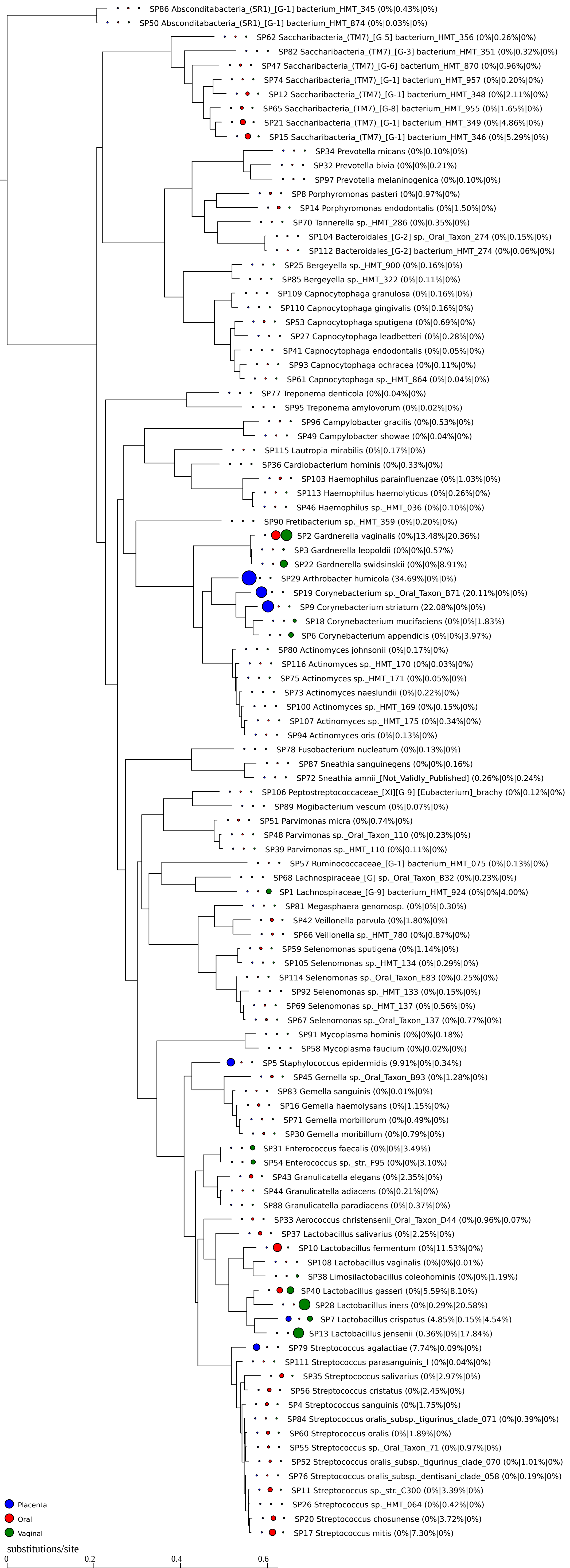
SP86 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345 (0%|0.43%|0%)
SP50 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_874 (0%|0.03%|0%)
SP62 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356 (0%|0.26%|0%)
SP82 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351 (0%|0.32%|0%)
SP47 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870 (0%|0.96%|0%)
SP74 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_957 (0%|0.20%|0%)
SP12 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348 (0%|2.11%|0%)
SP65 Saccharibacteria_(TM7)_[G-8] bacterium_HMT_955 (0%|1.65%|0%)
SP21 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349 (0%|4.86%|0%)
SP15 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346 (0%|5.29%|0%)
SP34 Prevotella micans (0%|0.10%|0%)
SP32 Prevotella bivia (0%|0%|0.21%)
SP97 Prevotella melaninogenica (0%|0.10%|0%)
SP8 Porphyromonas pasteri (0%|0.97%|0%)
SP14 Porphyromonas endodontalis (0%|1.50%|0%)
SP70 Tannerella sp._HMT_286 (0%|0.35%|0%)
SP104 Bacteroidales_[G-2] sp._Oral_Taxon_274 (0%|0.15%|0%)
SP112 Bacteroidales_[G-2] bacterium_HMT_274 (0%|0.06%|0%)
SP25 Bergeyella sp._HMT_900 (0%|0.16%|0%)
SP85 Bergeyella sp._HMT_322 (0%|0.11%|0%)
SP109 Capnocytophaga granulosa (0%|0.16%|0%)
SP110 Capnocytophaga gingivalis (0%|0.16%|0%)
SP53 Capnocytophaga sputigena (0%|0.69%|0%)
SP27 Capnocytophaga leadbetteri (0%|0.28%|0%)
SP41 Capnocytophaga endodontalis (0%|0.05%|0%)
SP93 Capnocytophaga ochracea (0%|0.11%|0%)
SP61 Capnocytophaga sp._HMT_864 (0%|0.04%|0%)
SP77 Treponema denticola (0%|0.04%|0%)
SP95 Treponema amylovorum (0%|0.02%|0%)
SP96 Campylobacter gracilis (0%|0.53%|0%)
SP49 Campylobacter showae (0%|0.04%|0%)
SP115 Lautropia mirabilis (0%|0.17%|0%)
SP36 Cardiobacterium hominis (0%|0.33%|0%)
SP103 Haemophilus parainfluenzae (0%|1.03%|0%)
SP113 Haemophilus haemolyticus (0%|0.26%|0%)
SP46 Haemophilus sp._HMT_036 (0%|0.10%|0%)
SP90 Fretibacterium sp._HMT_359 (0%|0.20%|0%)
SP2 Gardnerella vaginalis (0%|13.48%|20.36%)
SP3 Gardnerella leopoldii (0%|0%|0.57%)
SP22 Gardnerella swidsinskii (0%|0%|8.91%)
SP29 Arthrobacter humicola (34.69%|0%|0%)
SP19 Corynebacterium sp._Oral_Taxon_B71 (20.11%|0%|0%)
SP9 Corynebacterium striatum (22.08%|0%|0%)
SP18 Corynebacterium mucifaciens (0%|0%|1.83%)
SP6 Corynebacterium appendicis (0%|0%|3.97%)
SP80 Actinomyces johnsonii (0%|0.17%|0%)
SP116 Actinomyces sp._HMT_170 (0%|0.03%|0%)
SP75 Actinomyces sp._HMT_171 (0%|0.05%|0%)
SP73 Actinomyces naeslundii (0%|0.22%|0%)
SP100 Actinomyces sp._HMT_169 (0%|0.15%|0%)
SP107 Actinomyces sp._HMT_175 (0%|0.34%|0%)
SP94 Actinomyces oris (0%|0.13%|0%)
