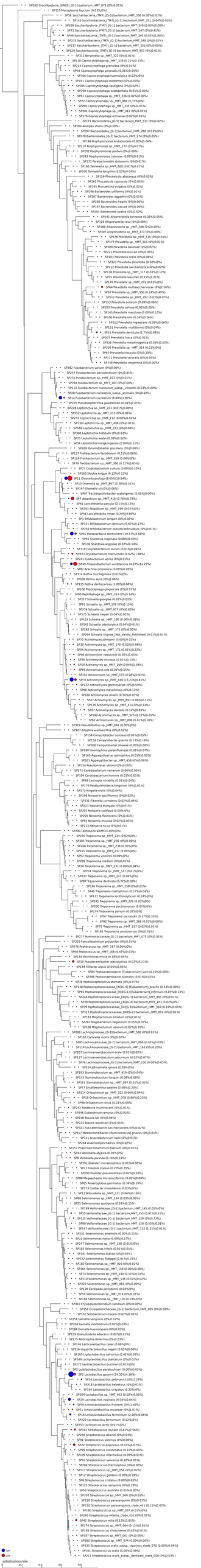
SP2162 Gracilibacteria_(GN02)_(G-1)_bacterium_HMT_952 (0%0.0%)
SP115 Mycoplasma faucium (0.03%0%)
SP51 Saccharofermentans_(TM7_(G-5))_bacterium_HMT_356 (0.30%0.03%)
SP143 Saccharofermentans_(TM7_(G-5))_bacterium_HMT_351 (0.04%0.03%)
SP209 Saccharofermentans_(TM7_(G-1))_bacterium_HMT_349 (0.03%0.00%)
SP51 Saccharofermentans_(TM7_(G-1))_bacterium_HMT_347 (0%0.00%)
SP46 Saccharofermentans_(TM7_(G-1))_bacterium_HMT_346 (0.05%1.80%)
SP25 Saccharofermentans_(TM7_(G-1))_bacterium_HMT_348 (0%0.02%)
SP237 Saccharofermentans_(TM7_(G-1))_bacterium_HMT_352 (0%0.00%)
SP129 Saccharofermentans_(TM7_(G-1))_bacterium_HMT_357 (0%0.01%)
SP92 Bergeyella sp._HMT_322 (0%0.01%)
SP118 Capnocytophaga sp._HMT_338 (0.21%0.15%)
SP216 Capnocytophaga granulosa (0%0.01%)
SP54 Capnocytophaga gingivalis (0.01%0.01%)
SP308 Capnocytophaga haemolytica (0%0.00%)
SP191 Capnocytophaga leadbetteri (0%0.00%)
SP285 Capnocytophaga ochracea (0%0.02%)
SP296 Capnocytophaga endodontalis (0.01%0.00%)
SP81 Capnocytophaga sp._HMT_336 (0.02%0.30%)
SP72 Capnocytophaga sp._HMT_864 (0.37%0%)
SP260 Capnocytophaga sp._HMT_335 (0%0.01%)
SP261 Capnocytophaga sp._HMT_412 (0%0.01%)
SP176 Capnocytophaga ochracea (0.01%0.01%)
SP172 Bacteroides_(G-5)_bacterium_HMT_511 (0%0.02%)
SP284 Alistipes (0%0.00%)
SP297 Bacteroides_(G-3)_bacterium_HMT_200 (0.03%0%)
SP278 Bacteroides_(G-2)_bacterium_HMT_274 (0%0.01%)
SP39 Porphyromonas endodontalis (0.00%0.04%)
SP210 Porphyromonas sp._HMT_277 (0%0)
SP202 Porphyromonas pasteri (0%0.00%)
SP243 Porphyromonas catoniae (0.00%0.01%)
SP133 Parabacteroides distasonis (0%0.01%)
SP186 Tannerella sp._HMT_808 (0.01%0.01%)
SP100 Tannerella forsythia (0.02%0.04%)
SP156 Phocaicola abscessus (0%0.02%)
SP162 Phocaicola coprocola (0%0.05%)
SP205 Phocaicola vulgatus (0%0.02%)
SP298 Bacteroides uniformis (0%0.01%)
SP287 Bacteroides eggerthii (0%0.01%)
SP286 Bacteroides fragilis (0%0.00%)
SP167 Bacteroides caccae (0%0.00%)
SP281 Bacteroides ovatus (0%0.00%)
SP141 Alloprevotella tannerae (0.02%0.05%)
SP16 Alloprevotella rava (0%0.00%)
SP188 Alloprevotella sp._HMT_308 (0%0.00%)
SP307 Alloprevotella sp._HMT_473 (0%0.00%)
SP178 Prevotella sp._HMT_515 (0%0.01%)
SP177 Prevotella sp._HMT_315 (0%0.01%)
SP309 Prevotella baroniae (0%0.02%)
SP221 Prevotella buccae (0%0.06%)
SP103 Prevotella oralis (0%0.06%)
SP212 Prevotella pleuritis (0.02%0%)
SP212 Prevotella saccharolytica (0%0.02%)
SP136 Prevotella sp._HMT_317 (0.03%0.17%)
SP35 Prevotella loescheii (0.15%0.01%)
SP170 Prevotella sp._HMT_472 (0.01%0%)
SP64 Prevotella multisaccharivorax (2.56%)
SP63 Prevotella sp._HMT_300 (2.24%0.40%)
SP112 Prevotella sp._HMT_292 (0.01%0.03%)
