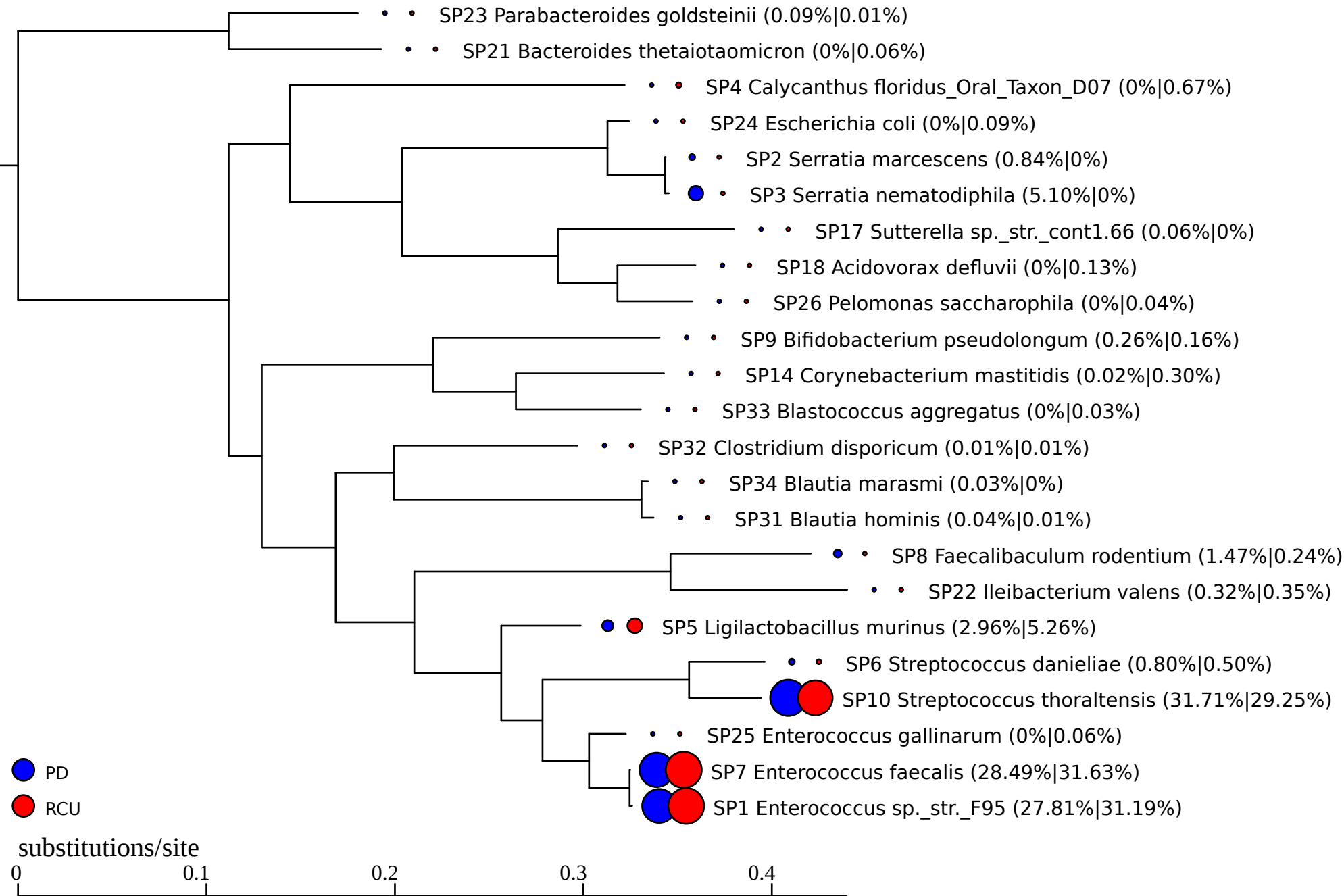
SP23 Parabacteroides goldsteinii (0.09%|0.01%)
SP21 Bacteroides thetaiotaomicron (0%|0.06%)
SP4 Calycanthus floridus_Oral_Taxon_D07 (0%|0.67%)
SP24 Escherichia coli (0%|0.09%)
SP2 Serratia marcescens (0.84%|0%)
SP3 Serratia nematodiphila (5.10%|0%)
SP17 Sutterella sp._str._cont1.66 (0.06%|0%)
SP18 Acidovorax defluvii (0%|0.13%)
SP26 Pelomonas saccharophila (0%|0.04%)
SP9 Bifidobacterium pseudolongum (0.26%|0.16%)
SP14 Corynebacterium mastitidis (0.02%|0.30%)
SP33 Blastococcus aggregatus (0%|0.03%)
SP32 Clostridium disporicum (0.01%|0.01%)
SP34 Blautia marasmi (0.03%|0%)
SP31 Blautia hominis (0.04%|0.01%)
SP8 Faecalibaculum rodentium (1.47%|0.24%)
SP22 Ileibacterium valens (0.32%|0.35%)
SP5 Ligilactobacillus murinus (2.96%|5.26%)
SP6 Streptococcus danieliae (0.80%|0.50%)
SP10 Streptococcus thoraltensis (31.71%|29.25%)
SP25 Enterococcus gallinarum (0%|0.06%)
SP7 Enterococcus faecalis (28.49%|31.63%)
SP1 Enterococcus sp._str._F95 (27.81%|31.19%)
PD
RCU
substitutions/site
0
0.1
0.2
0.3
0.4