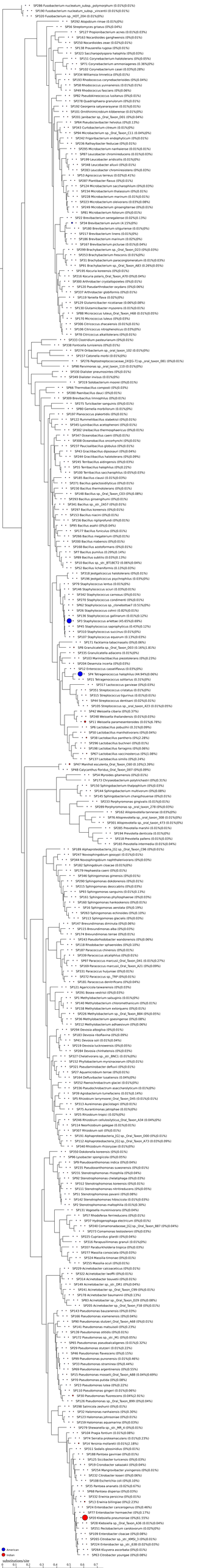
SP286 Fusobacterium nucleatum_subsp._polymorphum (0.01%|0.01%)
SP190 Fusobacterium nucleatum_subsp._vincentii (0.01%|0.01%)
SP320 Fusobacterium sp._HOT_204 (0.01%|0%)
SP292 Alloprevotella rimae (0.01%|0%)
SP56 Streptococcus gordonii (0%|0.04%)
SP127 Propionibacterium acnes (0.01%|0.03%)
SP163 Nocardioides sediminis (0%|0.01%)
SP250 Nocardioides zeae (0.02%|0.01%)
SP138 Pracaecia rugosa (0%|0.01%)
SP323 Saccharopolyspora halophila (0%|0.03%)
SP171 Corynebacterium glutamicum (0.05%|0%)
SP71 Corynebacterium ammoniagenes (0.36%|0%)
SP102 Corynebacterium case (0.03%|0.28%)
SP334 Williamsia limnetica (0%|0.01%)
SP193 Rhodococcus corynebacterioides (0%|0.04%)
SP58 Rhodococcus yunnanensis (0.01%|0.01%)
SP49 Rhodococcus fascians (0%|0%)
SP82 Pseudokineococcus lusitanus (0%|0.01%)
SP278 Quadrisphaera granulorum (0%|0.01%)
SP192 Georgenia satyanarayanai (0.01%|0.01%)
SP101 Ornithinimicrobium kibberense (0.01%|0%)
SP201 Janibacter sp._Oral_Taxon_001 (0%|0.04%)
SP64 Pseudoclavibacter helvolus (0%|0.13%)
SP343 Curtobacterium citreum (0.01%|0%)
SP94 Microbacterium sp._Oral_Taxon_211 (0.04%|0%)
SP242 Frigoribacterium endophyticum (0%|0.01%)
SP216 Rathayibacter festucae (0%|0.01%)
SP295 Microbacterium nanhaiense (0.01%|0.01%)
SP87 Leucobacter chromiireducens (0.01%|0.03%)
SP299 Leucobacter aridicola (0.01%|0%)
SP348 Leucobacter aluvii (0%|0.01%)
SP281 Leucobacter chromiiresistens (0%|0.03%)
SP53 Agrococcus terreus (0.02%|0.41%)
SP287 Plantibacter flavus (0%|0.01%)
SP224 Microbacterium saccharophilum (0%|0.03%)
SP234 Microbacterium thalassium (0%|0.01%)
SP228 Microbacterium marinum (0.01%|0.01%)
SP223 Microbacterium oleivorans (0.03%|0.08%)
SP249 Microbacterium ginsengiterrae (0%|0.01%)
SP61 Microbacterium foliorum (0%|0.01%)
SP22 Brevibacterium senegalense (0.02%|0.13%)
SP34 Brevibacterium avium (4.15%|0%)
SP180 Brevibacterium sloguiense (0.01%|0%)
SP217 Brevibacterium linens (0.01%|0%)
SP186 Brevibacterium marinum (0.02%|0%)
SP167 Brevibacterium pictum (0.01%|0.04%)
SP299 Brachybacterium sp._Oral_Taxon_023 (0%|0.03%)
SP253 Brachybacterium fresconense (0%|0%)
SP31 Brachybacterium paracongolenense (0.01%|0.03%)
SP91 Brachybacterium sp._Oral_Taxon_483 (0.26%|0.05%)
SP195 Kocuria kenensis (0%|0.01%)
SP216 Kocuria polaris_Oral_Taxon_A70 (0%|0.04%)
SP300 Arthrobacter crystallopoietes (0%|0.01%)
SP128 Pseudarthrobacter oxydans (0%|0.06%)
SP337 Arthrobacter globiformis (0%|0.01%)
SP119 Nesiella flava (0.02%|0%)
SP129 Glutamicibacter nicotianae (0.06%|0.08%)
SP130 Glutamicibacter mysorens (0.01%|0.01%)
SP98 Micrococcus lutea_Oral_Taxon_468 (0.01%|0.05%)
SP170 Micrococcus luteus (0%|0.03%)
SP306 Citricoccus zhacaiensis (0.01%|0.01%)
SP106 Citricoccus nitrophenolicus (0.03%|0%)
SP78 Citricoccus alkalitolerans (0%|0.01%)
