SP228 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_875 (0%|0%|0.05%|0%|0.18%|0%)
SP775 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_345 (0%|0%|0%|0%|0.01%|0%)
SP677 Absconditabacteria_(SR1)_[G-1] bacterium_HMT_874 (0%|0%|0.03%|0%|0%|0%)
SP176 Pinus sylvestris (0.01%|0%|0%|0%|0%|0.60%)
SP445 Triticum aestivum (0%|0.01%|0.00%|0.00%|0%|0.00%)
SP842 Gracilibacteria_(GN02)_[G-2] bacterium_HMT_873 (0%|0.00%|0%|0%|0%
SP615 Saccharibacteria_(TM7)_[G-5] bacterium_HMT_356 (0%|0%|0%|0.04%|0%|0.01%
SP765 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_350 (0%|0%|0%|0%|0%|0%)
SP305 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_870 (0.00%|0%|0%|0.13%|0.14%
SP462 Saccharibacteria_(TM7)_[G-4] bacterium_HMT_355 (0%|0%|0%|0.04%|0%|0%)
SP219 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_351 (0%|0%|0.63%|0%|0
SP704 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_348 (0%|0%|0%|0%|0%|0%)
SP229 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_352 (0%|0%|0.79%|0%|0.86%|0
SP639 Saccharibacteria_(TM7)_[G-8] bacterium_HMT_955 (0%|0%|0%|0%|0.00%|0
SP665 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_349 (0%|0%|0.02%|0%|0.01%|
SP407 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_346 (0%|0%|0.13%|0%|0.02%|
SP529 Saccharibacteria_(TM7)_[G-1] bacterium_HMT_347 (0%|0%|0.07%|0%|0.00%|
SP287 Hymenobacter rigui (0%|0%|0%|0%|0%|0.06%)
SP818 Sphingobacterium corticibacter (0.00%|0%|0%|0%|0%|0%)
SP460 Bergeyella sp._HMT_206 (0.03%|0%|0%|0.00%|0.04%|0%)
SP574 Cloacibacterium haliotis (0.00%|0%|0%|0%|0%|0%)
SP774 Epilithonimonas bovis (0.00%|0%|0%|0%|0%|0%)
SP184 Bergeyella sp._HMT_931 (0.03%|0%|0.19%|0.32%|0.00%|0%)
SP791 Bergeyella sp._HMT_900 (0%|0%|0%|0.01%|0%|0%)
SP332 Bergeyella sp._HMT_322 (0.02%|0.00%|0.06%|0.04%|0%|0%)
SP446 Capnocytophaga granulosa (0.02%|0%|0%|0.02%|0%|0%)
SP476 Capnocytophaga gingivalis (0%|0%|0%|0%|0.03%|0%)
SP153 Capnocytophaga sputigena (0%|0%|0%|0.02%|0.52%|0%)
SP312 Capnocytophaga leadbetteri (0%|0%|0.01%|0%|0.21%|0%)
SP638 Capnocytophaga sp._HMT_412 (0%|0%|0%|0%|0.01%|0%)
SP640 Capnocytophaga sp._HMT_864 (0.01%|0%|0%|0%|0%|0%)
SP837 Capnocytophaga sp._HMT_902 (0%|0%|0%|0.01%|0%|0%)
SP571 Bacteroidetes_[G-5] bacterium_HMT_511 (0%|0%|0%|0.00%|0%|0
SP559 Bacteroidetes_[G-5] bacterium_HMT_507 (0%|0%|0%|0%|0.00%|
SP811 Alistipes shahii (0%|0%|0%|0%|0%|0.00%)
SP656 Bacteroidetes_[G-3] bacterium_HMT_280 (0%|0%|0.01%|0%|0.00
SP822 Bacteroidetes_[G-3] bacterium_HMT_503 (0%|0%|0%|0%|0.00%|0
SP799 Bacteroidetes_[G-3] bacterium_HMT_365 (0%|0%|0%|0%|0%|0%)
SP528 Bacteroidetes_[G-3] bacterium_HMT_281 (0%|0%|0%|0%|0.01%|0
SP377 Bacteroidales_[G-2] bacterium_HMT_274 (0%|0%|0.21%|0%|0.01%|0
SP520 Parabacteroides merdae (0%|0%|0.03%|0%|0%|0%)
SP186 Parabacteroides distasonis (0.30%|0%|0%|0%|0%|0.04%)
SP351 Tannerella forsythia (0%|0%|0.19%|0%|0.03%|0.01%)
SP576 Tannerella sp._HMT_808 (0%|0%|0.05%|0%|0.00%|0%)
SP699 Tannerella sp._HMT_286 (0%|0%|0.00%|0%|0.05%|0%)
SP731 Tannerella serpentiformis (0%|0%|0.01%|0%|0.00%|0%)
SP682 Porphyromonas gingivalis (0.00%|0%|0.02%|0%|0.00%|0%)
SP159 Porphyromonas pasteri (0.00%|0%|0.67%|0%|1.47%|0%)
SP419 Porphyromonas sp._HMT_278 (0%|0%|0.03%|0%|0.03%|0%)
SP487 Porphyromonas bennonis (0%|0%|0%|0%|0%|0%)
SP485 Porphyromonas uenonis (0%|0%|0%|0%|0%|0.02%)
SP354 Porphyromonas endodontalis (0%|0%|0.30%|0%|0.09%|0%)
SP755 Odoribacter splanchnicus (0%|0%|0%|0%|0%|0%)
SP400 Bacteroides stercoris (0%|0%|0%|0%|0%|0.06%)
SP376 Bacteroides uniformis (0%|0%|0.03%|0%|0%|0.03%)
SP123 Phocaeicola vulgatus (0%|0%|0.75%|0%|0%|0.47%)
SP199 Phocaeicola dorei (0%|0%|0.03%|0%|0%|0.05%)
SP533 Bacteroides cellulosilyticus (0%|0%|0%|0%|0%|0%)
