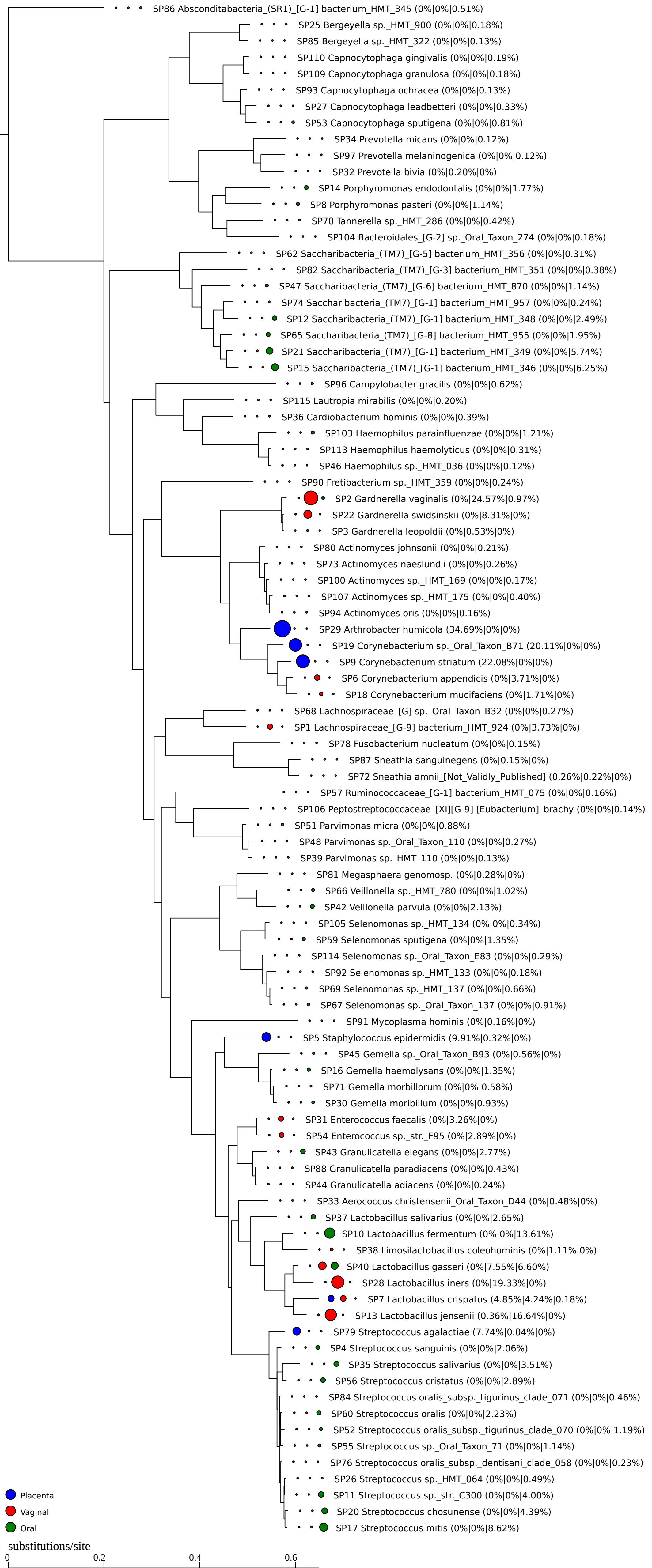
SP86 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345 (0%|0%|0.51%)
SP25 Bergeyella sp._HMT_900 (0%|0%|0.18%)
SP85 Bergeyella sp._HMT_322 (0%|0%|0.13%)
SP110 Capnocytophaga gingivalis (0%|0%|0.19%)
SP109 Capnocytophaga granulosa (0%|0%|0.18%)
SP93 Capnocytophaga ochracea (0%|0%|0.13%)
SP27 Capnocytophaga leadbetteri (0%|0%|0.33%)
SP53 Capnocytophaga sputigena (0%|0%|0.81%)
SP34 Prevotella micans (0%|0%|0.12%)
SP97 Prevotella melaninogenica (0%|0%|0.12%)
SP32 Prevotella bivia (0%|0.20%|0%)
SP14 Porphyromonas endodontalis (0%|0%|1.77%)
SP8 Porphyromonas pasteri (0%|0%|1.14%)
SP70 Tannerella sp._HMT_286 (0%|0%|0.42%)
SP104 Bacteroidales_[G-2] sp._Oral_Taxon_274 (0%|0%|0.18%)
SP62 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356 (0%|0%|0.31%)
SP82 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351 (0%|0%|0.38%)
SP47 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870 (0%|0%|1.14%)
SP74 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957 (0%|0%|0.24%)
SP12 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348 (0%|0%|2.49%)
SP65 Saccharibacteria_(TM7)_[G-8] bacterium_HMT_955 (0%|0%|1.95%)
SP21 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349 (0%|0%|5.74%)
SP15 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346 (0%|0%|6.25%)
SP96 Campylobacter gracilis (0%|0%|0.62%)
SP115 Lautropia mirabilis (0%|0%|0.20%)
SP36 Cardiobacterium hominis (0%|0%|0.39%)
SP103 Haemophilus parainfluenzae (0%|0%|1.21%)
SP113 Haemophilus haemolyticus (0%|0%|0.31%)
SP46 Haemophilus sp._HMT_036 (0%|0%|0.12%)
SP90 Fretibacterium sp._HMT_359 (0%|0%|0.24%)
SP2 Gardnerella vaginalis (0%|24.57%|0.97%)
SP22 Gardnerella swidsinskii (0%|8.31%|0%)
SP3 Gardnerella leopoldii (0%|0.53%|0%)
SP80 Actinomyces johnsonii (0%|0%|0.21%)
SP73 Actinomyces naeslundii (0%|0%|0.26%)
SP100 Actinomyces sp._HMT_169 (0%|0%|0.17%)
SP107 Actinomyces sp._HMT_175 (0%|0%|0.40%)
SP94 Actinomyces oris (0%|0%|0.16%)
SP29 Arthrobacter humicola (34.69%|0%|0%)
SP19 Corynebacterium sp._Oral_Taxon_B71 (20.11%|0%|0%)
SP9 Corynebacterium striatum (22.08%|0%|0%)
SP6 Corynebacterium appendicis (0%|3.71%|0%)
SP18 Corynebacterium mucifaciens (0%|1.71%|0%)
SP68 Lachnospiraceae_[G] sp._Oral_Taxon_B32 (0%|0%|0.27%)
SP1 Lachnospiraceae_[G-9] bacterium_HMT_924 (0%|3.73%|0%)