SP333 Cistribacter paderaiensis (0%|0.01%)
SP338 Fontibacillus tunisiensis (0%|0.01%)
SP274 Oribacterium sp._oral_taxon_102 (0.01%|0%)
SP157 Catonella morbi (0.01%|0%)
SP276 Peptostreptococcaceae_[XI][G-7]_sp._oral_taxon_081 (0%|0.01%)
SP98 Parvimonas sp._oral_taxon_110 (0.01%|0%)
SP330 Dialister pneumosintes (0%|0.01%)
SP349 Dialister invisus (0.01%|0%)
SP319 Solobacterium moorei (0%|0.01%)
SP66 Thermobacillus composti (0%|0.03%)
SP280 Paenibacillus dauci (0%|0.01%)
SP309 Brevibacillus limnophilus (0%|0.01%)
SP275 Turicibacter sanguinis (0%|0.01%)
SP80 Gemella morbillorum (0.01%|0%)
SP197 Planococcus plakortidis (0%|0.01%)
SP122 Rummeliibacillus stabekisii (0%|0.01%)
SP345 Lysinibacillus acetophenoni (0%|0.01%)
SP302 Urebacillus themophaeresus (0%|0.01%)
SP347 Oceanobacillus caeni (0%|0.01%)
SP308 Oceanobacillus oncorhynchi (0%|0.01%)
SP237 Piscicaillibacillus globulus (0%|0.01%)
SP43 Graciibacillus dipsosauri (0%|0.04%)
SP244 Graciibacillus halotolerans (0%|0%)
SP243 Terribacillus aidingensis (0%|0.03%)
SP65 Terribacillus halophilus (0%|0.22%)
SP100 Terribacillus saccharophilus (0.05%|0.03%)
SP185 Bacillus cleveli (0.01%|0.03%)
SP271 Bacillus gelatinilyticus (0%|0%)
SP230 Bacillus thermotolerans (0%|0.01%)
SP148 Bacillus sp._Oral_Taxon_220 (0.05%|0.08%)
SP293 Bacillus ginsengihumi (0%|0%)
SP341 Bacillus sp._str._JA57 (0%|0.01%)
SP297 Bacillus koreensis (0%|0.01%)
SP153 Bacillus niacini (0%|0.01%)
SP156 Bacillus rigiliprofundi (0%|0.01%)
SP95 Bacillus aiahi (0%|0.04%)
SP177 Bacillus funiculus (0%|0.01%)
SP266 Bacillus megaterium (0%|0.01%)
SP200 Bacillus niabensis (0%|0.01%)
SP168 Bacillus azotoformans (0%|0.01%)
SP7 Bacillus pumilus (0.29%|0.14%)
SP89 Bacillus subtilis (0.03%|0.13%)
SP10 Bacillus sp._str._BT1B/CT2 (0.06%|0.04%)
SP52 Bacillus licheniformis (0.13%|0.03%)
SP318 Jeotgalicoccus halotolerans (0%|0.01%)
SP36 Jeotgalicoccus psychrophilus (0.03%|0%)
SP79 Staphylococcus lentus (0.01%|0%)
SP146 Staphylococcus sciuri (0.03%|0.01%)
SP342 Staphylococcus carnosus (0%|0.01%)
SP270 Staphylococcus condimenti (0%|0.01%)
SP62 Staphylococcus sp._clonebotaa7 (0.01%|0%)
SP56 Staphylococcus cohnii (0.82%|0.01%)
SP136 Staphylococcus gallinarum (0.01%|0.12%)
SP3 Staphylococcus arlettae (45.45%|0.88%)
SP45 Staphylococcus saprophyticus (0.43%|0.12%)
SP310 Staphylococcus succinus (0.01%|0%)
SP107 Staphylococcus equorum (0.13%|0.03%)
SP171 Facklamia tabacinasalis (0%|0.06%)
SP8 Granulicatella sp._Oral_Taxon_003 (0.18%|1.81%)
SP335 Granulicatella adiacens (0.01%|0%)
SP103 Marinilactibacillus piezotolerans (0%|0.23%)
SP204 Desemzia incerta (0%|0%)
SP12 Enterococcus casseliflavus (0.03%|0%)
SP4 Tetragenococcus halophilus (44.94%|0.06%)
SP21 Tetragenococcus solitarius (0.13%|0%)
SP317 Lactococcus garvieae (0%|0.03%)
SP351 Streptococcus criceti (0.01%|0%)
SP315 Streptococcus ilguinus (0.01%|0.01%)
