|1.42%)
SP90 Veillonella tobetsuensis (0.02%|0.05%|0.06%|0.00%)
SP41 Solobacterium moorei (0.85%|0.92%|1.06%|1.11%)
SP33 Gemella sanguinis (1.31%|1.21%|1.33%|1.92%)
SP118 Gemella morbillorum (0.08%|0.07%|0.09%|0.09%)
SP51 Gemella haemolysans (2.27%|2.20%|1.54%|0.83%)
SP65 Abiotrophia defectiva (0.63%|0.46%|0.28%|0.20%)
SP2 Granulicatella adiacens (4.59%|4.82%|4.62%|7.81%)
SP48 Granulicatella elegans (1.04%|0.94%|0.90%|0.14%)
SP97 Streptococcus mutans (0.06%|0.00%|0.04%|0.02%)
SP82 Streptococcus intermedius (0.07%|0.10%|0.07%|0.02%)
SP105 Streptococcus gordonii (0.09%|0.05%|0.07%|0.18%)
SP109 Streptococcus cristatus_clade_578 (0.03%|0.04%|0.04%|0.00%)
SP27 Streptococcus cristatus (2.09%|3.21%|2.09%|0.62%)
SP21 Streptococcus peroris (0.09%|0.09%|0.06%|0.00%)
SP84 Streptococcus thermophilus (0.03%|0.04%|0.04%|0.05%)
SP14 Streptococcus salivarius (7.42%|6.71%|5.84%|7.41%)
SP4 Streptococcus rubneri (0.01%|0.05%|0%|0%)
SP23 Streptococcus australis (1.45%|1.68%|1.66%|1.97%)
SP37 Streptococcus sp._HMT_066 (1.75%|1.55%|1.35%|2.47%)
SP11 Streptococcus parasanguinis (1.06%|1.03%|1.38%|2.19%)
SP40 Streptococcus parasanguinis_clade_411 (2.84%|2.11%|2.66%|7.04%
SP55 Streptococcus sp._HMT_057 (0.21%|0.17%|0.20%|0.38%)
SP22 Streptococcus sanguinis (0.73%|0.61%|0.79%|0.49%)
SP63 Streptococcus oralis_subsp._tigurinus_clade_070 (0.08%|0.05%|0.03%
SP12 Streptococcus sp._HMT_061 (0.44%|0.69%|0.75%|0.89%)
SP39 Streptococcus sp._HMT_074 (0.46%|0.54%|0.82%|1.11%)
SP26 Streptococcus oralis_subsp._dentisani_clade_058 (0.42%|0.57%|0.24%
SP70 Streptococcus downii (0.18%|0.11%|0.23%|0.02%)
SP16 Streptococcus oralis (6.04%|4.84%|4.58%|3.85%)
SP1 Streptococcus chosunense (1.67%|2.71%|2.30%|0.67%)
SP17 Streptococcus mitis (12.21%|11.57%|12.73%|7.86%)
SP44 Streptococcus periodonticum (0.59%|0.66%|0.72%|1.04%)
SP30 Streptococcus infantis_clade_431 (1.09%|0.98%|1.08%|1.33%)
Caries_Active
Caries_Free
Caries_Static
Unknown
substitutions/site
0
0.2
0.4
0.6