SP121 Bacteroides fragilis (0%|0%|1.11%|0%|0%|0.05%)
SP193 Bacteroides thetaiotaomicron (0%|0%|0.56%|0%|0%|0.00%)
SP145 Bacteroides faecis (0%|0%|0.14%|0%|0%|0%)
SP331 Bacteroides ovatus (0%|0%|0.05%|0%|0%|0.02%)
SP567 Bacteroides xylanisolvens (0%|0%|0.00%|0%|0%|0.01%)
SP341 Bacteroides caccae (0%|0%|0.06%|0%|0%|0.04%)
SP569 Alloprevotella tannerae (0%|0%|0%|0%|0.01%|0%)
SP763 Alloprevotella rava (0%|0%|0%|0.00%|0%|0%)
SP802 Alloprevotella sp._HMT_912 (0%|0%|0%|0.00%|0%|0%)
SP438 Alloprevotella sp._HMT_473 (0%|0%|0.01%|0%|0.00%|0.00%)
SP784 Alloprevotella sp._HMT_308 (0.00%|0%|0.01%|0%|0.00%|0%)
SP483 Alloprevotella sp._HMT_914 (0.01%|0%|0%|0%|0.03%|0%)
SP847 Prevotella sp._HMT_304 (0%|0%|0.01%|0%|0%|0%)
SP669 Prevotella nanceiensis (0%|0%|0%|0%|0.02%|0%)
SP680 Prevotella micans (0%|0%|0%|0%|0.00%|0%)
SP179 Prevotella timonensis (0%|0%|0%|0%|0%|0.23%)
SP759 Prevotella buccalis (0%|0%|0%|0%|0%|0.01%)
SP698 Prevotella sp._HMT_301 (0%|0%|0.00%|0%|0%|0%)
SP453 Prevotella shahii (0%|0%|0%|0%|0.10%|0%)
SP808 Prevotella sp._HMT_472 (0%|0%|0%|0%|0.00%|0%)
SP807 Prevotella loescheii (0%|0%|0%|0%|0.00%|0%)
SP841 Prevotella sp._HMT_315 (0%|0%|0%|0%|0.00%|0%)
SP803 Prevotella baroniae (0%|0%|0%|0%|0.00%|0%)
SP705 Prevotella buccae (0%|0%|0.00%|0%|0%|0%)
SP645 Prevotella colorans (0%|0%|0%|0%|0.00%|0%)
SP688 Prevotella sp._HMT_305 (0%|0%|0.01%|0%|0.00%|0%)
SP821 Prevotella sp._HMT_300 (0%|0%|0%|0%|0.00%|0%)
SP813 Prevotella sp._HMT_292 (0%|0%|0%|0%|0.00%|0%
SP105 Prevotella bivia (0%|0%|0.66%|0%|0%|8.16%)
SP814 Prevotella sp._HMT_309 (0%|0%|0%|0%|0.00%|0%)
SP516 Prevotella oris (0%|0%|0.01%|0%|0.03%|0%)
SP605 Prevotella salivae (0.01%|0%|0.02%|0%|0.00%|0%)
SP737 Prevotella copri (0%|0%|0.00%|0%|0%|0%)
SP154 Prevotella corporis (0%|0%|0%|0%|0%|0.11%)
SP578 Prevotella disiens (0%|0%|0.01%|0%|0%|0.01%)
SP563 Prevotella pallens (0%|0%|0.01%|0%|0.03%|0%)
SP826 Prevotella nigrescens (0%|0%|0%|0%|0.01%|0%)
SP404 Prevotella intermedia (0%|0%|0%|0.01%|0.00%|
SP505 Prevotella multiformis (0%|0%|0%|0%|0.01%|0%)
SP401 Prevotella denticola (0%|0%|0.00%|0%|0.00%|0%)
SP265 Prevotella histicola (0.00%|0%|0.08%|0%|0.20%|0%
SP192 Prevotella melaninogenica (0.64%|0%|0.54%|0%|0.00%|1
SP352 Prevotella sp._HMT_306 (0%|0%|0.22%|0%|0%|0%)
SP777 Prevotella vespertina (0%|0%|0%|0%|0%|0%)
SP246 Campylobacter ureolyticus (0.04%|0%|0%|0%|0%|0.00%)
SP715 Campylobacter hominis (0%|0%|0%|0%|0%|0.00%)
SP340 Campylobacter concisus (0.00%|0%|0.08%|0%|0.04%|0%)
SP465 Campylobacter sp._HMT_044 (0%|0%|0%|0%|0.03%|0%)
SP629 Campylobacter gracilis (0%|0%|0.01%|0%|0.00%|0%)
SP480 Desulfobulbus sp._HMT_041 (0%|0%|0%|0%|0.01%|0%)
SP458 Bilophila wadsworthia (0%|0%|0.03%|0%|0%|0.03%)
SP786 Tistrella mobilis (0%|0%|0%|0%|0.00%|0%)
SP734 Methylobacterium nonmethylotrophicum (0%|0%|0%|0%|0%|0.00%)
SP304 Methylobacterium aerolatum (0%|0%|0%|0%|0%|0.03%)
SP736 Sphingomonas yunnanensis (0%|0%|0%|0%|0%|0.00%)
SP829 Pannonibacter indicus (0.00%|0%|0%|0%|0%|0%)
SP691 Polymorphum gilvum (0%|0.00%|0%|0%|0%|0%)
SP536 Amaricoccus kaplicensis (0.01%|0%|0%|0%|0%|0%)
SP449 Devosia yakushimensis (0.01%|0.01%|0%|0.01%|0.02%|0%)
SP788 Rhizobium rosettiformans (0%|0%|0%|0.01%|0%|0%)
SP756 Agrobacterium fabrum (0%|0%|0%|0.01%|0%|0%)
SP434 Cardiobacterium hominis (0.01%|0%|0.01%|0%|0.02%|0%)
SP321 Cardiobacterium valvarum (0%|0%|0%|0%|0.00%|0%)
SP787 Pseudoxanthomonas taiwanensis (0%|0%|0%|0%|0%|0.00%)
SP82 Stenotrophomonas geniculata (0.98%|0.04%|0.00%|0%|0%|2.03%)
SP204 Stenotrophomonas maltophilia (0.28%|0.00%|0%|0.00%|0%|0.51%)
SP535 Neisseriaceae_[G-1] bacterium_HMT_327 (0%|0%|0%|0%|0%|0.02%)
SP161 Neisseriaceae_[G-1] bacterium_HMT_174 (0.01%|0.00%|0%|0%|0%|0.
