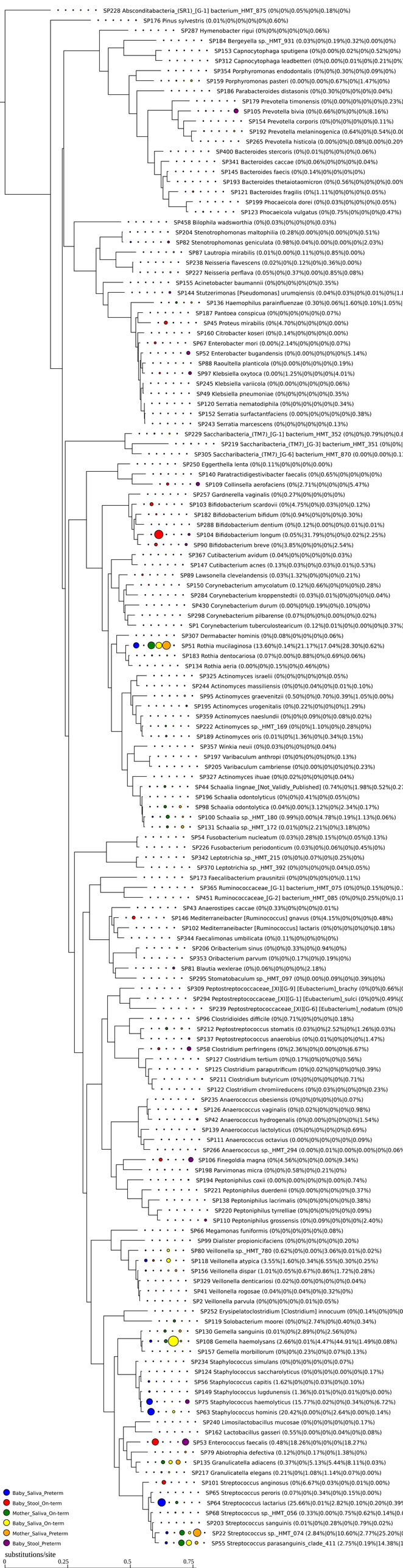
SP228 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_875 (0%|0%|0.05%|0%|0.18%|0%)
SP176 Pinus sylvestris (0.01%|0%|0%|0%|0%|0.60%)
SP287 Hymenobacter rigui (0%|0%|0%|0%|0%|0.06%)
SP184 Bergeyella sp._HMT_931 (0.03%|0%|0.19%|0.32%|0.00%|0%)
SP153 Capnocytophaga sputigena (0%|0.00%|0.02%|0%|0.52%|0%)
SP312 Capnocytophaga leadbetteri (0%|0.00%|0.01%|0%|0.21%|0%
SP354 Porphyromonas endodontalis (0%|0%|0.30%|0%|0.09%|0%)
SP159 Porphyromonas pasteri (0.00%|0.00%|0.67%|0%|1.47%|0%)
SP186 Parabacteroides distasonis (0%|0.30%|0%|0%|0%|0.04%)
SP179 Prevotella timonensis (0%|0.00%|0%|0%|0%|0.23%)
SP105 Prevotella bivia (0%|0.66%|0%|0%|0%|8.16%)
SP154 Prevotella corporis (0%|0%|0%|0%|0%|0.11%)
SP192 Prevotella melaninogenica (0.64%|0%|0.54%|0.00
SP265 Prevotella histicola (0.00%|0%|0.08%|0.00%|0.20%
SP400 Bacteroides stercoris (0%|0.01%|0%|0%|0%|0.06%)
SP341 Bacteroides caccae (0%|0.06%|0%|0%|0%|0.04%)
SP145 Bacteroides faecis (0%|0.14%|0%|0%|0%|0%)
SP193 Bacteroides thetaiotaomicron (0%|0.56%|0%|0%|0%|0.00%
SP121 Bacteroides fragilis (0%|1.11%|0%|0%|0%|0.05%)
SP199 Phocaeicola dorei (0%|0.03%|0%|0%|0%|0.05%)
SP123 Phocaeicola vulgatus (0%|0.75%|0%|0%|0%|0.47%)
SP458 Bilophila wadsworthia (0%|0.03%|0%|0%|0%|0.03%)
SP204 Stenotrophomonas maltophilia (0.28%|0.00%|0%|0.00%|0%|0.51%)
SP82 Stenotrophomonas geniculata (0.98%|0.04%|0.00%|0.00%|0%|2.03%)
SP87 Lautropia mirabilis (0.01%|0.00%|0.11%|0%|0.85%|0.00%)
SP238 Neisseria flavescens (0.02%|0%|0.12%|0%|0.36%|0.00%)
SP227 Neisseria perflava (0.05%|0%|0.37%|0.00%|0.85%|0.08%)
SP155 Acinetobacter baumannii (0%|0%|0%|0%|0%|0.35%)
SP144 Stutzerimonas [Pseudomonas] urumqiensis (0.04%|0.03%|0%|0.01%|0%|1.8
SP136 Haemophilus parainfluenzae (0.30%|0.06%|1.60%|0.10%|1.05%|
SP187 Pantoea conspicua (0%|0%|0%|0%|0%|0.07%)
SP45 Proteus mirabilis (0%|4.70%|0%|0%|0%|0.00%)
SP160 Citrobacter koseri (0%|0.14%|0%|0%|0%|0.00%)
SP67 Enterobacter mori (0.00%|2.14%|0%|0%|0%|0.07%)
SP52 Enterobacter bugandensis (0%|0.00%|0%|0%|0%|5.14%)
SP88 Raoultella planticola (0%|0.00%|0%|0%|0%|0.19%)
SP97 Klebsiella oxytoca (0.00%|1.25%|0%|0%|0%|4.01%)
SP245 Klebsiella variicola (0%|0.00%|0%|0%|0%|0.06%)
SP49 Klebsiella pneumoniae (0%|0%|0%|0%|0%|0.35%)
SP120 Serratia nematodiphila (0%|0%|0%|0%|0%|0.34%)
