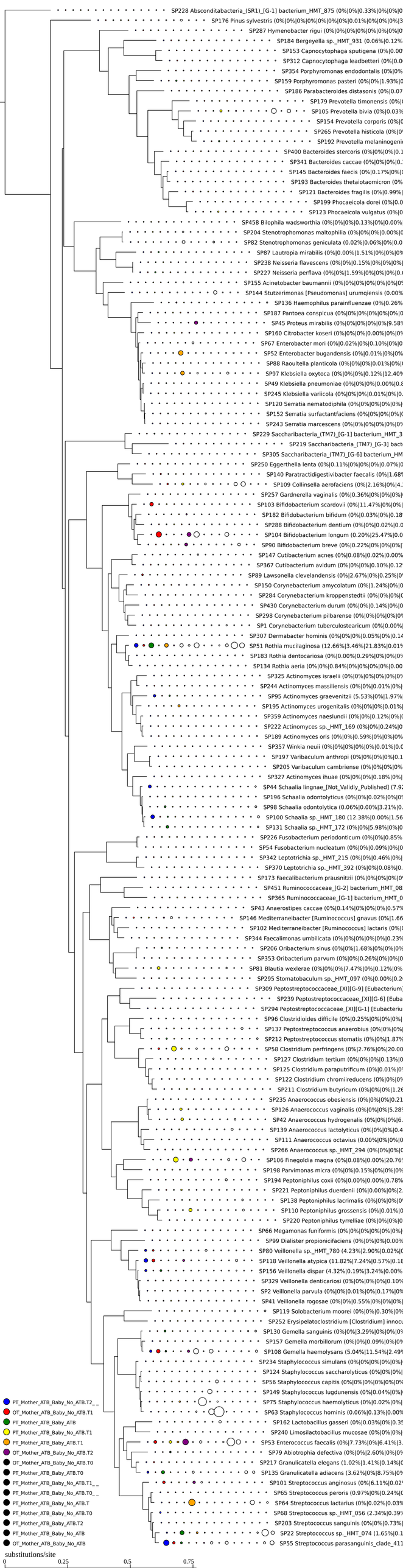
SP228 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_875 (0%|0%|0.33%|0%|0%|0
SP176 Pinus sylvestris (0%|0%|0%|0%|0%|0%|0.01%|0%|0%|0%|3
SP287 Hymenobacter rigui (0%|0%|0%|0%|0%|0%|0%|0
SP184 Bergeyella sp._HMT_931 (0.06%|0.12%
SP153 Capnocytophaga sputigena (0%|0.00
SP312 Capnocytophaga leadbetteri (0%|0.0
SP354 Porphyromonas endodontalis (0%|0%
SP159 Porphyromonas pasteri (0%|0%|1.93%|0
SP186 Parabacteroides distasonis (0%|0.07
SP179 Prevotella timonensis (0%|
SP105 Prevotella bivia (0%|0.03%
SP154 Prevotella corporis (0%|0
SP265 Prevotella histicola (0%|0
SP192 Prevotella melaninogenic
SP400 Bacteroides stercoris (0%|0%|0%|0.1
SP341 Bacteroides caccae (0%|0%|0%|0.
SP145 Bacteroides faecis (0%|0.17%|0%|
SP193 Bacteroides thetaiotaomicron (0%
SP121 Bacteroides fragilis (0%|0.99%|
SP199 Phocaeicola dorei (0%|0.0
SP123 Phocaeicola vulgatus (0%|0
SP458 Bilophila wadsworthia (0%|0%|0%|0.13%|0%|0.00%
SP204 Stenotrophomonas maltophilia (0%|0%|0%|0.00%|0
SP82 Stenotrophomonas geniculata (0.02%|0.06%|0%|0.0
SP87 Lautropia mirabilis (0%|0.00%|1.51%|0%|0%|0%
SP238 Neisseria flavescens (0%|0%|0.15%|0%|0%|0%|
SP227 Neisseria perflava (0%|0%|1.59%|0%|0%|0%|0.
SP155 Acinetobacter baumannii (0%|0%|0%|0%|0%|0%|0%
SP144 Stutzerimonas [Pseudomonas] urumqiensis (0.00%
SP136 Haemophilus parainfluenzae (0%|0.26%
SP187 Pantoea conspicua (0%|0%|0%|0%|0%|0%|0
SP45 Proteus mirabilis (0%|0%|0%|0%|0%|9.58%
SP160 Citrobacter koseri (0%|0%|0%|0.00%|0%|0%
SP67 Enterobacter mori (0%|0.02%|0%|0.10%|0%|0
SP52 Enterobacter bugandensis (0%|0.01%|0%|0%
SP88 Raoultella planticola (0%|0%|0%|0.01%|0%|
SP97 Klebsiella oxytoca (0%|0%|0%|0.12%|12.40%
SP49 Klebsiella pneumoniae (0%|0%|0%|0.00%|0.8
SP245 Klebsiella variicola (0%|0%|0%|0.01%|0%|0.
