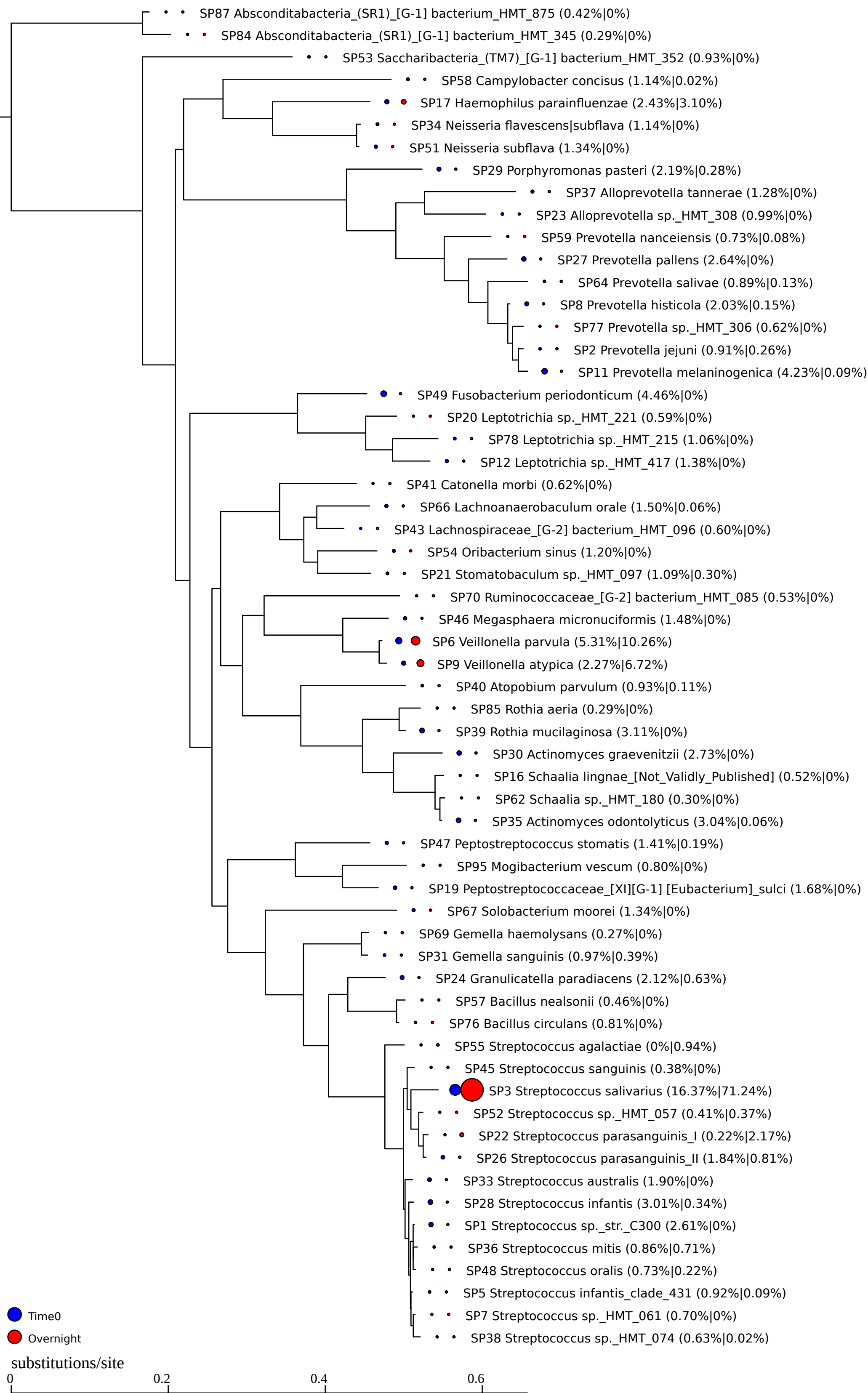
SP87 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_875 (0.42%|0%)
SP84 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345 (0.29%|0%)
SP53 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_352 (0.93%|0%)
SP58 Campylobacter concisus (1.14%|0.02%)
SP17 Haemophilus parainfluenzae (2.43%|3.10%)
SP34 Neisseria flavescens|subflava (1.14%|0%)
SP51 Neisseria subflava (1.34%|0%)
SP29 Porphyromonas pasteri (2.19%|0.28%)
SP37 Alloprevotella tannerae (1.28%|0%)
SP23 Alloprevotella sp._HMT_308 (0.99%|0%)
SP59 Prevotella nanceiensis (0.73%|0.08%)
SP27 Prevotella pallens (2.64%|0%)
SP64 Prevotella salivae (0.89%|0.13%)
SP8 Prevotella histicola (2.03%|0.15%)
SP77 Prevotella sp._HMT_306 (0.62%|0%)
SP2 Prevotella jejuni (0.91%|0.26%)
SP11 Prevotella melaninogenica (4.23%|0.09%)
SP49 Fusobacterium periodonticum (4.46%|0%)
SP20 Leptotrichia sp._HMT_221 (0.59%|0%)
SP78 Leptotrichia sp._HMT_215 (1.06%|0%)
SP12 Leptotrichia sp._HMT_417 (1.38%|0%)
SP41 Catonella morbi (0.62%|0%)
SP66 Lachnoanaerobaculum orale (1.50%|0.06%)
SP43 Lachnospiraceae_[G-2] bacterium_HMT_096 (0.60%|0%)
SP54 Oribacterium sinus (1.20%|0%)
SP21 Stomatobaculum sp._HMT_097 (1.09%|0.30%)
SP70 Ruminococcaceae_[G-2] bacterium_HMT_085 (0.53%|0%)
SP46 Megasphaera micronuciformis (1.48%|0%)
SP6 Veillonella parvula (5.31%|10.26%)