SP667 Kingella oralis (0%|0%|0.00%|0%|0.01%|0%)
SP310 Neisseria oralis (0%|0%|0%|0%|0.07%|0%)
SP555 Neisseria elongata (0%|0%|0.01%|0%|0.00%|0%)
SP730 Neisseria subflava (0%|0%|0%|0%|0%|0.00%)
SP227 Neisseria perflava (0.05%|0%|0.37%|0.00%|0.85%|0.08%)
SP455 Neisseria cinerea (0.01%|0%|0%|0.00%|0%|0%)
SP238 Neisseria flavescens (0.02%|0%|0.12%|0%|0.36%|0.00%)
SP758 Hydrogenophilus thermoluteolus (0%|0%|0%|0%|0%|0.00%)
SP764 Schlegelella aquatica (0%|0%|0%|0%|0%|0.00%)
SP760 Leptothrix sp._HMT_266 (0.00%|0%|0%|0%|0%|0%)
SP328 Hydrogenophaga aquatica (0.00%|0%|0%|0%|0%|0.05%)
SP477 Delftia acidovorans (0.01%|0%|0%|0%|0%|0%)
SP748 Ottowia sp._HMT_894 (0%|0%|0%|0.00%|0%|0%)
SP87 Lautropia mirabilis (0.01%|0%|0.11%|0%|0.85%|0.00%)
SP654 Sutterella stercoricanis (0%|0%|0.01%|0%|0%|0%)
SP579 Sutterella timonensis (0%|0%|0%|0%|0%|0.01%)
SP286 Castellaniella defragrans (0%|0%|0%|0.01%|0%|0%)
SP355 Bordetella petrii (0%|0%|0%|0%|0%|0.03%)
SP812 Noviherbaspirillum aurantiacum (0%|0%|0%|0%|0%|0%)
SP776 Noviherbaspirillum suwonense (0.00%|0%|0%|0%|0.00%|0%)
SP790 Massilia aurea (0%|0%|0%|0%|0.00%|0%)
SP820 Massilia arvi (0%|0%|0%|0%|0%|0%)
SP322 Massilia pinisoli (0%|0%|0%|0%|0%|0.02%)
SP742 Pseudomonas hutmensis (0%|0%|0%|0%|0%|0.00%)
SP679 Pseudomonas koreensis (0%|0.00%|0%|0%|0%|0%)
SP343 Pseudomonas sihuiensis (0.00%|0%|0%|0%|0%|0.03%)
SP683 Pseudomonas aeruginosa (0.01%|0%|0%|0%|0%|0%)
SP539 Pseudomonas matsuisoli (0.00%|0%|0%|0%|0%|0%)
SP144 Stutzerimonas [Pseudomonas] urumqiensis (0.04%|0.03%|0%|0.01%|0.01%|1
SP561 Moraxella osloensis (0.01%|0.00%|0%|0%|0%|0%)
SP467 Acinetobacter johnsonii (0.00%|0%|0%|0%|0%|0.01%)
SP155 Acinetobacter baumannii (0.01%|0%|0%|0%|0%|0.35%)
SP603 Acinetobacter ursingii (0.00%|0%|0%|0%|0%|0%)
SP548 Marinomonas arctica (0.01%|0%|0%|0%|0%|0%)
SP701 Vibrio litoralis (0.00%|0%|0%|0%|0%|0%)
SP767 Haemophilus sputorum (0%|0%|0.01%|0%|0%|0%)
SP136 Haemophilus parainfluenzae (0.30%|0.06%|1.60%|0.10%|1.05%|0
SP631 Haemophilus pittmaniae (0.00%|0%|0.03%|0%|0.01%)
SP566 Haemophilus parahaemolyticus (0%|0%|0.14%|0%|0%|0%)
SP658 Aggregatibacter sp._HMT_898 (0%|0%|0.00%|0%|0%|0%)
SP564 Aggregatibacter actinomycetemcomitans (0%|0%|0.00%|0%
SP782 Aggregatibacter aphrophilus (0%|0%|0%|0.00%|0%|0%)
SP530 Aggregatibacter sp._HMT_513 (0%|0%|0%|0%|0.03%|0%)
SP362 Aggregatibacter sp._HMT_458 (0%|0.03%|0.02%|0.02%|0%)
SP281 Haemophilus haemolyticus (0%|0%|0.00%|0%|0.02%|0%)
SP241 Haemophilus sp._HMT_908 (0%|0%|0%|0.00%|0%|0%)
SP596 Morganella morganii (0%|0%|0%|0.01%|0%|0%)
SP45 Proteus mirabilis (0%|4.70%|0%|0%|0%|0%)
SP323 Proteus columbae (0%|0.01%|0%|0%|0%|0%)
SP457 Proteus alimentorum (0%|0%|0%|0%|0%|0.02%)
SP282 Proteus vulgaris (0%|0.00%|0%|0%|0%|0%)
SP527 Serratia quinivorans (0%|0%|0%|0%|0%|0%)
SP120 Serratia nematodiphila (0%|0%|0%|0%|0%|0.34%)
SP243 Serratia marcescens (0%|0%|0%|0%|0%|0.13%)
SP152 Serratia surfactantfaciens (0.00%|0%|0%|0%|0%|0.38%)
SP435 Cedecea lapagei (0%|0.00%|0%|0%|0%|0%)
SP67 Enterobacter mori (0.00%|2.14%|0%|0%|0%|0.07%)
SP261 Klebsiella aerogenes (0%|0.01%|0%|0%|0%|0%)
SP88 Raoultella planticola (0%|0.00%|0%|0%|0%|0.19%)
SP50 Klebsiella quasipneumoniae (0%|0%|0%|0%|0%|0%)
SP49 Klebsiella pneumoniae (0%|0%|0%|0%|0%|0.35%)
SP245 Klebsiella variicola (0%|0.00%|0%|0%|0%|0.06%)
SP52 Enterobacter bugandensis (0%|0.00%|0%|0%|0%|5.14%)
SP97 Klebsiella oxytoca (0.00%|1.25%|0%|0%|0%|4.01%)
SP816 Citrobacter europaeus (0%|0%|0%|0%|0%|0%)
SP437 Citrobacter freundii (0%|0%|0%|0%|0%|0.00%)
SP381 Kosakonia sacchari (0%|0.01%|0%|0%|0%|0%)
SP838 Pluralibacter gergoviae (0%|0.00%|0%|0%|0%|0%)
SP187 Pantoea conspicua (0%|0%|0%|0%|0%|0.07%)
SP399 Pantoea septica (0%|0.00%|0%|0%|0%|0%)
SP380 Mixta theicola (0%|0%|0%|0%|0%|0.01%)
SP160 Citrobacter koseri (0%|0.14%|0%|0%|0%|0.00%)
