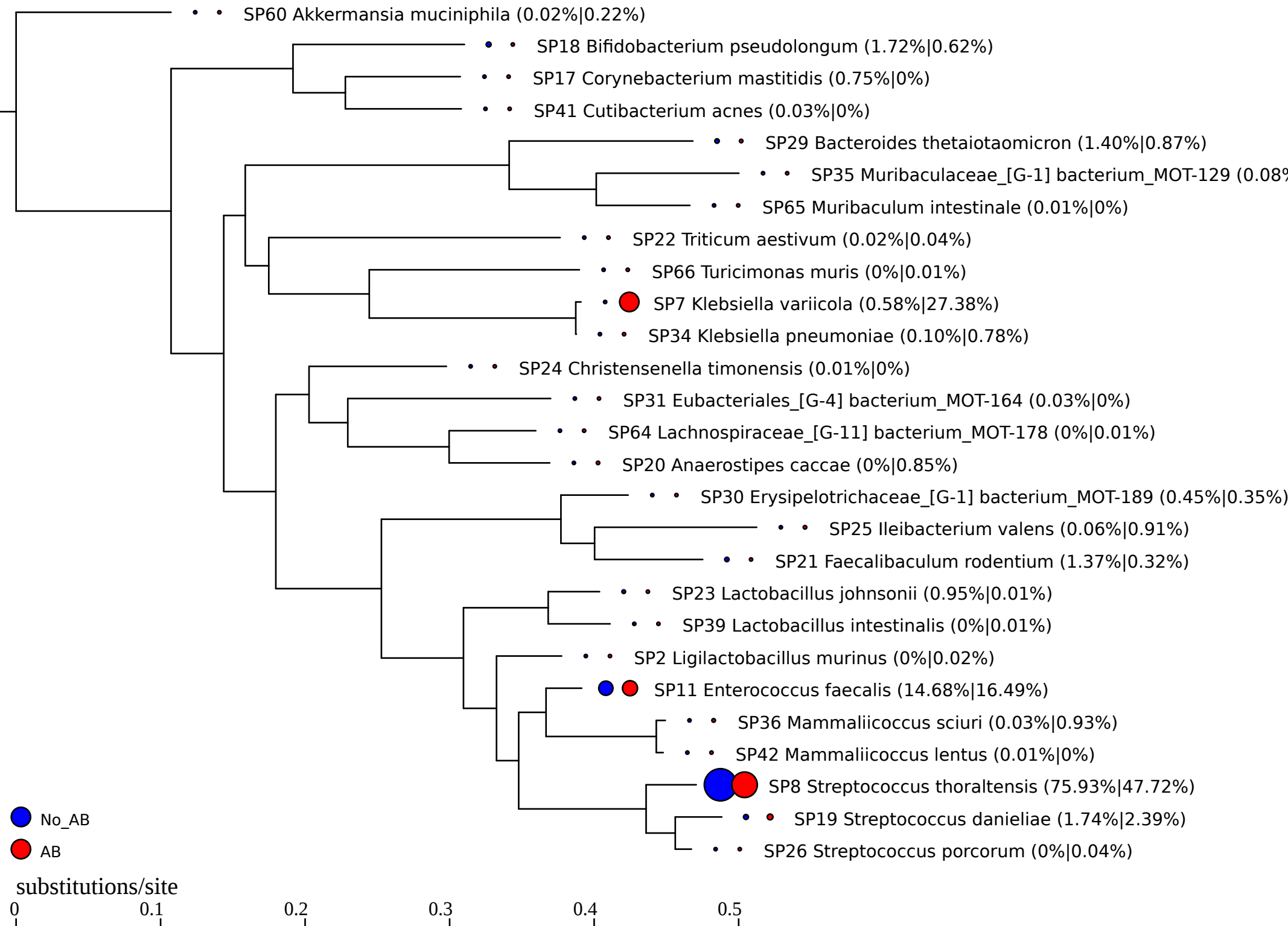
SP60 Akkermansia muciniphila (0.02%|0.22%)
SP18 Bifidobacterium pseudolongum (1.72%|0.62%)
SP17 Corynebacterium mastitidis (0.75%|0%)
SP41 Cutibacterium acnes (0.03%|0%)
SP29 Bacteroides thetaiotaomicron (1.40%|0.87%)
SP35 Muribaculaceae_[G-1] bacterium_MOT-129 (0.08%
SP65 Muribaculum intestinale (0.01%|0%)
SP22 Triticum aestivum (0.02%|0.04%)
SP66 Turicimonas muris (0%|0.01%)
SP7 Klebsiella variicola (0.58%|27.38%)
SP34 Klebsiella pneumoniae (0.10%|0.78%)
SP24 Christensenella timonensis (0.01%|0%)
SP31 Eubacteriales_[G-4] bacterium_MOT-164 (0.03%|0%)
SP64 Lachnospiraceae_[G-11] bacterium_MOT-178 (0%|0.01%)
SP20 Anaerostipes caccae (0%|0.85%)
SP30 Erysipelotrichaceae_[G-1] bacterium_MOT-189 (0.45%|0.35%)
SP25 Ileibacterium valens (0.06%|0.91%)
SP21 Faecalibaculum rodentium (1.37%|0.32%)
SP23 Lactobacillus johnsonii (0.95%|0.01%)
SP39 Lactobacillus intestinalis (0%|0.01%)
SP2 Ligilactobacillus murinus (0%|0.02%)
SP11 Enterococcus faecalis (14.68%|16.49%)
SP36 Mammaliicoccus sciuri (0.03%|0.93%)
SP42 Mammaliicoccus lentus (0.01%|0%)
SP8 Streptococcus thoraltensis (75.93%|47.72%)
SP19 Streptococcus danieliae (1.74%|2.39%)
SP26 Streptococcus porcorum (0%|0.04%)
No_AB
AB
substitutions/site
0
0.1
0.2
0.3
0.4
0.5