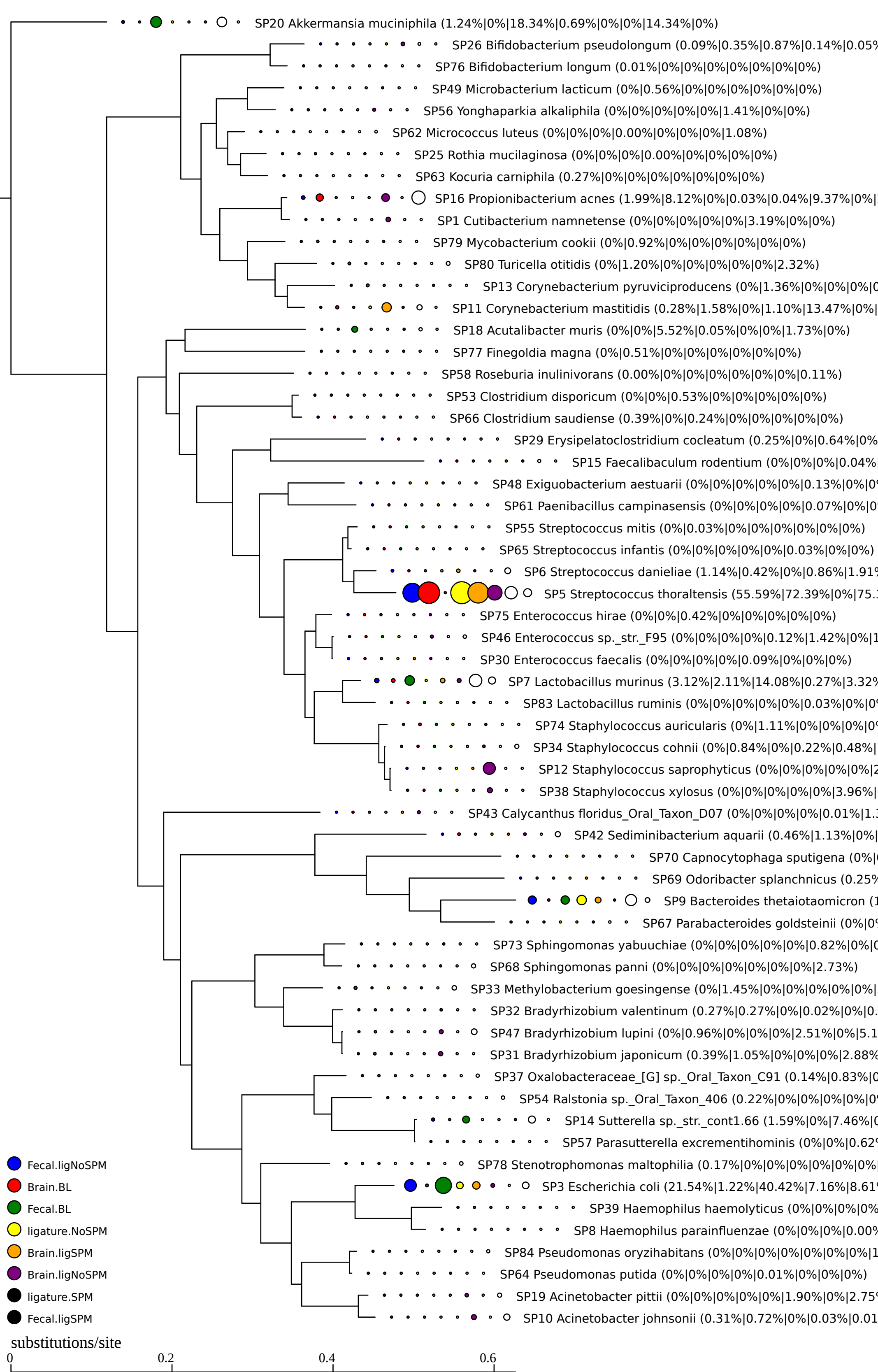
SP20 Akkermansia muciniphila (1.24%|0%|18.34%|0.69%|0%|0%|14.34%|0%)
SP26 Bifidobacterium pseudolongum (0.09%|0.35%|0.87%|0.14%|0.05%
SP76 Bifidobacterium longum (0.01%|0%|0%|0%|0%|0%|0%|0%)
SP49 Microbacterium lacticum (0%|0.56%|0%|0%|0%|0%|0%|0%)
SP56 Yonghaparkia alkaliphila (0%|0%|0%|0%|0%|1.41%|0%|0%)
SP62 Micrococcus luteus (0%|0%|0%|0.00%|0%|0%|0%|1.08%)
SP25 Rothia mucilaginosa (0%|0%|0%|0.00%|0%|0%|0%|0%)
SP63 Kocuria carniphila (0.27%|0%|0%|0%|0%|0%|0%|0%)
SP16 Propionibacterium acnes (1.99%|8.12%|0%|0.03%|0.04%|9.37%|0%|
SP1 Cutibacterium namnetense (0%|0%|0%|0%|0%|3.19%|0%|0%)
SP79 Mycobacterium cookii (0%|0.92%|0%|0%|0%|0%|0%|0%)
SP80 Turicella otitidis (0%|1.20%|0%|0%|0%|0%|0%|2.32%)
SP13 Corynebacterium pyruviciproducens (0%|1.36%|0%|0%|0%|0
SP11 Corynebacterium mastitidis (0.28%|1.58%|0%|1.10%|13.47%|0%|
SP18 Acutalibacter muris (0%|0%|5.52%|0.05%|0%|0%|1.73%|0%)
SP77 Finegoldia magna (0%|0.51%|0%|0%|0%|0%|0%|0%)
SP58 Roseburia inulinivorans (0.00%|0%|0%|0%|0%|0%|0%|0.11%)
SP53 Clostridium disporicum (0%|0%|0.53%|0%|0%|0%|0%|0%)
SP66 Clostridium saudiense (0.39%|0%|0.24%|0%|0%|0%|0%|0%)
SP29 Erysipelatoclostridium cocleatum (0.25%|0%|0.64%|0%
SP15 Faecalibaculum rodentium (0%|0%|0%|0.04%|
SP48 Exiguobacterium aestuarii (0%|0%|0%|0%|0%|0.13%|0%|0
SP61 Paenibacillus campinasensis (0%|0%|0%|0%|0.07%|0%|0
SP55 Streptococcus mitis (0%|0.03%|0%|0%|0%|0%|0%|0%)
SP65 Streptococcus infantis (0%|0%|0%|0%|0%|0.03%|0%|0%)
SP6 Streptococcus danieliae (1.14%|0.42%|0%|0.86%|1.91%
SP5 Streptococcus thoraltensis (55.59%|72.39%|0%|75.
SP75 Enterococcus hirae (0%|0%|0.42%|0%|0%|0%|0%|0%)
SP46 Enterococcus sp._str._F95 (0%|0%|0%|0%|0.12%|1.42%|0%|