SP78 Fusobacterium nucleatum (0%|0.13%|0%)
SP87 Sneathia sanguinegens (0%|0%|0.16%)
SP72 Sneathia amnii_[Not_Validly_Published] (0.26%|0%|0.24%)
SP106 Peptostreptococcaceae_[XI][G-9] [Eubacterium]_brachy (0%|0.12%|0%)
SP89 Mogibacterium vescum (0%|0.07%|0%)
SP51 Parvimonas micra (0%|0.74%|0%)
SP48 Parvimonas sp._Oral_Taxon_110 (0%|0.23%|0%)
SP39 Parvimonas sp._HMT_110 (0%|0.11%|0%)
SP57 Ruminococcaceae_[G-1] bacterium_HMT_075 (0%|0.13%|0%)
SP68 Lachnospiraceae_[G] sp._Oral_Taxon_B32 (0%|0.23%|0%)
SP1 Lachnospiraceae_[G-9] bacterium_HMT_924 (0%|0%|4.00%)
SP81 Megasphaera genomosp. (0%|0%|0.30%)
SP42 Veillonella parvula (0%|1.80%|0%)
SP66 Veillonella sp._HMT_780 (0%|0.87%|0%)
SP59 Selenomonas sputigena (0%|1.14%|0%)
SP105 Selenomonas sp._HMT_134 (0%|0.29%|0%)
SP114 Selenomonas sp._Oral_Taxon_E83 (0%|0.25%|0%)
SP92 Selenomonas sp._HMT_133 (0%|0.15%|0%)
SP69 Selenomonas sp._HMT_137 (0%|0.56%|0%)
SP67 Selenomonas sp._Oral_Taxon_137 (0%|0.77%|0%)
SP91 Mycoplasma hominis (0%|0%|0.18%)
SP58 Mycoplasma faucium (0%|0.02%|0%)
SP5 Staphylococcus epidermidis (9.91%|0%|0.34%)
SP45 Gemella sp._Oral_Taxon_B93 (0%|1.28%|0%)
SP83 Gemella sanguinis (0%|0.01%|0%)
SP16 Gemella haemolysans (0%|1.15%|0%)
SP71 Gemella morbillorum (0%|0.49%|0%)
SP30 Gemella moribillum (0%|0.79%|0%)
SP31 Enterococcus faecalis (0%|0%|3.49%)
SP54 Enterococcus sp._str._F95 (0%|0%|3.10%)
SP43 Granulicatella elegans (0%|2.35%|0%)
SP44 Granulicatella adiacens (0%|0.21%|0%)
SP88 Granulicatella paradiacens (0%|0.37%|0%)
SP33 Aerococcus christensenii_Oral_Taxon_D44 (0%|0.96%|0.07%)
SP37 Lactobacillus salivarius (0%|2.25%|0%)
SP10 Lactobacillus fermentum (0%|11.53%|0%)
SP108 Lactobacillus vaginalis (0%|0%|0.01%)
SP38 Limosilactobacillus coleohominis (0%|0%|1.19%)
SP40 Lactobacillus gasseri (0%|5.59%|8.10%)
SP28 Lactobacillus iners (0%|0.29%|20.58%)
SP7 Lactobacillus crispatus (4.85%|0.15%|4.54%)
SP13 Lactobacillus jensenii (0.36%|0%|17.84%)
SP79 Streptococcus agalactiae (7.74%|0.09%|0%)
SP111 Streptococcus parasanguinis_I (0%|0.04%|0%)
SP35 Streptococcus salivarius (0%|2.97%|0%)
SP56 Streptococcus cristatus (0%|2.45%|0%)
SP4 Streptococcus sanguinis (0%|1.75%|0%)
SP84 Streptococcus oralis_subsp._tigurinus_clade_071 (0%|0.39%|0%)
SP60 Streptococcus oralis (0%|1.89%|0%)
SP55 Streptococcus sp._Oral_Taxon_71 (0%|0.97%|0%)
SP52 Streptococcus oralis_subsp._tigurinus_clade_070 (0%|1.01%|0%)
SP76 Streptococcus oralis_subsp._dentisani_clade_058 (0%|0.19%|0%)
SP11 Streptococcus sp._str._C300 (0%|3.39%|0%)
SP26 Streptococcus sp._HMT_064 (0%|0.42%|0%)
SP20 Streptococcus chosunense (0%|3.72%|0%)
SP17 Streptococcus mitis (0%|7.30%|0%)
Placenta
Oral
Vaginal
substitutions/site
0
0.2
0.4
0.6