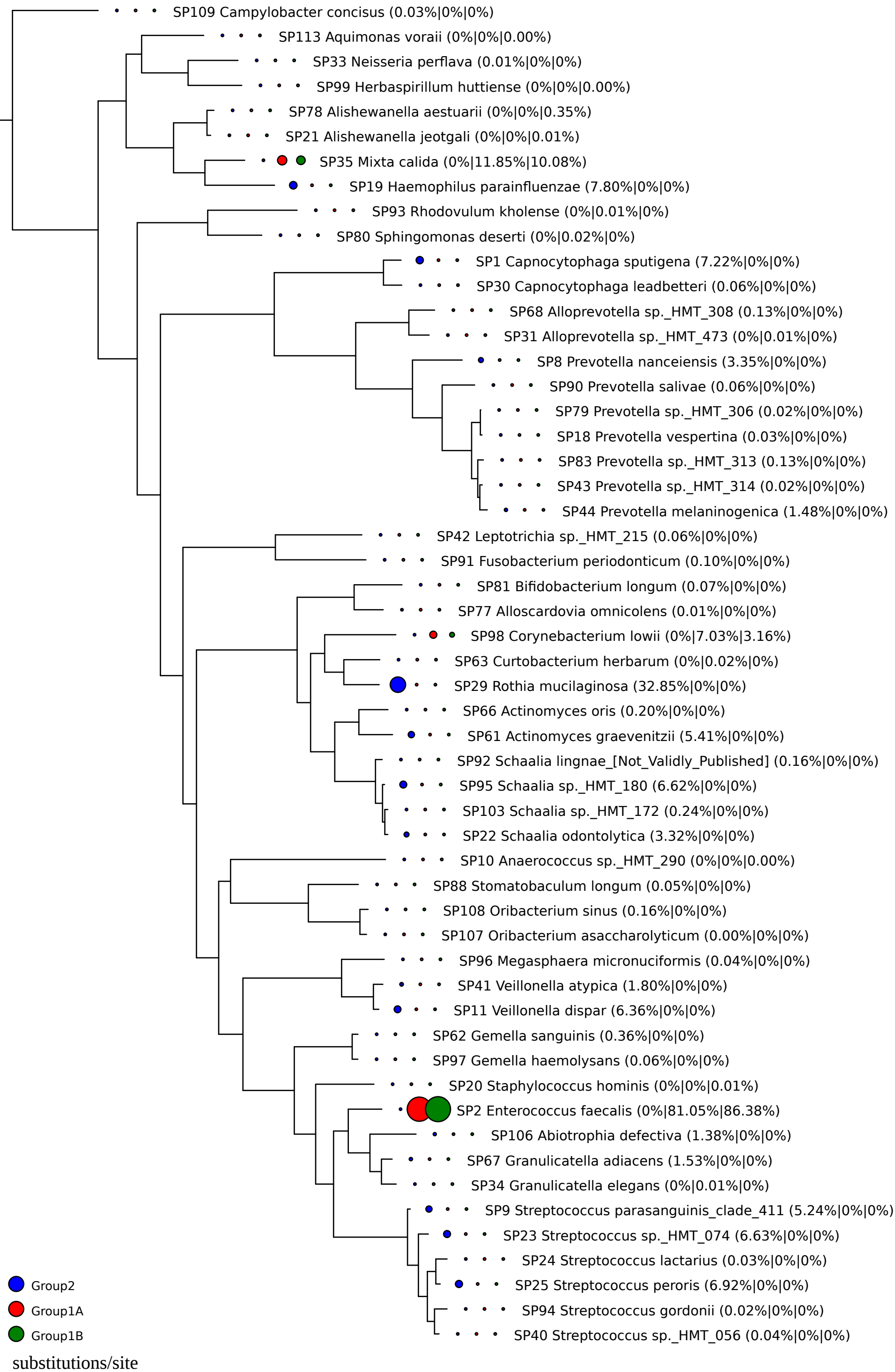
SP109 Campylobacter concisus (0.03%|0%|0%)
SP113 Aquimonas voraii (0%|0%|0.00%)
SP33 Neisseria perflava (0.01%|0%|0%)
SP99 Herbaspirillum huttiense (0%|0%|0.00%)
SP78 Alishewanella aestuarii (0%|0%|0.35%)
SP21 Alishewanella jeotgali (0%|0%|0.01%)
SP35 Mixta calida (0%|11.85%|10.08%)
SP19 Haemophilus parainfluenzae (7.80%|0%|0%)
SP93 Rhodovulum kholense (0%|0.01%|0%)
SP80 Sphingomonas deserti (0%|0.02%|0%)
SP1 Capnocytophaga sputigena (7.22%|0%|0%)
SP30 Capnocytophaga leadbetteri (0.06%|0%|0%)
SP68 Alloprevotella sp._HMT_308 (0.13%|0%|0%)
SP31 Alloprevotella sp._HMT_473 (0%|0.01%|0%)
SP8 Prevotella nanceiensis (3.35%|0%|0%)
SP90 Prevotella salivae (0.06%|0%|0%)
SP79 Prevotella sp._HMT_306 (0.02%|0%|0%)
SP18 Prevotella vespertina (0.03%|0%|0%)
SP83 Prevotella sp._HMT_313 (0.13%|0%|0%)
SP43 Prevotella sp._HMT_314 (0.02%|0%|0%)
SP44 Prevotella melaninogenica (1.48%|0%|0%)
SP42 Leptotrichia sp._HMT_215 (0.06%|0%|0%)
SP91 Fusobacterium periodonticum (0.10%|0%|0%)
SP81 Bifidobacterium longum (0.07%|0%|0%)
SP77 Alloscardovia omnicolens (0.01%|0%|0%)
SP98 Corynebacterium lowii (0%|7.03%|3.16%)
SP63 Curtobacterium herbarum (0%|0.02%|0%)
