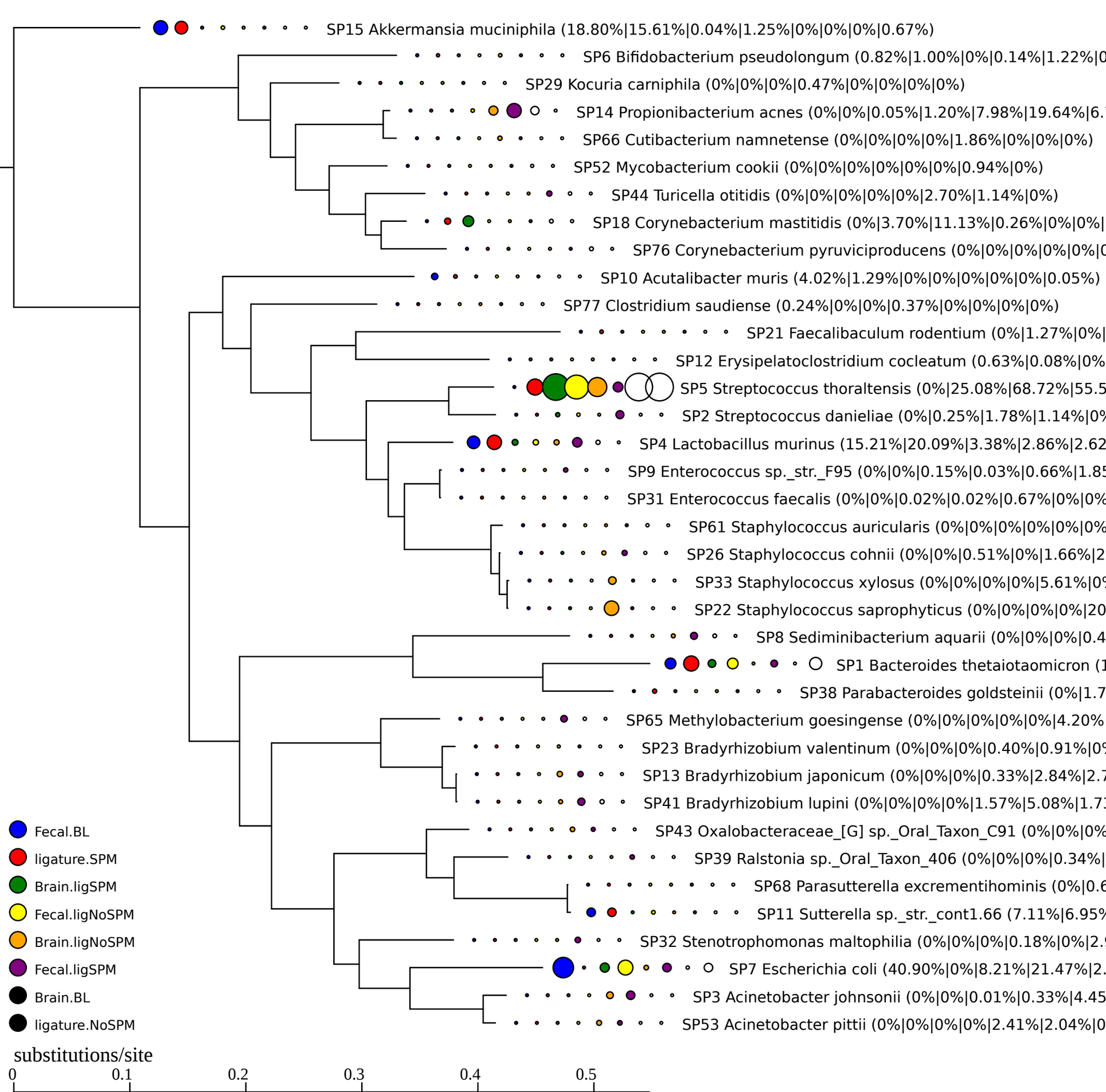
SP15 Akkermansia muciniphila (18.80%|15.61%|0.04%|1.25%|0%|0%|0%|0.67%)
SP6 Bifidobacterium pseudolongum (0.82%|1.00%|0%|0.14%|1.22%|0
SP29 Kocuria carniphila (0%|0%|0%|0.47%|0%|0%|0%|0%)
SP14 Propionibacterium acnes (0%|0%|0.05%|1.20%|7.98%|19.64%|6.
SP66 Cutibacterium namnetense (0%|0%|0%|0%|1.86%|0%|0%|0%)
SP52 Mycobacterium cookii (0%|0%|0%|0%|0%|0%|0.94%|0%)
SP44 Turicella otitidis (0%|0%|0%|0%|0%|2.70%|1.14%|0%)
SP18 Corynebacterium mastitidis (0%|3.70%|11.13%|0.26%|0%|0%|
SP76 Corynebacterium pyruviciproducens (0%|0%|0%|0%|0%|0
SP10 Acutalibacter muris (4.02%|1.29%|0%|0%|0%|0%|0%|0.05%)
SP77 Clostridium saudiense (0.24%|0%|0%|0.37%|0%|0%|0%|0%)
SP21 Faecalibaculum rodentium (0%|1.27%|0%|
SP12 Erysipelatoclostridium cocleatum (0.63%|0.08%|0%
SP5 Streptococcus thoraltensis (0%|25.08%|68.72%|55.5
SP2 Streptococcus danieliae (0%|0.25%|1.78%|1.14%|0%
SP4 Lactobacillus murinus (15.21%|20.09%|3.38%|2.86%|2.62
SP9 Enterococcus sp._str._F95 (0%|0%|0.15%|0.03%|0.66%|1.85
SP31 Enterococcus faecalis (0%|0%|0.02%|0.02%|0.67%|0%|0%
SP61 Staphylococcus auricularis (0%|0%|0%|0%|0%|0%
SP26 Staphylococcus cohnii (0%|0%|0.51%|0%|1.66%|2
SP33 Staphylococcus xylosus (0%|0%|0%|0%|5.61%|0
SP22 Staphylococcus saprophyticus (0%|0%|0%|0%|20
SP8 Sediminibacterium aquarii (0%|0%|0%|0.4
SP1 Bacteroides thetaiotaomicron (1
SP38 Parabacteroides goldsteinii (0%|1.7
SP65 Methylobacterium goesingense (0%|0%|0%|0%|0%|4.20%
SP23 Bradyrhizobium valentinum (0%|0%|0%|0.40%|0.91%|0
SP13 Bradyrhizobium japonicum (0%|0%|0%|0.33%|2.84%|2.7
SP41 Bradyrhizobium lupini (0%|0%|0%|0%|1.57%|5.08%|1.7
SP43 Oxalobacteraceae_[G] sp._Oral_Taxon_C91 (0%|0%|0%
SP39 Ralstonia sp._Oral_Taxon_406 (0%|0%|0%|0.34%|
SP68 Parasutterella excrementihominis (0%|0.6
SP11 Sutterella sp._str._cont1.66 (7.11%|6.95%
SP32 Stenotrophomonas maltophilia (0%|0%|0%|0.18%|0%|2.
SP7 Escherichia coli (40.90%|0%|8.21%|21.47%|2.
SP3 Acinetobacter johnsonii (0%|0%|0.01%|0.33%|4.45
SP53 Acinetobacter pittii (0%|0%|0%|0%|2.41%|2.04%|0
Fecal.BL
ligature.SPM
Brain.ligSPM
Fecal.ligNoSPM
Brain.ligNoSPM
Fecal.ligSPM
Brain.BL
ligature.NoSPM
substitutions/site
0
0.1
0.2
0.3
0.4
0.5