SP152 Serratia surfactantfaciens (0.00%|0%|0%|0%|0%|0.38%)
SP243 Serratia marcescens (0%|0%|0%|0%|0%|0.13%)
SP229 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_352 (0%|0%|0.79%|0%|0.8
SP219 Saccharibacteria_(TM7)_[G-3] bacterium_HMT_351 (0%|0%|
SP305 Saccharibacteria_(TM7)_[G-6] bacterium_HMT_870 (0.00%|0.00%|0.1
SP250 Eggerthella lenta (0%|0.11%|0%|0%|0%|0.00%)
SP140 Paratractidigestivibacter faecalis (0%|0.65%|0%|0%|0%|0%)
SP109 Collinsella aerofaciens (0%|2.71%|0%|0%|0%|5.47%)
SP257 Gardnerella vaginalis (0%|0.27%|0%|0%|0%|0%)
SP103 Bifidobacterium scardovii (0%|4.75%|0%|0.03%|0%|0.12%)
SP182 Bifidobacterium bifidum (0%|0.94%|0%|0%|0%|0.30%)
SP288 Bifidobacterium dentium (0%|0.12%|0.00%|0%|0.01%|0.01%)
SP104 Bifidobacterium longum (0.05%|31.79%|0%|0%|0.02%|2.25%)
SP90 Bifidobacterium breve (0%|3.85%|0%|0%|0%|2.54%)
SP367 Cutibacterium avidum (0.04%|0%|0%|0%|0%|0.03%)
SP147 Cutibacterium acnes (0.13%|0.03%|0%|0.03%|0.01%|0.53%)
SP89 Lawsonella clevelandensis (0.03%|1.32%|0%|0%|0%|0.21%)
SP150 Corynebacterium amycolatum (0.12%|0.66%|0%|0%|0%|0.28%)
SP284 Corynebacterium kroppenstedtii (0.03%|0.01%|0%|0%|0%|0.04%)
SP430 Corynebacterium durum (0.00%|0%|0.19%|0%|0.10%|0%)
SP298 Corynebacterium pilbarense (0.07%|0%|0%|0.00%|0%|0.02%)
SP1 Corynebacterium tuberculostearicum (0.12%|0.01%|0%|0.00%|0%|0.37%)
SP307 Dermabacter hominis (0%|0.08%|0%|0%|0%|0.06%)
SP51 Rothia mucilaginosa (13.60%|0.14%|21.17%|17.04%|28.30%|0.62%)
SP183 Rothia dentocariosa (0.07%|0.00%|0.88%|0%|0.69%|0.06%)
SP134 Rothia aeria (0.00%|0%|0.15%|0%|0.46%|0%)
SP325 Actinomyces israelii (0%|0%|0%|0%|0%|0.05%)
SP244 Actinomyces massiliensis (0%|0%|0.04%|0%|0.01%|0.10%)
SP95 Actinomyces graevenitzii (0.50%|0%|0.70%|0.39%|1.05%|0.00%)
SP195 Actinomyces urogenitalis (0%|0.22%|0%|0%|0%|1.29%)
SP359 Actinomyces naeslundii (0%|0%|0.09%|0%|0.08%|0.02%)
SP222 Actinomyces sp._HMT_169 (0%|0%|1.10%|0%|0.28%|0%)
SP189 Actinomyces oris (0.01%|0%|1.36%|0%|0.34%|0.15%)
SP357 Winkia neuii (0%|0.03%|0%|0%|0%|0.04%)
SP197 Varibaculum anthropi (0%|0%|0%|0%|0%|0.13%)
SP205 Varibaculum cambriense (0%|0.00%|0%|0%|0%|0.23%)
SP327 Actinomyces ihuae (0%|0.02%|0%|0%|0%|0.04%)
SP44 Schaalia lingnae_[Not_Validly_Published] (0.74%|0%|1.98%|0.52%|0.27
SP196 Schaalia odontolyticus (0%|0%|0.41%|0%|0.05%|0%)