SP155 Prevotella oulorum (0.09%0.06%)
SP203 Prevotella salivae (0.02%0.05%)
SP145 Prevotella maculosa (0.50%0.13%)
SP106 Prevotella oris (0.34%0.26%)
SP113 Prevotella nigrescens (0.05%0.06%)
SP151 Prevotella multiformis (0%0.04%)
SP1 Prevotella denticola (1.75%0.69%)
SP303 Prevotella fusca (0%0.01%)
SP226 Prevotella melaninogenica (0.03%0.02%)
SP126 Prevotella sp._HMT_314 (0.01%0%)
SP67 Prevotella histicola (0%0.19%)
SP171 Prevotella veroralis (0%0.00%)
SP189 Prevotella enoeca (0%0.00%)
SP282 Fusobacterium varium (0%0.00%)
SP257 Fusobacterium periodonticum (0%0.01%)
SP231 Fusobacterium sp._HMT_203 (0%0.01%)
SP294 Fusobacterium sp._HMT_204 (0%0.00%)
SP110 Fusobacterium nucleatum_subsp._vincentii (0.03%0.00%)
SP28 Fusobacterium nucleatum_subsp._animalis (0%0.02%)
SP14 Fusobacterium nucleatum (0.89%0.80%)
SP220 Pseudoleptotrichia goodfellowii (0.04%0.01%)
SP228 Leptotrichia sp._HMT_221 (0.01%0.00%)
SP252 Leptotrichia sp._HMT_215 (0%0.01%)
SP255 Leptotrichia sp._HMT_212 (0.00%0.02%)
SP238 Leptotrichia sp._HMT_498 (0%0.01%)
SP168 Leptotrichia sp._HMT_223 (0%0.08%)
SP300 Leptotrichia hofstadii (0%0.01%)
SP73 Leptotrichia wadei (0.09%0.02%)
SP54 Leptotrichia hongkongensis (0.00%0.11%)
SP269 Pyramidobacter piscolens (0%0.00%)
SP117 Helcococcus fastidiosus (0.51%0.06%)
SP124 Fretibacterium sp._HMT_359 (0%0.00%)
SP79 Fretibacterium sp._HMT_360 (0.12%0.05%)
SP57 Cryptobacterium curtum (0.09%0.10%)
SP189 Slackia exigua (0.12%0.01%)
SP11 Olsenella profusa (8.05%19.84%)
SP15 Olsenella sp._HMT_807 (0.38%0.31%)
SP197 Olsenella uli (0%0.06%)
SP47 Thermoleptotrichaceae scatologenes (0.05%0.00%)
SP7 Atopobium sp._HMT_416 (0.76%8.73%)
SP41 Lancefieldella parauria (0.15%0.13%)
SP293 Atopobium sp._HMT_199 (0.02%0%)
SP48 Lancefieldella rimae (0.24%0.60%)
SP1 Bifidobacterium longum (0%0.00%)
SP121 Bifidobacterium dentium (0.67%0.17%)
SP259 Bifidobacterium pseudocatenulatum (0%0.01%)
SP45 Paracoccus denitrificans (13.33%3.68%)
SP55 Scardovia inopinata (0.08%0.40%)
SP130 Scardovia wiggsiae (0.01%0.14%)
SP119 Corynebacterium durum (0.01%0.09%)
SP43 Corynebacterium matruchotii (0.02%1.86%)
SP141 Cutibacterium acnes (0%0.01%)
SP49 Propionibacterium acidifaciens (4.67%12.27%)
SP62 Arachnia propionica (0.08%0.36%)
SP114 Rothia mucilaginosa (0.01%0%)
SP109 Rothia aeria (0%0.06%)
SP115 Rothia dentocariosa (1.09%0.68%)
SP206 Pedobacter gingivicola (0%0.03%)
SP66 Pedobacter sp._HMT_183 (0%0.01%)
SP217 Schaalia georgiae (0.02%0.02%)
SP61 Schaalia sp._HMT_178 (0%0%)
SP239 Schaalia sp._HMT_877 (0%0.00%)
SP175 Schaalia meyeri (0.04%0.02%)
SP123 Schaalia sp._HMT_180 (0.06%0.08%)
SP142 Schaalia odontolytica (0.04%0.01%)