SP120 Serratia nematodiphila (0%|0%|0%|0%|0%|
SP152 Serratia surfactantfaciens (0%|0%|0%|0%|0
SP243 Serratia marcescens (0%|0%|0%|0%|0%|0%
SP229 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_3
SP219 Saccharibacteria_(TM7)_[G-3] bact
SP305 Saccharibacteria_(TM7)_[G-6] bacterium_HM
SP250 Eggerthella lenta (0%|0.11%|0%|0%|0%|0.07%|0
SP140 Paratractidigestivibacter faecalis (0%|1.68%
SP109 Collinsella aerofaciens (0%|2.16%|0%|4.
SP257 Gardnerella vaginalis (0%|0.36%|0%|0%|0%|
SP103 Bifidobacterium scardovii (0%|11.47%|0%|0%|
SP182 Bifidobacterium bifidum (0%|0.03%|0%|0.18
SP288 Bifidobacterium dentium (0%|0%|0.02%|0.0
SP104 Bifidobacterium longum (0.20%|25.47%|0.0
SP90 Bifidobacterium breve (0%|0.22%|0%|0%|0%|
SP147 Cutibacterium acnes (0%|0.08%|0.02%|0.00%
SP367 Cutibacterium avidum (0%|0%|0%|0.10%|0.12
SP89 Lawsonella clevelandensis (0%|2.67%|0%|0.25%|0
SP150 Corynebacterium amycolatum (0%|1.24%|0%|0
SP284 Corynebacterium kroppenstedtii (0%|0%|0%|0
SP430 Corynebacterium durum (0%|0%|0.14%|0%|0
SP298 Corynebacterium pilbarense (0%|0%|0%|0%|0
SP1 Corynebacterium tuberculostearicum (0%|0.00%
SP307 Dermabacter hominis (0%|0%|0%|0.05%|0%|0.14
SP51 Rothia mucilaginosa (12.66%|3.46%|21.83%|0.01%
SP183 Rothia dentocariosa (0%|0.00%|0.29%|0%|0%|0
SP134 Rothia aeria (0%|0%|0.84%|0%|0%|0%|0.00
SP325 Actinomyces israelii (0%|0%|0%|0%|0%|0
SP244 Actinomyces massiliensis (0%|0%|0.01%|0%|
SP95 Actinomyces graevenitzii (5.53%|0%|1.97%
SP195 Actinomyces urogenitalis (0%|0%|0%|0.01%
SP359 Actinomyces naeslundii (0%|0%|0.12%|0%|0
SP222 Actinomyces sp._HMT_169 (0%|0%|0.24%|0
SP189 Actinomyces oris (0%|0%|0.59%|0%|0%|0%
SP357 Winkia neuii (0%|0%|0%|0%|0%|0.01%|0.0
SP197 Varibaculum anthropi (0%|0%|0%|0%|0.1
SP205 Varibaculum cambriense (0%|0%|0%|0%
SP327 Actinomyces ihuae (0%|0%|0%|0.18%|0%|
SP44 Schaalia lingnae_[Not_Validly_Published] (7.92
SP196 Schaalia odontolyticus (0%|0%|0.02%|0%|0
SP98 Schaalia odontolytica (0.06%|0.00%|3.21%|0.
SP100 Schaalia sp._HMT_180 (12.38%|0.00%|1.56
SP131 Schaalia sp._HMT_172 (0%|0%|5.98%|0%|0
SP226 Fusobacterium periodonticum (0%|0%|0.85%
SP54 Fusobacterium nucleatum (0%|0%|0.09%|0%|0
SP342 Leptotrichia sp._HMT_215 (0%|0%|0.46%|0%|
SP370 Leptotrichia sp._HMT_392 (0%|0%|0.08%|0.
SP173 Faecalibacterium prausnitzii (0%|0%|0%|0%|0
SP451 Ruminococcaceae_[G-2] bacterium_HMT_08
SP365 Ruminococcaceae_[G-1] bacterium_HMT_0
SP43 Anaerostipes caccae (0%|0.14%|0%|0%|0%|0.57%
SP146 Mediterraneibacter [Ruminococcus] gnavus (0%|1.66
SP102 Mediterraneibacter [Ruminococcus] lactaris (0%|
SP344 Faecalimonas umbilicata (0%|0%|0%|0%|0%|0.23%
SP206 Oribacterium sinus (0%|0%|1.68%|0%|0%|0%
SP353 Oribacterium parvum (0%|0%|0.26%|0%|0%|0
SP81 Blautia wexlerae (0%|0%|0%|7.47%|0%|0.12%|0%
SP295 Stomatobaculum sp._HMT_097 (0%|0.00%|0.2
SP309 Peptostreptococcaceae_[XI][G-9] [Eubacterium_
SP239 Peptostreptococcaceae_[XI][G-6] [Euba
SP294 Peptostreptococcaceae_[XI][G-1] [Eubacteriu
SP96 Clostridioides difficile (0%|0.25%|0%|0%|0%|
SP137 Peptostreptococcus anaerobius (0%|0%|0%|
SP212 Peptostreptococcus stomatis (0%|0%|1.87%
SP58 Clostridium perfringens (0%|2.76%|0%|20.00
SP127 Clostridium tertium (0%|0%|0%|0.13%|0
SP125 Clostridium paraputrificum (0%|0.01%|0
SP122 Clostridium chromiireducens (0%|0%|0%
SP211 Clostridium butyricum (0%|0%|0%|1.26
SP235 Anaerococcus obesiensis (0%|0%|0%|0.21
SP126 Anaerococcus vaginalis (0%|0%|0%|5.28
SP42 Anaerococcus hydrogenalis (0%|0%|0%|6.