SP98 Schaalia odontolytica (0.04%|0.00%|3.12%|0%|2.34%|0.17%)
SP100 Schaalia sp._HMT_180 (0.99%|0.00%|4.78%|0.19%|1.13%|0.06%)
SP131 Schaalia sp._HMT_172 (0.01%|0%|2.21%|0%|3.18%|0%)
SP54 Fusobacterium nucleatum (0.03%|0.28%|0.15%|0%|0.05%|0.13%)
SP226 Fusobacterium periodonticum (0.03%|0%|0.06%|0%|0.45%|0%)
SP342 Leptotrichia sp._HMT_215 (0%|0%|0.07%|0%|0.25%|0%)
SP370 Leptotrichia sp._HMT_392 (0%|0%|0%|0%|0.04%|0.05%)
SP173 Faecalibacterium prausnitzii (0%|0%|0%|0%|0%|0.11%)
SP365 Ruminococcaceae_[G-1] bacterium_HMT_075 (0%|0%|0.15%|0%|0.1
SP451 Ruminococcaceae_[G-2] bacterium_HMT_085 (0%|0%|0.25%|0%|0.17
SP43 Anaerostipes caccae (0%|0.33%|0%|0%|0%|0.01%)
SP146 Mediterraneibacter [Ruminococcus] gnavus (0%|4.15%|0%|0%|0%|0.48%)
SP102 Mediterraneibacter [Ruminococcus] lactaris (0%|0%|0%|0%|0%|0.18%)
SP344 Faecalimonas umbilicata (0%|0.11%|0%|0%|0%|0%)
SP206 Oribacterium sinus (0%|0%|0.33%|0%|0.94%|0%)
SP353 Oribacterium parvum (0%|0%|0.17%|0%|0.19%|0%)
SP81 Blautia wexlerae (0%|0.06%|0%|0%|0%|2.18%)
SP295 Stomatobaculum sp._HMT_097 (0%|0.00%|0.09%|0%|0.39%|0%)
SP309 Peptostreptococcaceae_[XI][G-9] [Eubacterium]_brachy (0%|0%|0.66%|0
SP294 Peptostreptococcaceae_[XI][G-1] [Eubacterium]_sulci (0%|0%|0.49%|0
SP239 Peptostreptococcaceae_[XI][G-6] [Eubacterium]_nodatum (0%|0
SP96 Clostridioides difficile (0%|0.71%|0%|0%|0%|0.18%)
SP212 Peptostreptococcus stomatis (0.03%|0%|2.52%|0%|1.26%|0.03%)
SP137 Peptostreptococcus anaerobius (0%|0.01%|0%|0%|0%|1.47%)
SP58 Clostridium perfringens (0%|2.36%|0%|0.00%|0%|6.67%)
SP127 Clostridium tertium (0%|0.17%|0%|0%|0%|0.56%)
SP125 Clostridium paraputrificum (0%|0.02%|0%|0%|0%|0.39%)
SP211 Clostridium butyricum (0%|0%|0%|0%|0%|0.71%)
SP122 Clostridium chromiireducens (0%|0.03%|0%|0%|0%|0.23%)
SP235 Anaerococcus obesiensis (0%|0%|0%|0%|0%|0.07%)
SP126 Anaerococcus vaginalis (0%|0.02%|0%|0%|0%|0.98%)
SP42 Anaerococcus hydrogenalis (0%|0.00%|0%|0%|0%|1.54%)
SP139 Anaerococcus lactolyticus (0%|0%|0%|0%|0%|0.69%)
SP111 Anaerococcus octavius (0.00%|0%|0%|0%|0%|0.09%)
SP266 Anaerococcus sp._HMT_294 (0.00%|0.01%|0.00%|0.00%|0%|0.06%
SP106 Finegoldia magna (0%|4.56%|0%|0%|0.00%|9.34%)
SP198 Parvimonas micra (0%|0%|0.58%|0%|0.21%|0%)
SP194 Peptoniphilus coxii (0.00%|0.00%|0%|0%|0.00%|0.74%)