SP265 Schaalia sp._HMT_172 (0%0.00%)
SP264 Schaalia lingnae_(Not_Validly_Published) (0.01%0.01%)
SP70 Actinomyces johnsonii (0.00%0.02%)
SP50 Actinomyces sp._HMT_170 (0.02%0.48%)
SP64 Actinomyces sp._HMT_171 (0.01%0.22%)
SP66 Actinomyces naeslundii (0.05%0.65%)
SP26 Actinomyces viscosus (0.01%0.14%)
SP19 Actinomyces sp._HMT_169 (0.04%1.38%)
SP69 Actinomyces oris (0.04%0.55%)
SP182 Actinomyces sp._HMT_175 (0.08%0.03%)
SP78 Actinomyces sp._HMT_448 (11.12%8.81%)
SP132 Actinomyces graevenitzae (0%0.50%)
SP86 Actinomyces massiliensis (0%0.13%)
SP160 Actinomyces israelii (0.50%0.05%)
SP47 Actinomyces sp._HMT_897 (0.06%0.11%)
SP126 Actinomyces sp._HMT_414 (0%0.31%)
SP27 Actinomyces denticolens (0.12%0.83%)
SP140 Actinomyces sp._HMT_525 (0.17%0.01%)
SP62 Actinomyces sp._HMT_896 (0.01%0.18%)
SP119 Desulfobulbus sp._HMT_041 (0.00%0%)
SP107 Bilophila wadsworthia (0%0.02%)
SP154 Campylobacter concisus (0.01%0.05%)
SP158 Campylobacter gracilis (0.11%0.18%)
SP308 Campylobacter showae (0.00%0.00%)
SP180 Haemophilus parainfluenzae (0.01%0.07%)
SP165 Aggregatibacter aphrophilus (0.01%0.00%)
SP261 Aggregatibacter sp._HMT_458 (0%0.06%)
SP254 Pseudomonas veronii (0%0.00%)
SP271 Cardiobacterium valvarum (0.00%0.00%)
SP104 Cardiobacterium hominis (0.01%0.01%)
SP89 Lautropia mirabilis (0.01%0.04%)
SP179 Paraburkholderia fungorum (0%0.01%)
SP173 Kingella oralis (0%0.04%)
SP198 Neisseria bacilliformis (0%0.02%)
SP111 Eikenella corrodens (0.02%0.08%)
SP23 Neisseria elongata (0%0.01%)
SP292 Neisseria subflava (0.00%0%)
SP205 Neisseria flavescens (0%0.02%)
SP83 Neisseria mucosa (0.02%0.03%)
SP213 Neisseria sicca (0%0.01%)
SP290 Leptospira wolffi (0.00%0%)
SP270 Treponema sp._HMT_234 (0.02%0%)
SP305 Treponema sp._HMT_230 (0.00%0%)
SP268 Treponema sp._HMT_238 (0.00%0%)
SP25 Treponema sp._HMT_231 (0.09%0%)
SP52 Treponema vincentii (0.09%0%)
SP289 Treponema medium (0%0.01%)
SP20 Treponema sp._HMT_231 (0.00%0.04%)
SP274 Treponema sp._HMT_517 (0.01%0%)
SP227 Treponema sp._HMT_247 (0.04%0%)
SP87 Treponema denticola (0.72%0.03%)
SP196 Treponema sp._HMT_758 (0%0.01%)
SP40 Treponema maltophilum (1.17%0.04%)
SP111 Treponema lecithinolyticum (0.24%0%)
SP245 Treponema sp._HMT_270 (0.03%0%)
SP250 Treponema pectinovorum (0.01%0%)
SP278 Treponema parvum (0%0%)
SP57 Treponema socranskii (0.37%0.33%)
SP92 Treponema sp._HMT_268 (0.03%0.08%)
SP75 Treponema sp._HMT_257 (0.02%0.01%)
SP195 Treponema amylovorum (0%0.01%)
SP277 Ruminococcaceae_(G-1)_bacterium_HMT_075 (0%0.01%)
SP29 Faecalibacterium prausnitzii (0%0.03%)
SP276 Peptococcus sp._HMT_167 (0.00%0%)
SP60 Peptococcus sp._HMT_168 (0.47%0.21%)
SP134 Parvimonas micra (0.18%0.04%)
