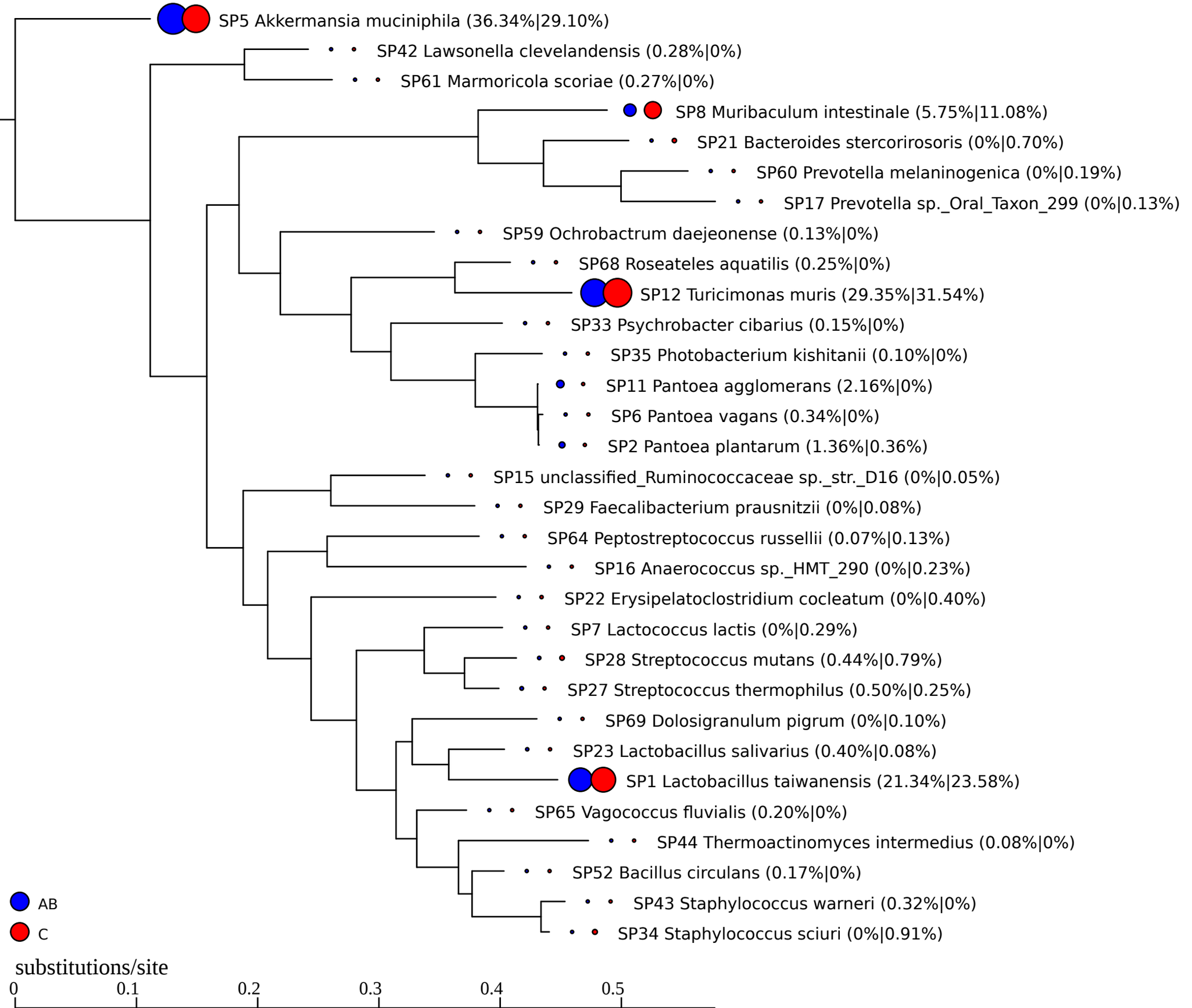
SP5 Akkermansia muciniphila (36.34%|29.10%)
SP42 Lawsonella clevelandensis (0.28%|0%)
SP61 Marmoricola scoriae (0.27%|0%)
SP8 Muribaculum intestinale (5.75%|11.08%)
SP21 Bacteroides stercorirosoris (0%|0.70%)
SP60 Prevotella melaninogenica (0%|0.19%)
SP17 Prevotella sp._Oral_Taxon_299 (0%|0.13%)
SP59 Ochrobactrum daejeonense (0.13%|0%)
SP68 Roseateles aquatilis (0.25%|0%)
SP12 Turicimonas muris (29.35%|31.54%)
SP33 Psychrobacter cibarius (0.15%|0%)
SP35 Photobacterium kishitanii (0.10%|0%)
SP11 Pantoea agglomerans (2.16%|0%)
SP6 Pantoea vagans (0.34%|0%)
SP2 Pantoea plantarum (1.36%|0.36%)
SP15 unclassified_Ruminococcaceae sp._str._D16 (0%|0.05%)
SP29 Faecalibacterium prausnitzii (0%|0.08%)
SP64 Peptostreptococcus russellii (0.07%|0.13%)
SP16 Anaerococcus sp._HMT_290 (0%|0.23%)
SP22 Erysipelatoclostridium cocleatum (0%|0.40%)
SP7 Lactococcus lactis (0%|0.29%)
SP28 Streptococcus mutans (0.44%|0.79%)
SP27 Streptococcus thermophilus (0.50%|0.25%)
SP69 Dolosigranulum pigrum (0%|0.10%)
SP23 Lactobacillus salivarius (0.40%|0.08%)
SP1 Lactobacillus taiwanensis (21.34%|23.58%)
SP65 Vagococcus fluvialis (0.20%|0%)
SP44 Thermoactinomyces intermedius (0.08%|0%)
SP52 Bacillus circulans (0.17%|0%)
SP43 Staphylococcus warneri (0.32%|0%)
SP34 Staphylococcus sciuri (0%|0.91%)
AB
C
substitutions/site
0
0.1
0.2
0.3
0.4
0.5