SP221 Peptoniphilus duerdenii (0%|0.00%|0%|0%|0%|0.37%)
SP138 Peptoniphilus lacrimalis (0%|0%|0%|0%|0%|0.38%)
SP220 Peptoniphilus tyrrelliae (0%|0%|0%|0%|0%|0.09%)
SP110 Peptoniphilus grossensis (0%|0.09%|0%|0%|0%|2.40%)
SP66 Megamonas funiformis (0%|0%|0%|0%|0%|0.08%)
SP99 Dialister propionicifaciens (0%|0%|0%|0%|0%|0.20%)
SP80 Veillonella sp._HMT_780 (0.62%|0%|0.00%|3.06%|0.01%|0.02%)
SP118 Veillonella atypica (3.55%|1.60%|0.34%|6.55%|0.30%|0.25%)
SP156 Veillonella dispar (1.01%|0.05%|0.67%|0.86%|1.72%|0.28%)
SP329 Veillonella denticariosi (0.02%|0.00%|0%|0%|0%|0.04%)
SP41 Veillonella rogosae (0.04%|0%|0.04%|0%|0.32%|0%)
SP2 Veillonella parvula (0%|0%|0%|0%|0.01%|0.05%)
SP252 Erysipelatoclostridium [Clostridium] innocuum (0%|0.14%|0%|0%|0
SP119 Solobacterium moorei (0%|0%|2.74%|0%|0.40%|0.34%)
SP130 Gemella sanguinis (0.01%|0%|2.89%|0%|2.56%|0%)
SP108 Gemella haemolysans (2.66%|0.01%|4.47%|44.91%|1.49%|0.08%)
SP157 Gemella morbillorum (0%|0%|0.23%|0%|0.07%|0.13%)
SP234 Staphylococcus simulans (0%|0%|0%|0%|0%|0.07%)
SP124 Staphylococcus saccharolyticus (0%|0%|0%|0.00%|0%|0.17%)
SP56 Staphylococcus capitis (1.62%|0%|0%|0.03%|0%|0.10%)
SP149 Staphylococcus lugdunensis (1.36%|0.01%|0%|0.01%|0%|0.00%)
SP75 Staphylococcus haemolyticus (15.77%|0.02%|0%|0.34%|0%|6.72%)
SP63 Staphylococcus hominis (20.42%|0.00%|0%|2.64%|0.00%|0.14%)
SP240 Limosilactobacillus mucosae (0%|0%|0%|0%|0%|0.17%)
SP162 Lactobacillus gasseri (0.55%|0.00%|0%|0.04%|0%|0.08%)
SP53 Enterococcus faecalis (0.48%|18.26%|0%|0%|0%|18.27%)
SP79 Abiotrophia defectiva (0.12%|0%|0.17%|0%|1.38%|0%)
SP135 Granulicatella adiacens (0.37%|0%|5.13%|5.44%|8.11%|0.03%)
SP217 Granulicatella elegans (0.21%|0%|1.08%|1.14%|0.07%|0.00%)
SP101 Streptococcus anginosus (0%|6.67%|0.03%|0%|0.01%|0.00%)
SP65 Streptococcus peroris (0.07%|0%|0.34%|0%|0.15%|0.00%)
SP64 Streptococcus lactarius (25.66%|0.01%|2.82%|0.10%|0.20%|0.39%
SP68 Streptococcus sp._HMT_056 (0.33%|0.00%|0.75%|0.62%|0.14%|0.0
SP203 Streptococcus sanguinis (0.01%|0%|0.28%|0%|0.79%|0.02%)
SP22 Streptococcus sp._HMT_074 (2.84%|0%|10.60%|2.77%|25.20%|0
SP55 Streptococcus parasanguinis_clade_411 (2.75%|0.19%|14.38%|1
Baby_Saliva_Preterm
Baby_Stool_On-term
Mother_Saliva_On-term
Baby_Saliva_On-term
Mother_Saliva_Preterm
Baby_Stool_Preterm
substitutions/site
0
0.25
0.5
0.75