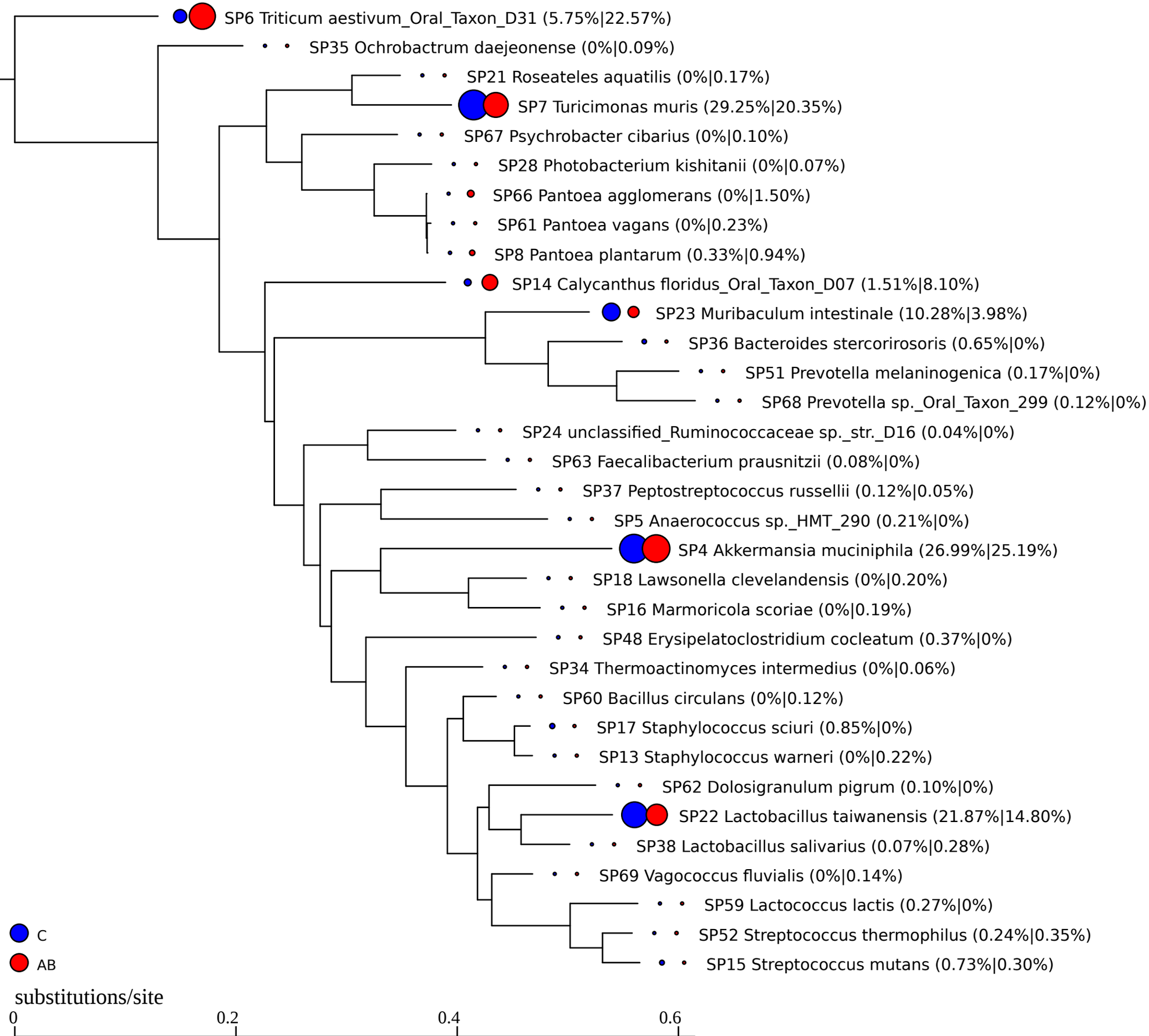
SP6 Triticum aestivum_Oral_Taxon_D31 (5.75%|22.57%)
SP35 Ochrobactrum daejeonense (0%|0.09%)
SP21 Roseateles aquatilis (0%|0.17%)
SP7 Turicimonas muris (29.25%|20.35%)
SP67 Psychrobacter cibarius (0%|0.10%)
SP28 Photobacterium kishitanii (0%|0.07%)
SP66 Pantoea agglomerans (0%|1.50%)
SP61 Pantoea vagans (0%|0.23%)
SP8 Pantoea plantarum (0.33%|0.94%)
SP14 Calycanthus floridus_Oral_Taxon_D07 (1.51%|8.10%)
SP23 Muribaculum intestinale (10.28%|3.98%)
SP36 Bacteroides stercorirosoris (0.65%|0%)
SP51 Prevotella melaninogenica (0.17%|0%)
SP68 Prevotella sp._Oral_Taxon_299 (0.12%|0%)
SP24 unclassified_Ruminococcaceae sp._str._D16 (0.04%|0%)
SP63 Faecalibacterium prausnitzii (0.08%|0%)
SP37 Peptostreptococcus russellii (0.12%|0.05%)
SP5 Anaerococcus sp._HMT_290 (0.21%|0%)
SP4 Akkermansia muciniphila (26.99%|25.19%)
SP18 Lawsonella clevelandensis (0%|0.20%)
SP16 Marmoricola scoriae (0%|0.19%)
SP48 Erysipelatoclostridium cocleatum (0.37%|0%)
SP34 Thermoactinomyces intermedius (0%|0.06%)
SP60 Bacillus circulans (0%|0.12%)
SP17 Staphylococcus sciuri (0.85%|0%)
SP13 Staphylococcus warneri (0%|0.22%)
SP62 Dolosigranulum pigrum (0.10%|0%)
SP22 Lactobacillus taiwanensis (21.87%|14.80%)
SP38 Lactobacillus salivarius (0.07%|0.28%)
SP69 Vagococcus fluvialis (0%|0.14%)
SP59 Lactococcus lactis (0.27%|0%)
SP52 Streptococcus thermophilus (0.24%|0.35%)
SP15 Streptococcus mutans (0.73%|0.30%)
C
AB
substitutions/site
0
0.2
0.4
0.6