SP9 Veillonella atypica (2.27%|6.72%)
SP40 Atopobium parvulum (0.93%|0.11%)
SP85 Rothia aeria (0.29%|0%)
SP39 Rothia mucilaginosa (3.11%|0%)
SP30 Actinomyces graevenitzii (2.73%|0%)
SP16 Schaalia lingnae_[Not_Validly_Published] (0.52%|0%)
SP62 Schaalia sp._HMT_180 (0.30%|0%)
SP35 Actinomyces odontolyticus (3.04%|0.06%)
SP47 Peptostreptococcus stomatis (1.41%|0.19%)
SP95 Mogibacterium vescum (0.80%|0%)
SP19 Peptostreptococcaceae_[XI][G-1] [Eubacterium]_sulci (1.68%|0%)
SP67 Solobacterium moorei (1.34%|0%)
SP69 Gemella haemolysans (0.27%|0%)
SP31 Gemella sanguinis (0.97%|0.39%)
SP24 Granulicatella paradiacens (2.12%|0.63%)
SP57 Bacillus nealsonii (0.46%|0%)
SP76 Bacillus circulans (0.81%|0%)
SP55 Streptococcus agalactiae (0%|0.94%)
SP45 Streptococcus sanguinis (0.38%|0%)
SP3 Streptococcus salivarius (16.37%|71.24%)
SP52 Streptococcus sp._HMT_057 (0.41%|0.37%)
SP22 Streptococcus parasanguinis_I (0.22%|2.17%)
SP26 Streptococcus parasanguinis_II (1.84%|0.81%)
SP33 Streptococcus australis (1.90%|0%)
SP28 Streptococcus infantis (3.01%|0.34%)
SP1 Streptococcus sp._str._C300 (2.61%|0%)
SP36 Streptococcus mitis (0.86%|0.71%)
SP48 Streptococcus oralis (0.73%|0.22%)
SP5 Streptococcus infantis_clade_431 (0.92%|0.09%)
SP7 Streptococcus sp._HMT_061 (0.70%|0%)
SP38 Streptococcus sp._HMT_074 (0.63%|0.02%)
Time0
Overnight
substitutions/site
0
0.2
0.4
0.6