SP78 Fusobacterium nucleatum (0%|0%|0.15%)
SP87 Sneathia sanguinegens (0%|0.15%|0%)
SP72 Sneathia amnii_[Not_Validly_Published] (0.26%|0.22%|0%)
SP57 Ruminococcaceae_[G-1] bacterium_HMT_075 (0%|0%|0.16%)
SP106 Peptostreptococcaceae_[XI][G-9] [Eubacterium]_brachy (0%|0%|0.14%)
SP51 Parvimonas micra (0%|0%|0.88%)
SP48 Parvimonas sp._Oral_Taxon_110 (0%|0%|0.27%)
SP39 Parvimonas sp._HMT_110 (0%|0%|0.13%)
SP81 Megasphaera genomosp. (0%|0.28%|0%)
SP66 Veillonella sp._HMT_780 (0%|0%|1.02%)
SP42 Veillonella parvula (0%|0%|2.13%)
SP105 Selenomonas sp._HMT_134 (0%|0%|0.34%)
SP59 Selenomonas sputigena (0%|0%|1.35%)
SP114 Selenomonas sp._Oral_Taxon_E83 (0%|0%|0.29%)
SP92 Selenomonas sp._HMT_133 (0%|0%|0.18%)
SP69 Selenomonas sp._HMT_137 (0%|0%|0.66%)
SP67 Selenomonas sp._Oral_Taxon_137 (0%|0%|0.91%)
SP91 Mycoplasma hominis (0%|0.16%|0%)
SP5 Staphylococcus epidermidis (9.91%|0.32%|0%)
SP45 Gemella sp._Oral_Taxon_B93 (0%|0.56%|0%)
SP16 Gemella haemolysans (0%|0%|1.35%)
SP71 Gemella morbillorum (0%|0%|0.58%)
SP30 Gemella moribillum (0%|0%|0.93%)
SP31 Enterococcus faecalis (0%|3.26%|0%)
SP54 Enterococcus sp._str._F95 (0%|2.89%|0%)
SP43 Granulicatella elegans (0%|0%|2.77%)
SP88 Granulicatella paradiacens (0%|0%|0.43%)
SP44 Granulicatella adiacens (0%|0%|0.24%)
SP33 Aerococcus christensenii_Oral_Taxon_D44 (0%|0.48%|0%)
SP37 Lactobacillus salivarius (0%|0%|2.65%)
SP10 Lactobacillus fermentum (0%|0%|13.61%)
SP38 Limosilactobacillus coleohominis (0%|1.11%|0%)
SP40 Lactobacillus gasseri (0%|7.55%|6.60%)
SP28 Lactobacillus iners (0%|19.33%|0%)
SP7 Lactobacillus crispatus (4.85%|4.24%|0.18%)
SP13 Lactobacillus jensenii (0.36%|16.64%|0%)
SP79 Streptococcus agalactiae (7.74%|0.04%|0%)
SP4 Streptococcus sanguinis (0%|0%|2.06%)
SP35 Streptococcus salivarius (0%|0%|3.51%)
SP56 Streptococcus cristatus (0%|0%|2.89%)
SP84 Streptococcus oralis_subsp._tigurinus_clade_071 (0%|0%|0.46%)
SP60 Streptococcus oralis (0%|0%|2.23%)
SP52 Streptococcus oralis_subsp._tigurinus_clade_070 (0%|0%|1.19%)
SP55 Streptococcus sp._Oral_Taxon_71 (0%|0%|1.14%)
SP76 Streptococcus oralis_subsp._dentisani_clade_058 (0%|0%|0.23%)
SP26 Streptococcus sp._HMT_064 (0%|0%|0.49%)
SP11 Streptococcus sp._str._C300 (0%|0%|4.00%)
SP20 Streptococcus chosunense (0%|0%|4.39%)
SP17 Streptococcus mitis (0%|0%|8.62%)
Placenta
Vaginal
Oral
substitutions/site
0
0.2
0.4
0.6