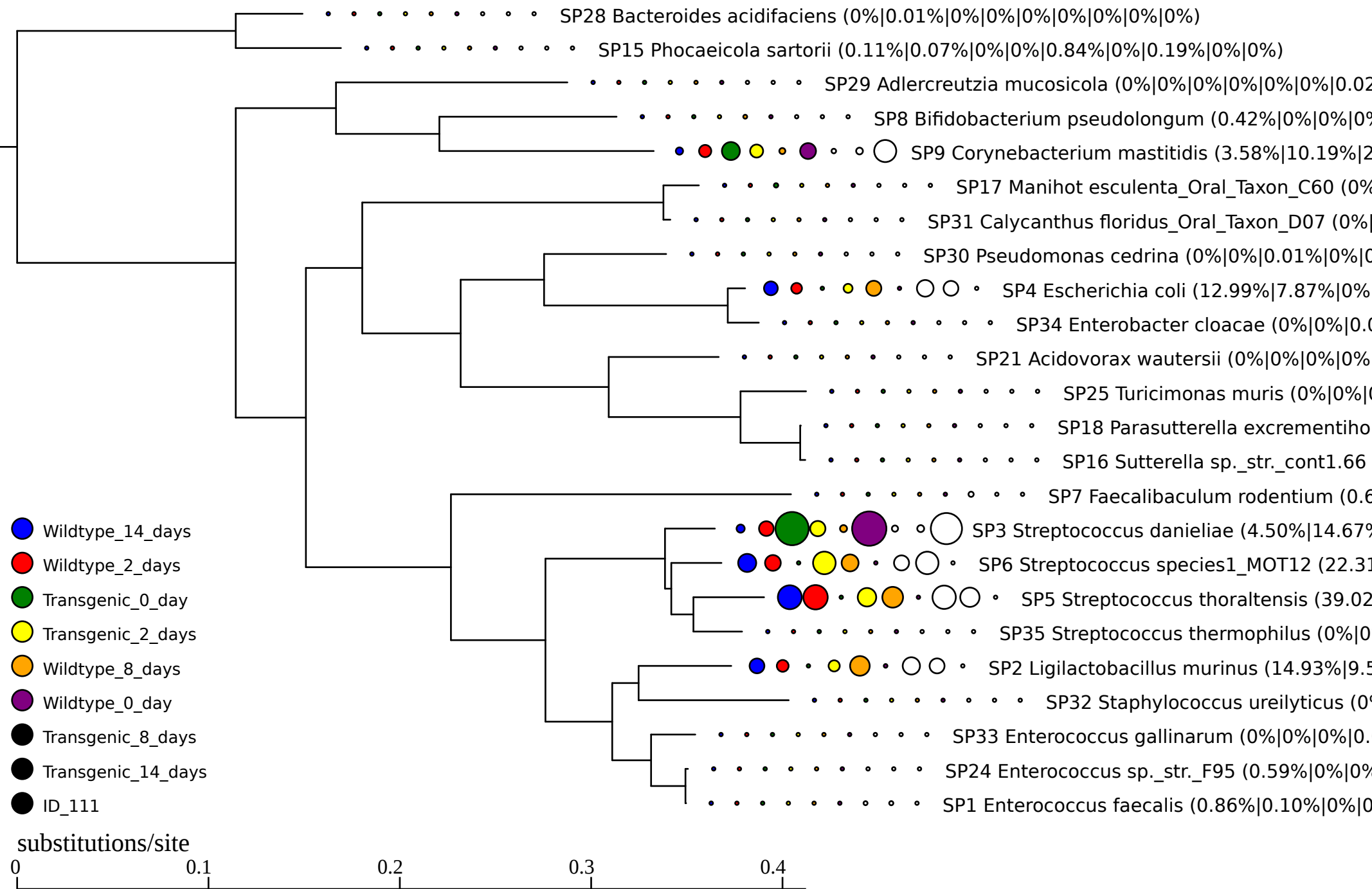
SP28 Bacteroides acidifaciens (0%|0.01%|0%|0%|0%|0%|0%|0%|0%)
SP15 Phocaeicola sartorii (0.11%|0.07%|0%|0%|0.84%|0%|0.19%|0%|0%)
SP29 Adlercreutzia mucosicola (0%|0%|0%|0%|0%|0%|0.02
SP8 Bifidobacterium pseudolongum (0.42%|0%|0%|0%
SP9 Corynebacterium mastitidis (3.58%|10.19%|2
SP17 Manihot esculenta_Oral_Taxon_C60 (0%
SP31 Calycanthus floridus_Oral_Taxon_D07 (0%|
SP30 Pseudomonas cedrina (0%|0%|0.01%|0%|0
SP4 Escherichia coli (12.99%|7.87%|0%
SP34 Enterobacter cloacae (0%|0%|0.0
SP21 Acidovorax wautersii (0%|0%|0%|0%
SP25 Turicimonas muris (0%|0%|0
SP18 Parasutterella excrementiho
SP16 Sutterella sp._str._cont1.66
SP7 Faecalibaculum rodentium (0.6
SP3 Streptococcus danieliae (4.50%|14.67%
SP6 Streptococcus species1_MOT12 (22.31
SP5 Streptococcus thoraltensis (39.02
SP35 Streptococcus thermophilus (0%|0
SP2 Ligilactobacillus murinus (14.93%|9.5
SP32 Staphylococcus ureilyticus (0%
SP33 Enterococcus gallinarum (0%|0%|0%|0.
SP24 Enterococcus sp._str._F95 (0.59%|0%|0%
SP1 Enterococcus faecalis (0.86%|0.10%|0%|0
Wildtype_14_days
Wildtype_2_days
Transgenic_0_day
Transgenic_2_days
Wildtype_8_days
Wildtype_0_day
Transgenic_8_days
Transgenic_14_days
ID_111
substitutions/site
0
0.1
0.2
0.3
0.4