SP29 Rothia mucilaginosa (32.85%|0%|0%)
SP66 Actinomyces oris (0.20%|0%|0%)
SP61 Actinomyces graevenitzii (5.41%|0%|0%)
SP92 Schaalia lingnae_[Not_Validly_Published] (0.16%|0%|0%)
SP95 Schaalia sp._HMT_180 (6.62%|0%|0%)
SP103 Schaalia sp._HMT_172 (0.24%|0%|0%)
SP22 Schaalia odontolytica (3.32%|0%|0%)
SP10 Anaerococcus sp._HMT_290 (0%|0%|0.00%)
SP88 Stomatobaculum longum (0.05%|0%|0%)
SP108 Oribacterium sinus (0.16%|0%|0%)
SP107 Oribacterium asaccharolyticum (0.00%|0%|0%)
SP96 Megasphaera micronuciformis (0.04%|0%|0%)
SP41 Veillonella atypica (1.80%|0%|0%)
SP11 Veillonella dispar (6.36%|0%|0%)
SP62 Gemella sanguinis (0.36%|0%|0%)
SP97 Gemella haemolysans (0.06%|0%|0%)
SP20 Staphylococcus hominis (0%|0%|0.01%)
SP2 Enterococcus faecalis (0%|81.05%|86.38%)
SP106 Abiotrophia defectiva (1.38%|0%|0%)
SP67 Granulicatella adiacens (1.53%|0%|0%)
SP34 Granulicatella elegans (0%|0.01%|0%)
SP9 Streptococcus parasanguinis_clade_411 (5.24%|0%|0%)
SP23 Streptococcus sp._HMT_074 (6.63%|0%|0%)
SP24 Streptococcus lactarius (0.03%|0%|0%)
SP25 Streptococcus peroris (6.92%|0%|0%)
SP94 Streptococcus gordonii (0.02%|0%|0%)
SP40 Streptococcus sp._HMT_056 (0.04%|0%|0%)
Group2
Group1A
Group1B
substitutions/site
0
0.2
0.4
0.6