SP139 Anaerococcus lactolyticus (0%|0%|0%|0.4
SP111 Anaerococcus octavius (0.00%|0%|0%|0
SP266 Anaerococcus sp._HMT_294 (0%|0%|0%|0
SP106 Finegoldia magna (0%|0.08%|0.00%|20.76
SP198 Parvimonas micra (0%|0%|0.15%|0%|0%|0%
SP194 Peptoniphilus coxii (0%|0.00%|0.00%|0.78%
SP221 Peptoniphilus duerdenii (0%|0.00%|0%|2
SP138 Peptoniphilus lacrimalis (0%|0%|0%|0
SP110 Peptoniphilus grossensis (0%|0.01%|0
SP220 Peptoniphilus tyrrelliae (0%|0%|0%|0
SP66 Megamonas funiformis (0%|0%|0%|0%|0%|0%|
SP99 Dialister propionicifaciens (0%|0%|0%|0.00%
SP80 Veillonella sp._HMT_780 (4.23%|2.90%|0.02%|0
SP118 Veillonella atypica (11.82%|7.24%|0.57%|0.18
SP156 Veillonella dispar (4.32%|0.19%|3.24%|0.00%
SP329 Veillonella denticariosi (0%|0%|0%|0%|0.10%
SP2 Veillonella parvula (0%|0%|0.01%|0%|0.17%|0%
SP41 Veillonella rogosae (0%|0%|0.55%|0%|0%|0%|
SP119 Solobacterium moorei (0%|0%|0.30%|0%
SP252 Erysipelatoclostridium [Clostridium] innocu
SP130 Gemella sanguinis (0%|0%|3.29%|0%|0%|0
SP157 Gemella morbillorum (0%|0%|0.09%|0%|0
SP108 Gemella haemolysans (5.04%|11.54%|2.49
SP234 Staphylococcus simulans (0%|0%|0%|0%|
SP124 Staphylococcus saccharolyticus (0%|0%|0%|
SP56 Staphylococcus capitis (0%|0%|0%|0%|0%|0%
SP149 Staphylococcus lugdunensis (0%|0.04%|0%|
SP75 Staphylococcus haemolyticus (0%|0.02%|0%
SP63 Staphylococcus hominis (0.06%|0.13%|0.00%
SP162 Lactobacillus gasseri (0%|0.03%|0%|0.35
SP240 Limosilactobacillus mucosae (0%|0%|0%|
SP53 Enterococcus faecalis (0%|7.73%|0%|6.41%|3.
SP79 Abiotrophia defectiva (0%|0%|2.60%|0%|0%
SP217 Granulicatella elegans (1.02%|1.41%|0.14%|0
SP135 Granulicatella adiacens (3.62%|0%|8.75%|0%
SP101 Streptococcus anginosus (0%|6.11%|0.02
SP65 Streptococcus peroris (0.97%|0%|0.24%|0
SP64 Streptococcus lactarius (0%|0.02%|0.03%
SP68 Streptococcus sp._HMT_056 (2.34%|0.39%
SP203 Streptococcus sanguinis (0%|0%|0.73%|
SP22 Streptococcus sp._HMT_074 (1.65%|0.1
SP55 Streptococcus parasanguinis_clade_411
PT_Mother_ATB_Baby_No_ATB.T2_ _
OT_Mother_ATB_Baby_No_ATB.T1
PT_Mother_ATB_Baby_ATB
PT_Mother_ATB_Baby_No_ATB.T1
PT_Mother_ATB_Baby_ATB.T1
OT_Mother_ATB_Baby_No_ATB.T2
OT_Mother_ATB_Baby_No_ATB.T0
PT_Mother_ATB_Baby_ATB.T0
PT_Mother_ATB_Baby_No_ATB.T1_ _
PT_Mother_ATB_Baby_No_ATB.T0_ _
PT_Mother_ATB_Baby_No_ATB.T
PT_Mother_ATB_Baby_No_ATB.T0
PT_Mother_ATB_Baby_ATB.T2
PT_Mother_ATB_Baby_No_ATB
OT_Mother_ATB_Baby_No_ATB
substitutions/site
0
0.25
0.5
0.75