SP230 Salmonella bongori (0%|0.01%|0%|0%|0%|0.04%)
SP171 Pantoea dispersa (0%|0%|0%|0%|0%|0.01%)
SP609 Pantoea wallisii (0%|0%|0%|0%|0%|0%)
SP356 Erwinia iniecta (0%|0%|0%|0%|0%|0.01%)
SP185 Erwinia toletana (0%|0%|0%|0%|0%|0.00%)
SP762 Thermus thermophilus (0%|0%|0%|0%|0%|0.00%)
SP396 Gordonibacter pamelaeae (0%|0%|0.00%|0%|0%|0%)
SP250 Eggerthella lenta (0%|0.11%|0%|0%|0%|0.00%)
SP614 Cryptobacterium curtum (0%|0%|0.01%|0%|0%|0%)
SP594 Slackia exigua (0%|0%|0.02%|0%|0.01%|0%)
SP109 Collinsella aerofaciens (0%|0%|2.71%|0%|0%|5.47%)
SP140 Paratractidigestivibacter faecalis (0%|0%|0.65%|0%|0%|0%)
SP652 Parolsenella massiliensis (0%|0.01%|0%|0%|0%|0%)
SP541 Olsenella sp._HMT_807 (0%|0%|0.03%|0%|0.04%|0%)
SP382 Atopobium minutum (0%|0%|0%|0%|0%|0.04%)
SP582 Olsenella uli (0%|0%|0.01%|0%|0.04%|0%)
SP600 Alloscardovia omnicolens (0%|0%|0%|0%|0.03%|0%)
SP448 Parascardovia denticolens (0%|0%|0%|0%|0.01%|0%)
SP664 Scardovia wiggsiae (0%|0%|0.02%|0%|0.02%|0%)
SP257 Gardnerella vaginalis (0%|0%|2.7%|0%|0%|0.57%)
SP374 Gardnerella swidsinskii (0%|0%|0%|0%|0%|0.04%)
SP103 Bifidobacterium scardovii (0%|0%|4.75%|0%|0.03%|0.12%)
SP90 Bifidobacterium breve (0%|3.85%|0%|0%|0%|2.54%)
SP104 Bifidobacterium longum (0.05%|31.79%|0%|0%|0.02%|2.25%)
SP288 Bifidobacterium dentium (0%|0.12%|0.00%|0%|0.01%|0.01%)
SP182 Bifidobacterium bifidum (0%|0.94%|0%|0%|0%|0.30%)
SP267 Bifidobacterium apri (0%|0%|0%|0%|0%|0.00%)
SP655 Actinotignum schaalii (0%|0%|0.01%|0%|0%|0%)
SP218 Peptidiphaga gingivicola (0%|0%|0.01%|0%|0.01%|0%)
SP225 Peptidiphaga sp._HMT_183 (0.02%|0%|0.02%|0%|0.05%|0%)
SP357 Winkia neuii (0%|0%|0.03%|0%|0%|0.04%)
SP205 Varibaculum cambriense (0%|0%|0.00%|0%|0%|0.23%)
SP197 Varibaculum anthropi (0%|0%|0%|0%|0%|0.13%)
SP327 Actinomyces ihuae (0%|0%|0.02%|0%|0%|0.04%)
SP604 Schaalia sp._HMT_178 (0.01%|0%|0.03%|0%|0.02%|0%)
SP402 Schaalia meyeri (0%|0%|0.00%|0%|0%|0%)
SP425 Schaalia turicensis (0%|0%|0%|0%|0.01%|0%)
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%)
SP98 Schaalia odontolytica (0.04%|0.00%|3.12%|0%|2.34%|0.17%)
SP196 Schaalia odontolytica (0%|0%|0.41%|0%|0.05%|0%)
SP44 Schaalia lingnae_[Not_Validly_Published] (0.74%|0%|1.98%|0.52%|0.27
SP95 Actinomyces graevenitzii (0%|0%|0.39%|0%|0.05%|0.00%)
SP583 Actinomyces sp._HMT_448 (0%|0%|0.03%|0%|0.01%|0.01%)
SP733 Actinomyces dentalis (0%|0%|0%|0%|0.00%|0%)
SP557 Actinomyces sp._HMT_897 (0%|0.00%|0%|0%|0%|0%)
SP244 Actinomyces massiliensis (0%|0%|0.04%|0%|0.01%|0.10%)
SP325 Actinomyces israelii (0%|0%|0%|0%|0.05%|0%)
SP347 Actinomyces gerencseriae (0.02%|0%|0.04%|0%|0.01%|0.02%)
SP195 Actinomyces urogenitalis (0%|0%|0.22%|0%|0%|1.29%)
SP666 Actinomyces timonensis (0%|0%|0.01%|0%|0.01%|0%)
SP696 Actinomyces bowdenii (0%|0%|0%|0%|0.00%|0%)
SP757 Actinomyces johnsonii (0%|0%|0%|0%|0.00%|0%)
SP443 Actinomyces sp._HMT_170 (0%|0%|0.03%|0%|0.03%|0%)
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%)
SP270 Actinomyces sp._HMT_175 (0%|0%|0.05%|0%|0.04%|0%)
SP702 Arthrobacter agilis (0.01%|0%|0%|0%|0%|0%)
SP766 Pseudoglutamicibacter cumminsii (0.00%|0%|0%|0%|0%|0%)
SP694 Rothia kristinae (0%|0%|0%|0%|0%|0.00%)
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%)
SP633 Curtobacterium oceanosedimentum (0%|0%|0%|0%|0%|0.01%)
SP817 Microbacterium saccharophilum (0%|0%|0%|0%|0%|0.00%)
SP729 Micrococcus hominis (0.00%|0%|0%|0%|0%|0%)
SP753 Pedococcus aerophilus (0.00%|0%|0%|0%|0%|0%)
SP846 Phycicoccus jejuensis (0%|0%|0%|0%|0%|0.00%)
SP825 Brachybacterium squillarum (0%|0.01%|0%|0%|0%|0%)
SP307 Dermabacter hominis (0%|0.08%|0%|0%|0%|0.06%)
SP433 Dermabacter jinjuensis (0%|0%|0%|0%|0%|0.00%)
SP518 Nakamurella multipartita (0%|0%|0%|0.00%|0%|0%)
SP447 Nocardioides marinus (0%|0%|0%|0%|0%|0.02%)
SP608 Marmoricola scoriae (0%|0%|0%|0%|0%|0.00%)