SP30 Enterococcus faecalis (0%|0%|0%|0%|0.09%|0%|0%|0%)
SP7 Lactobacillus murinus (3.12%|2.11%|14.08%|0.27%|3.32%
SP83 Lactobacillus ruminis (0%|0%|0%|0%|0%|0.03%|0%|0
SP74 Staphylococcus auricularis (0%|1.11%|0%|0%|0%|0
SP34 Staphylococcus cohnii (0%|0.84%|0%|0.22%|0.48%|
SP12 Staphylococcus saprophyticus (0%|0%|0%|0%|0%|2
SP38 Staphylococcus xylosus (0%|0%|0%|0%|0%|3.96%|
SP43 Calycanthus floridus_Oral_Taxon_D07 (0%|0%|0%|0%|0.01%|1.
SP42 Sediminibacterium aquarii (0.46%|1.13%|0%|
SP70 Capnocytophaga sputigena (0%|
SP69 Odoribacter splanchnicus (0.25%
SP9 Bacteroides thetaiotaomicron (1
SP67 Parabacteroides goldsteinii (0%|0
SP73 Sphingomonas yabuuchiae (0%|0%|0%|0%|0%|0.82%|0%|0
SP68 Sphingomonas panni (0%|0%|0%|0%|0%|0%|0%|2.73%)
SP33 Methylobacterium goesingense (0%|1.45%|0%|0%|0%|0%|0%|
SP32 Bradyrhizobium valentinum (0.27%|0.27%|0%|0.02%|0%|0.
SP47 Bradyrhizobium lupini (0%|0.96%|0%|0%|0%|2.51%|0%|5.1
SP31 Bradyrhizobium japonicum (0.39%|1.05%|0%|0%|0%|2.88%
SP37 Oxalobacteraceae_[G] sp._Oral_Taxon_C91 (0.14%|0.83%|0
SP54 Ralstonia sp._Oral_Taxon_406 (0.22%|0%|0%|0%|0%|0%|0
SP14 Sutterella sp._str._cont1.66 (1.59%|0%|7.46%|0
SP57 Parasutterella excrementihominis (0%|0%|0.62
SP78 Stenotrophomonas maltophilia (0.17%|0%|0%|0%|0%|0%|0%|
SP3 Escherichia coli (21.54%|1.22%|40.42%|7.16%|8.61
SP39 Haemophilus haemolyticus (0%|0%|0%|0%
SP8 Haemophilus parainfluenzae (0%|0%|0%|0.00%
SP84 Pseudomonas oryzihabitans (0%|0%|0%|0%|0%|0%|0%|1
SP64 Pseudomonas putida (0%|0%|0%|0%|0.01%|0%|0%|0%)
SP19 Acinetobacter pittii (0%|0%|0%|0%|0%|1.90%|0%|2.75
SP10 Acinetobacter johnsonii (0.31%|0.72%|0%|0.03%|0.01
Fecal.ligNoSPM
Brain.BL
Fecal.BL
ligature.NoSPM
Brain.ligSPM
Brain.ligNoSPM
ligature.SPM
Fecal.ligSPM
substitutions/site
0
0.2
0.4
0.6