SP10 Peptostreptococcus stomatis (0.03%4.21%)
SP144 Filifactor alocis (0.05%0.05%)
SP84 Peptoanaerobacter [Eubacterium] yurii (0.10%0.00%)
SP298 Peptoanaerobacter stomatis (0.01%0.02%)
SP36 Peptostreptococcus stomatis (0%0.01%)
SP148 Peptostreptococcaceae_(XI)-(3)-bacterium] brachy (0.43%0.00%)
SP65 Peptostreptococcaceae_(XI)[G-11]bacterium]_infirmum (0.05%0.13%)
SP248 Peptostreptococcaceae_(XI)[G-3]_bacterium_HMT_495 (0%0.01%)
SP258 Peptostreptococcaceae_(XI)[G-4]_bacterium_HMT_103 (0.04%0%)
SP39 Peptostreptococcaceae_(XI)[G-4]_bacterium_HMT_291 (0%0.01%)
SP313 Peptostreptococcaceae_(XI)[G-6]_bacterium_HMT_091 (0%0.01%)
SP183 Mogibacterium timidum (0%0.01%)
SP203 Mogibacterium neglectum (0.00%0.02%)
SP108 Mogibacterium vescum (0.42%0.16%)
SP208 Lachnospiraceae_[G-8]_bacterium_HMT_500 (0%0.01%)
SP168 Catonella morbi (0%0.02%)
SP85 Lachnospiraceae_[G-7]_bacterium_HMT_086 (0.03%0.03%)
SP114 Lachnospiraceae_[G-1]_bacterium_HMT_081 (0%0.20%)
SP207 Lachnoanaerobaculum oral (0.01%0.02%)
SP247 Lachnoanaerobaculum saburreum (0.03%0.07%)
SP76 Lachnospiraceae_[G-3]_bacterium_HMT_100 (0.00%0.05%)
SP224 Johnsonella ignava (0.03%0%)
SP193 Stomatobaculum sp._HMT_097 (0%0.00%)
SP155 Stomatobaculum longum (0.09%0.08%)
SP161 Stomatobaculum sp._HMT_097 (0.01%0.02%)
SP77 Shuttleworthia satelles (0.08%0.23%)
SP214 Oribacterium sp._HMT_102 (0.00%0.04%)
SP18 Oribacterium sp._HMT_078 (0.86%0.23%)
SP98 Oribacterium sinus (0.01%0.00%)
SP42 Roseburia inulinivorans (0%0.01%)
SP146 Eubacterium ramulus (0%0%)
SP218 Blautia luti (0%0.04%)
SP25 Blautia wexlerae (0%0.01%)
SP321 Fusicatenibacter saccharivorans (0%0.02%)
SP313 Mediterraneibacter [Ruminococcus] gnavus (0%0.01%)
SP111 Anaerobiluticum hallii (0%0.01%)
SP244 Anaerostipes hadrus (0%0.02%)
SP157 Phascolarctobacterium faecium (0%0.01%)
SP44 Veillonella atypica (0.05%0%)
SP8 Veillonella parvula (0.16%0.13%)
SP291 Dialister micraerophilus (0.01%0.00%)
SP17 Dialister invisus (0.20%0.35%)
SP200 Dialister pneumosintes (0.62%0.03%)
SP88 Megasphaera micronuciformis (0.05%0.09%)
SP60 Anaerogloboides genmaiea (0.34%0.19%)
SP273 Colibacter massiliensis (0.03%0%)
SP13 Mitsuokella sp._HMT_131 (0.06%0.19%)
SP68 Selenomonas sp._HMT_134 (0.01%0%)
SP25 Selenomonas spurigena (0.29%0.15%)
SP189 Veillonellaceae_[G-1]_bacterium_HMT_145 (0.01%0%)
SP50 Veillonellaceae_[G-1]_bacterium_HMT_155 (0.91%0.11%)
SP122 Veillonellaceae_[G-1]_bacterium_HMT_129 (0%0.73%)
SP60 Veillonellaceae_[G-1]_bacterium_HMT_150 (0.55%0%)
SP147 Veillonellaceae_[G-1]_bacterium_HMT_132 (1.21%0.01%)
SP151 Selenomonas artemidis (0.00%0%)
SP23 Selenomonas noxia (0.09%0.17%)
SP247 Selenomonas sp._HMT_136 (0.07%0%)