SP844 Propionibacteriaceae_[G-1] bacterium_HMT_915 (0%|0%|0%|0%|0.00%
SP651 Propionibacterium acidifaciens (0%|0%|0.01%|0%|0.01%|0%)
SP738 Arachnia rubra (0%|0%|0%|0%|0.00%|0%)
SP358 Arachnia propionica (0.01%|0%|0.03%|0%|0.01%|0%)
SP819 Cutibacterium granulosum (0%|0%|0%|0%|0%|0.00%)
SP367 Cutibacterium avidum (0.04%|0%|0%|0%|0%|0.00%)
SP147 Cutibacterium acnes (0.13%|0.03%|0%|0.03%|0.01%|0.53%)
SP703 Gordonia amarae (0.01%|0%|0%|0%|0%|0%)
SP823 Mycolicibacterium obuense (0%|0%|0.00%|0%|0%|0%)
SP624 Mycolicibacterium iranicum (0%|0%|0%|0%|0.00%|0%)
SP89 Lawsonella clevelandensis (0.03%|1.32%|0%|0%|0%|0.21%)
SP284 Corynebacterium kroppenstedtii (0.03%|0%|0%|0.02%|0%|0.04%)
SP387 Corynebacterium sphenisci (0%|0%|0%|0%|0%|0%)
SP150 Corynebacterium amycolatum (0.12%|0.66%|0%|0%|0%|0.28%)
SP556 Corynebacterium falsenii (0.01%|0%|0%|0%|0%|0%)
SP441 Corynebacterium epidermidicanis (0%|0%|0%|0%|0%|0%)
SP597 Corynebacterium matruchotii (0%|0%|0.00%|0%|0.02%|0%)
SP728 Corynebacterium appendicis (0%|0%|0%|0%|0%|0.00%)
SP298 Corynebacterium pilbarense (0.07%|0%|0%|0.00%|0.02%|0%)
SP330 Corynebacterium pyruviciproducens (0%|0%|0%|0%|0%|0.03%)
SP796 Corynebacterium frankenforstense (0%|0%|0.00%|0%|0%|0%)
SP430 Corynebacterium durum (0%|0%|0.01%|0.19%|0.10%|0%)
SP481 Corynebacterium simulans (0%|0%|0%|0%|0%|0.00%)
SP632 Corynebacterium singulare (0%|0.01%|0%|0%|0%|0%)
SP506 Corynebacterium propinquum (0%|0%|0%|0%|0%|0.00%)
SP456 Corynebacterium macginleyi (0.02%|0%|0%|0%|0%|0%)
SP1 Corynebacterium tuberculostearicum (0.12%|0.01%|0%|0.00%|0%|0.37%)
SP442 Anaerolineae_[G-1] bacterium_HMT_439 (0%|0%|0.02%|0%|0.00%|0%)
SP751 Akkermansia muciniphila (0%|0%|0.00%|0%|0%|0%)
SP659 Fretibacterium fastidiosum (0%|0%|0.00%|0%|0.01%|0%)
SP700 Fretibacterium sp._HMT_359 (0%|0%|0.03%|0%|0.00%|0%)
SP626 Fretibacterium sp._HMT_362 (0%|0%|0.00%|0%|0%|0%)
SP317 Fretibacterium sp._HMT_360 (0%|0%|0.01%|0%|0.00%|0%)
SP570 Treponema maltophilum (0%|0%|0.01%|0%|0%|0%)
SP792 Treponema sp._HMT_258 (0%|0%|0.00%|0%|0%|0%)
SP735 Treponema sp._HMT_270 (0%|0%|0%|0%|0.00%|0%)
SP450 Treponema socranskii (0%|0%|0.02%|0%|0.01%|0%)
SP521 Treponema amylovorum (0%|0%|0%|0%|0.00%|0%)
SP747 Treponema sp._HMT_264 (0%|0%|0%|0%|0.00%|0%)
SP560 Treponema sp._HMT_238 (0%|0%|0.01%|0%|0%|0%)
SP427 Treponema sp._HMT_237 (0.01%|0%|0.00%|0%|0%|0.01%)
SP684 Treponema sp._HMT_231 (0%|0%|0.01%|0%|0.00%|0%)
SP721 Treponema vincentii (0%|0%|0.00%|0%|0%|0%)
SP413 Treponema sp._HMT_249 (0%|0%|0.00%|0%|0.00%|0%)
SP540 Treponema denticola (0%|0%|0.06%|0%|0.00%|0%)
SP534 Treponema sp._HMT_508 (0%|0%|0%|0%|0.00%|0%)
SP746 Treponema sp._HMT_518 (0%|0%|0%|0%|0.00%|0%)
SP466 Cetobacterium somerae (0%|0%|0%|0%|0%|0.02%)
SP749 Fusobacterium sp._HMT_248 (0%|0%|0.00%|0%|0%|0%)
SP226 Fusobacterium periodonticum (0.03%|0%|0.06%|0%|0.45%|0%)
SP472 Fusobacterium hwasookii (0%|0%|0%|0%|0.02%|0%)
SP54 Fusobacterium nucleatum (0.03%|0%|0.28%|0%|0.15%|0.13%)
SP158 Fusobacterium nucleatum_subsp._vincentii (0%|0%|0.05%|0%|0.02%|0%
SP303 Fusobacterium sp._HMT_203 (0%|0%|0%|0%|0.04%|0%)
SP339 Leptotrichia sp._HMT_221 (0.01%|0%|0.00%|0%|0.06%|0%)
SP642 Leptotrichia sp._HMT_219 (0%|0%|0%|0%|0.00%|0%)
SP370 Leptotrichia sp._HMT_392 (0%|0%|0.00%|0%|0.04%|0.05%)
SP342 Leptotrichia sp._HMT_215 (0%|0%|0.07%|0%|0.25%|0%)
SP833 Leptotrichia sp._HMT_212 (0%|0%|0%|0%|0.00%|0%)
SP549 Leptotrichia sp._HMT_218 (0%|0%|0.04%|0%|0.01%|0%)
SP739 Leptotrichia sp._HMT_498 (0%|0%|0%|0%|0.00%|0%)
SP565 Leptotrichia sp._HMT_225 (0.01%|0%|0.00%|0%|0.01%|0%)
SP554 Leptotrichia hofstadii (0%|0%|0%|0%|0.00%|0%)
SP444 Leptotrichia hongkongensis (0%|0%|0.01%|0%|0.06%|0%)
SP815 Leptotrichia sp._HMT_417 (0%|0%|0.02%|0%|0.01%|0%)
SP713 Leptotrichia wadei (0%|0%|0.00%|0%|0.01%|0%)
SP365 Ruminococcaceae_[G-1] bacterium_HMT_075 (0%|0%|0.15%|0%|0.16%|0.
SP451 Ruminococcaceae_[G-2] bacterium_HMT_085 (0%|0%|0.25%|0%|0.17%|0
SP411 Flavonifractor plautii (0%|0%|0%|0%|0%|0.02%)
SP216 Ruthenibacterium lactatiformans (0%|0%|0%|0%|0%|0.02%)
SP845 Subdoligranulum variabile (0%|0%|0.00%|0%|0%|0%)
SP173 Faecalibacterium prausnitzii (0%|0%|0.00%|0%|0%|0.11%)
SP695 Lachnospiraceae_[G-8] bacterium_HMT_500 (0%|0%|0.01%|0%|0.02%|0%)
SP517 Catonella morbi (0%|0%|0.03%|0%|0.02%|0%)
SP789 Butyrivibrio sp._HMT_080 (0%|0%|0.01%|0%|0%|0%)
SP552 Butyrivibrio sp._HMT_455 (0%|0%|0.01%|0%|0.02%|0%)
SP302 Shuttleworthia satelles (0%|0%|0.05%|0%|0.00%|0%)
SP602 Lachnospiraceae_[G-7] bacterium_HMT_163 (0%|0%|0.00%|0%|0%|0%)
SP773 Lachnospiraceae_[G-7] bacterium_HMT_086 (0%|0%|0.00%|0%|0.00%|0%)
SP575 Lachnospiraceae_[G-2] bacterium_HMT_096 (0%|0%|0.05%|0%|0.01%|0%)
SP251 Lachnospiraceae_[G-3] bacterium_HMT_100 (0%|0%|0.02%|0%|0.02%|0%
SP685 Johnsonella sp._HMT_166 (0%|0%|0.00%|0%|0.00%|0%)
SP595 Lachnoanaerobaculum sp._HMT_496 (0%|0%|0.00%|0%|0.01%|0%)
SP621 Lachnoanaerobaculum sp._HMT_089 (0%|0%|0.03%|0%|0.00%|0%)
SP490 Lachnoanaerobaculum orale (0%|0%|0.09%|0%|0.05%|0%)
SP754 Lachnoanaerobaculum saburreum (0%|0%|0.00%|0%|0.00%|0%)
SP834 Moryella indoligenes (0%|0%|0%|0%|0.00%|0%)
SP436 Stomatobaculum longum (0%|0%|0.01%|0%|0.08%|0%)
SP295 Stomatobaculum sp._HMT_097 (0%|0%|0.09%|0%|0.39%|0%)
SP804 Oribacterium sp._HMT_102 (0%|0%|0.00%|0%|0.00%|0%)
SP646 Oribacterium sp._HMT_078 (0%|0%|0.01%|0%|0.03%|0%)
SP206 Oribacterium sinus (0%|0%|0.33%|0%|0.94%|0%)
SP353 Oribacterium parvum (0%|0%|0.17%|0%|0.19%|0%)
SP403 Oribacterium asaccharolyticum (0%|0%|0.15%|0%|0.01%|0%)
SP628 Roseburia inulinivorans (0%|0%|0.00%|0%|0%|0%)
SP778 Roseburia faecis (0%|0%|0%|0%|0%|0%)
SP81 Blautia wexlerae (0%|0.06%|0%|0%|0%|2.18%)
SP716 Enterocloster aldenensis (0%|0%|0%|0%|0%|0.00%)
SP209 Hungatella hathewayi (0%|0%|0%|0%|0%|0.00%)
SP344 Faecalimonas umbilicata (0%|0%|0.11%|0%|0%|0%)
SP783 Tyzzerella nexilis (0%|0%|0%|0%|0.00%|0%)
SP43 Anaerostipes caccae (0%|0%|0.33%|0%|0%|0.01%)
SP522 Dorea longicatena (0.01%|0%|0%|0%|0%|0.00%)
SP146 Mediterraneibacter [Ruminococcus] gnavus (0%|0%|4.15%|0%|0%|0.48%)
SP102 Mediterraneibacter [Ruminococcus] lactaris (0%|0%|0%|0%|0%|0.18%)
SP368 Mediterraneibacter faecis (0%|0%|0%|0%|0%|0.01%)
SP714 Mediterraneibacter [Ruminococcus] torques (0%|0%|0%|0%|0%|0.01%)
SP429 Clostridiales_[F-1]_[G-1] bacterium_HMT_093 (0%|0%|0.00%|0%|0.01%
SP366 Clostridium tarantellae (0%|0%|0.01%|0%|0%|0%)
SP127 Clostridium tertium (0%|0%|0.17%|0%|0%|0.56%)
SP58 Clostridium perfringens (0%|0%|2.36%|0%|0%|6.67%)
SP627 Clostridium baratii (0%|0%|0.02%|0%|0%|0.01%)
SP125 Clostridium paraputrificum (0%|0%|0.02%|0%|0%|0.39%)
SP122 Clostridium chromiireducens (0%|0%|0.03%|0%|0%|0.23%)
SP211 Clostridium butyricum (0%|0%|0.06%|0%|0%|0.71%)
SP625 Clostridium saccharobutylicum (0%|0%|0.00%|0%|0%|0%)
SP637 Eubacterium limosum (0%|0%|0%|0%|0%|0.00%)
SP843 Pseudoramibacter alactolyticus (0%|0%|0%|0%|0.01%|0%)
SP369 Filifactor alocis (0%|0%|0.08%|0%|0.04%|0%)
SP137 Peptostreptococcus anaerobius (0%|0%|0.01%|0%|0%|1.47%)
SP212 Peptostreptococcus stomatis (0.03%|0%|2.52%|0%|1.26%|0.03%)
SP96 Clostridioides difficile (0%|0%|0.71%|0%|0%|0.18%)
SP547 Intestinibacter bartlettii (0%|0%|0%|0%|0%|0.01%)
SP805 Paeniclostridium [Eubacterium] tenue (0%|0%|0%|0%|0%|0.00%)
SP309 Peptostreptococcaceae_[XI][G-9] [Eubacterium]_brachy (0%|0%|0.66%|0%|0
SP653 Peptostreptococcaceae_[XI][G-1] bacterium_HMT_091 (0%|0%|0.01%|0%|
SP388 Peptostreptococcaceae_[XI][G-4] bacterium_HMT_369 (0%|0%|0.00%|0.18%