SP185 Selenomonas infelix (0.01%0.01%)
SP181 Selenomonas dianae (0%0.02%)
SP232 Selenomonas flueggei (0.01%0.01%)
SP182 Selenomonas sp._HMT_920 (0%0.01%)
SP164 Selenomonas sp._HMT_146 (0.00%0.00%)
SP74 Selenomonas sp._HMT_149 (1.11%0.01%)
SP119 Selenomonas sp._HMT_136 (0.02%0.02%)
SP322 Selenomonas sp._HMT_481 (0%0.00%)
SP128 Centipeda periodontii (0.04%0%)
SP59 Selenomonas sp._HMT_919 (0%0.01%)
SP166 Selenomonas sp._HMT_126 (0.03%0%)
SP110 Erysipelatotrichaceae_bacterium (0%0%)
SP150 Erysipelatotrichaceae_[G-1]_bacterium_HMT_905 (0%0.02%)
SP158 Gemella morbillorum (0.02%0.03%)
SP159 Gemella sanguinis (0%0.02%)
SP206 Gemella haemolysans (0.02%0.03%)
SP139 Granulicatella adiacens (0.04%0%)
SP275 Abiotrophia defectiva (0%0.03%)
SP146 Lactobacillus iners (0.04%0%)
SP116 Liquorilactobacillus nagelii (0.05%0.00%)
SP105 Lipilactobacillus salivarius (0.01%0.00%)
SP249 Lactiplantibacillus plantarum (0%0.01%)
SP272 Lentilactobacillus buchneri (0.03%0%)
SP5 Lentilactobacillus parabuchneri (0.00%0.02%)
SP2 Lactobacillus gasseri (24.30%4.16%)
SP34 Lactobacillus delbrueckii (0%1.28%)
SP318 Lactobacillus helveticus (0%0.01%)
SP194 Lactobacillus crispatus (0.32%0%)
SP304 Lactobacillus sp._HMT_052 (0.02%0.00%)
SP3 Lactobacillus vaginalis (6.04%0.04%)
SP56 Limosilactobacillus mucosae (0%0.01%)
SP21 Limosilactobacillus mucosae (0%0.01%)
SP18 Limosilactobacillus fermentum (2.90%0.06%)
SP222 Lactobacillus fermentum (0.02%0%)
SP253 Lactococcus lactis (0.01%0%)
SP192 Streptococcus mutans (0.81%2.56%)
SP256 Streptococcus downei (0%0.03%)
SP91 Streptococcus sobrinus (0%0.06%)
SP22 Streptococcus angiosus (0.07%0.07%)
SP99 Streptococcus constellatus (0.10%0.00%)
SP139 Streptococcus intermedius (0.03%0.01%)
SP93 Streptococcus salivarius (0.10%0.01%)
SP288 Streptococcus thermophilus (0%0%)
SP117 Streptococcus sp._HMT_056 (0%0.02%)
SP12 Streptococcus gordonii (0.09%0.29%)
SP6 Streptococcus cristatus (0.00%0.02%)
SP125 Streptococcus sanguinis (0%0.28%)
SP53 Streptococcus australis (0.01%0.00%)
SP320 Streptococcus sp._HMT_066 (0%0.01%)
SP230 Streptococcus parasanguinis (0%0.01%)
SP120 Streptococcus parasanguinis (0.02%0.02%)
SP190 Streptococcus sp._HMT_057 (0.01%0%)
SP240 Streptococcus infantis_clade_431 (0%0.01%)
SP42 Streptococcus mitis (0.13%2.81%)
SP174 Streptococcus sp._HMT_064 (0.12%0.01%)
SP94 Streptococcus chosuneae (0.03%0.01%)
SP187 Streptococcus sp._HMT_061 (0%0.00%)
SP180 Streptococcus sp._HMT_074 (0%0.00%)
SP135 Streptococcus oralis_subsp._tigurinus_clade_070 (0.00%0.04%)
SP101 Streptococcus oralis (0%0.49%)
SP127 Streptococcus oralis_subsp._dentisani_clade_058 (0%0.03%)
SP
NP
Institutions/site
0.4
0.6
0.8
1.0