SP239 Peptostreptococcaceae_[XI][G-6] [Eubacterium]_nodatum (0%|0%|0.0
SP424 Peptostreptococcaceae_[XI][G-1] [Eubacterium]_infirmum (0%|0%|0%|0%
SP294 Peptostreptococcaceae_[XI][G-1] [Eubacterium]_sulci (0%|0%|0.49%|0%|
SP644 Mogibacterium timidum (0%|0%|0.03%|0%|0.00%|0%)
SP259 Peptostreptococcaceae_[XI][G-5] [Eubacterium]_saphenum (0%|0%|0%|0.2
SP634 Peptostreptococcaceae_[XI][G-5] bacterium_HMT_493 (0%|0%|0.02%|0
SP311 Anaerococcus provencensis (0%|0%|0%|0%|0%|0.03%)
SP111 Anaerococcus octavius (0.00%|0%|0%|0%|0%|0.09%)
SP139 Anaerococcus lactolyticus (0%|0%|0%|0%|0%|0.05%)
SP266 Anaerococcus sp._HMT_294 (0.00%|0%|0.01%|0%|0.00%|0.06%)
SP235 Anaerococcus obesiensis (0%|0%|0%|0%|0%|0.07%)
SP126 Anaerococcus vaginalis (0%|0%|0.02%|0%|0%|0.98%)
SP361 Anaerococcus jeddahensis (0%|0%|0%|0%|0%|0.01%)
SP42 Anaerococcus hydrogenalis (0%|0%|0.00%|0%|0%|1.54%)
SP670 Anaerococcus senegalensis (0%|0%|0%|0%|0%|0.01%)
SP724 Murdochiella asaccharolytica (0%|0%|0%|0%|0%|0.00%)
SP198 Parvimonas micra (0%|0%|0.58%|0%|0.21%|0%)
SP106 Finegoldia magna (0%|4.56%|0%|0%|0%|9.34%)
SP508 Peptoniphilaceae_[G-1] bacterium_HMT_113 (0%|0%|0%|0%|0.00%|0%)
SP194 Peptoniphilus coxii (0.00%|0%|0.00%|0%|0%|0.74%)
SP360 Peptoniphilus sp._HMT_375 (0%|0%|0%|0%|0%|0.02%)
SP221 Peptoniphilus duerdenii (0%|0%|0.00%|0%|0%|0.37%)
SP138 Peptoniphilus lacrimalis (0%|0%|0.03%|0%|0%|0.38%)
SP577 Peptoniphilus harei (0%|0%|0%|0%|0%|0.00%)
SP110 Peptoniphilus grossensis (0%|0%|0.09%|0%|0%|2.40%)
SP657 Peptoniphilus senegalensis (0%|0%|0%|0%|0%|0.00%)
SP220 Peptoniphilus tyrrelliae (0%|0%|0%|0%|0%|0.09%)
SP553 Peptococcus sp._HMT_167 (0%|0%|0.02%|0%|0.00%|0%)
SP697 Peptococcus sp._HMT_168 (0%|0%|0.05%|0%|0.01%|0%)
SP262 Peptococcus simiae (0%|0%|0%|0%|0.03%|0%)
SP308 Peptococcus niger (0%|0%|0%|0%|0.01%|0%)
SP285 Phascolarctobacterium faecium (0%|0%|0.05%|0%|0%|0%)
SP66 Megamonas funiformis (0%|0%|0%|0%|0%|0.08%)
SP648 Selenomonas sputigena (0%|0%|0.02%|0%|0.01%|0%)
SP768 Veillonellaceae_[G-1] bacterium_HMT_129 (0%|0%|0.00%|0%|0%|0%)
SP781 Veillonellaceae_[G-1] bacterium_HMT_145 (0%|0%|0.00%|0%|0.00%|0%)
SP473 Veillonellaceae_[G-1] bacterium_HMT_155 (0%|0%|0.01%|0%|0.01%|0%)
SP519 Mitsuokella sp._HMT_521 (0%|0%|0.00%|0%|0.00%|0%)
SP526 Selenomonas sp._HMT_136 (0.00%|0%|0.03%|0%|0.03%|0%)
SP824 Selenomonas sp._HMT_478 (0%|0%|0.00%|0%|0.00%|0%)
SP623 Selenomonas flueggei (0%|0%|0.00%|0%|0.00%|0%)
SP681 Selenomonas noxia (0%|0%|0.00%|0%|0.02%|0%)
SP568 Selenomonas artemidis (0%|0%|0%|0%|0.01%|0%)
SP647 Selenomonas sp._HMT_137 (0%|0%|0.00%|0%|0.01%|0%)
SP80 Veillonella sp._HMT_780 (0.62%|0%|0.00%|3.06%|0.01%|0.02%)
SP690 Veillonella sp._HMT_917 (0.01%|0%|0.00%|0%|0%|0%)
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%|0.05%|0%)
SP620 Megasphaera micronuciformis (0%|0%|0.03%|0%|0.01%|0.00%)
SP678 Megasphaera cerevisiae (0%|0%|0%|0%|0%|0.00%)
SP761 Anaeroglobus geminatus (0%|0%|0.02%|0%|0.00%|0%)
SP406 Negativicoccus massiliensis (0%|0%|0%|0%|0%|0.02%)
SP482 Negativicoccus succinicivorans (0%|0%|0%|0%|0%|0.01%)
SP660 Dialister micraerophilus (0%|0%|0%|0%|0%|0.00%)
SP99 Dialister propionicifaciens (0%|0%|0%|0%|0%|0.20%)
SP806 Dialister pneumosintes (0%|0%|0.00%|0%|0.00%|0%)
SP459 Dialister invisus (0%|0%|0.01%|0%|0.01%|0%)
SP668 Mollicutes_[G-1] bacterium_HMT_504 (0%|0%|0.01%|0%|0.00%|0%)
SP601 Mycoplasma faucium (0%|0%|0%|0%|0.05%|0%)
SP542 Mycoplasma salivarium (0%|0%|0.05%|0%|0.00%|0%)
SP828 Eggerthia catenaformis (0%|0%|0.01%|0%|0.00%|0%)
SP454 Erysipelatoclostridium ramosum (0%|0%|0.00%|0%|0%|0.02%)
SP252 Erysipelatoclostridium [Clostridium] innocuum (0%|0.14%|0%|0%|0%|0.
SP610 Erysipelotrichaceae_[G-1] bacterium_HMT_904 (0%|0%|0%|0%|0.0
SP827 Erysipelotrichaceae_[G-1] bacterium_HMT_905 (0%|0%|0%|0%|0.01%|
SP525 Bulleidia extructa (0%|0%|0.05%|0%|0.01%|0%)
SP119 Solobacterium moorei (0%|0%|2.74%|0%|0.40%|0.34%)
SP832 Paenibacillus thailandensis (0%|0%|0%|0%|0.00%|0%)
SP130 Gemella sanguinis (0.01%|0%|2.89%|0%|2.56%|0%)
SP157 Gemella morbillorum (0%|0%|0.23%|0%|0.07%|0.13%)
SP108 Gemella haemolysans (2.66%|0.01%|4.47%|44.91%|1.49%|0.08%)
SP643 Dolosigranulum pigrum (0%|0%|0.00%|0%|0%|0%)
SP785 Atopostipes sp._HOT_200 (0%|0%|0.00%|0%|0%|0%)
SP79 Abiotrophia defectiva (0.12%|0%|0.17%|0%|1.38%|0%)
SP732 Facklamia hominis (0.00%|0%|0%|0%|0%|0%)
SP217 Granulicatella elegans (0.21%|0%|1.08%|1.14%|0.07%|0.00%)
SP135 Granulicatella adiacens (0.37%|0%|5.13%|5.44%|8.11%|0.03%)
SP53 Enterococcus faecalis (0.48%|18.26%|0%|0%|0%|18.27%)
SP326 Brochothrix thermosphacta (0.00%|0%|0%|0%|0%|0%)
SP395 Weizmannia ginsengihumi (0%|0%|0%|0%|0%|0.03%)
SP752 Bacillus swezeyi (0%|0%|0%|0%|0%|0.00%)
SP718 Bacillus carboniphilus (0%|0%|0%|0%|0.00%|0%)
SP234 Staphylococcus simulans (0%|0%|0%|0%|0%|0.07%)
SP717 Staphylococcus auricularis (0.00%|0%|0%|0%|0%|0%)
SP373 Staphylococcus ureilyticus (0%|0%|0%|0%|0%|0.01%)
SP306 Staphylococcus equorum (0%|0%|0%|0%|0%|0.00%)
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%)
SP215 Staphylococcus warneri (0.02%|0%|0%|0%|0%|0.03%)
SP148 Staphylococcus pasteuri (0.00%|0%|0%|0%|0%|0%)
SP56 Staphylococcus capitis (1.62%|0%|0%|0.03%|0%|0.10%)
SP124 Staphylococcus saccharolyticus (0%|0%|0%|0%|0%|0.17%)
SP663 Leuconostoc carnosum (0.00%|0%|0%|0%|0%|0%)
SP720 Ligilactobacillus saerimneri (0%|0%|0%|0%|0%|0%)
SP240 Limosilactobacillus mucosae (0%|0%|0%|0%|0%|0.17%)
SP345 Limosilactobacillus reuteri (0%|0%|0%|0%|0%|0.00%)
SP750 Lactobacillus iners (0.00%|0%|0%|0%|0%|0%)
SP162 Lactobacillus gasseri (0.55%|0%|0%|0.00%|0%|0.08%)
SP386 Lactobacillus johnsonii (0.00%|0%|0%|0%|0%|0%)
SP461 Lactobacillus hominis (0.00%|0%|0%|0%|0%|0%)
SP452 Lactococcus raffinolactis (0%|0%|0%|0%|0%|0.02%)
SP622 Streptococcus mutans (0%|0%|0%|0%|0.00%|0%)
SP426 Streptococcus thermophilus (0%|0%|0%|0%|0.00%|0.02%)
SP64 Streptococcus lactarius (25.66%|0.01%|2.82%|0.10%|0.20%|0.39%)
SP65 Streptococcus peroris (0.07%|0%|0.34%|0%|0.15%|0.00%)
SP203 Streptococcus sanguinis (0.01%|0%|0.28%|0%|0.79%|0.02%)
SP22 Streptococcus sp._HMT_074 (2.84%|0%|16.60%|2.77%|25.20%|0.2
SP55 Streptococcus parasanguinis_clade_411 (2.75%|0.19%|14.38%|12.6
SP68 Streptococcus sp._HMT_056 (0.33%|0.00%|0.75%|0.62%|0.14%|0.00%)
SP247 Streptococcus gordonii (0.01%|0.00%|0.02%|0.06%|0.01%|0%)
SP107 Streptococcus sinensis (0.00%|0%|0.17%|0.00%|0.01%|0.00%)
SP101 Streptococcus anginosus (0%|6.67%|0.03%|0%|0.01%|0.00%)
SP428 Streptococcus intermedius (0.01%|0.00%|0.02%|0%|0.01%|0%)
Baby_Saliva_Preterm
Baby_Stool_On-term
Mother_Saliva_On-term
Baby_Saliva_On-term
Mother_Saliva_Preterm
Baby_Stool_Preterm
0
0.25
0.5
0.75
1
substitutions/site