SP44 Streptococcus dentisani (0.02%|0.01%)
SP15 Streptococcus sp._oral_taxon_423 (0.01%|0.05%)
SP42 Weissella cibaria (0%|0.37%)
SP248 Weissella thailandensis (0.01%|0.03%)
SP11 Weissella paramesenteroides (0.01%|8.78%)
SP6 Lactobacillus pobuzihi (0.31%|0.09%)
SP161 Lactobacillus mainhofiae (0%|0.06%)
SP38 Lactobacillus panthetis (0%|2.28%)
SP296 Lactobacillus buchneri (0%|0.01%)
SP198 Lactobacillus farraginis (0%|0.06%)
SP67 Lactobacillus vaccinostercus (0%|1.08%)
SP137 Lactobacillus similis (0%|0.24%)
SP47 Weinhut eiuolenta_Oral_Taxon_C60 (0.10%|3.39%)
SP48 Cryptanthrum floridum_Oral_Taxon_D07 (0%|0.09%)
SP54 Mycobacterium gilanyonis (0%|0.01%)
SP173 Chryseobacterium gleanjinkiethi (0%|0.31%)
SP150 Sphingobacterium thalpophilum (0%|0.03%)
SP144 Sphingobacterium multivorum (0%|0.08%)
SP145 Sphingobacterium changhouense (0%|0.01%)
SP233 Porphyromonas gingivalis (0.01%|0.01%)
SP289 Porphyromonas sp._oral_taxon_278 (0%|0%)
SP162 Alloprevotella tannerae (0.03%|0%)
SP76 Alloprevotella sp._oral_taxon_308 (0.01%|0%)
SP301 Alloprevotella sp._oral_taxon_473 (0.01%|0%)
SP285 Prevotella marshii (0.01%|0.01%)
SP184 Prevotella denticola (0.01%|0%)
SP126 Prevotella pallens (0.01%|0.03%)
SP165 Prevotella intermedia (0.01%|0.04%)
SP189 Alphaproteobacteria_[G]_sp._Oral_Taxon_CW (0%|0%)
SP247 Novosphingobium gossypii (0.01%|0.01%)
SP184 Novosphingobium naphthalenivorans (0%|0.03%)
SP182 Sphingobium cloacae (0.01%|0%)
SP179 Hephaestia caeni (0%|0.01%)
SP346 Sphingomonas ginsenosi (0%|0.01%)
SP090 Sphingomonas dokdonensis (0%|0.01%)
SP215 Sphingomonas desiccabilis (0%|0.03%)
SP91 Sphingomonas sanguinis (0.01%|0.13%)
SP161 Sphingomonas phyllosphaerae (0%|0.03%)
SP160 Sphingomonas haneokwonii (0%|0.01%)
SP16 Sphingomonas aerolata (0%|0.19%)
SP263 Sphingomonas echinoides (0%|0.10%)
SP113 Sphingomonas glacialis (0%|0.03%)
SP147 Brevundimonas diminuta (0%|0.06%)
SP115 Brevundimonas alba (0%|0.03%)
SP174 Brevundimonas terrae (0%|0.01%)
SP243 Pseudoholdoaster wandonensis (0%|0.06%)
SP118 Rhodobacter sphaeroides (0%|0.10%)
SP187 Paracoccus chinensis (0%|0.01%)
SP339 Paracoccus alcaliphilus (0%|0.01%)
SP97 Paracoccus marcusii_Oral_Taxon_041 (0.01%|0.27%)
SP169 Paracoccus marcusii_Oral_Taxon_A21 (0%|0.09%)
SP331 Paracoccus hujuinae (0%|0.01%)
SP272 Paracoccus sp._TNF (0%|0.01%)
SP181 Paracoccus denitrificans (0%|0.04%)
SP231 Agaricicola taiwanensis (0%|0%)
SP291 Bosea vestrisii (0%|0.03%)
SP1 Methylobacterium sabugina (0.01%|0%)
SP140 Methylobacterium chloromethanicum (0%|0.01%)
SP158 Methylobacterium extorquens (0%|0.01%)
SP224 Methylobacterium sp._Oral_Taxon_ba (0%|0.05%)
SP36 Methylobacterium gossypiorum (0%|0.08%)
SP212 Methylobacterium adhaesivum (0%|0.06%)
SP264 Devosia elbofgha (0%|0.01%)
SP183 Devosia riboflavina (0%|0.09%)
SP41 Devosia soli (0.01%|0.04%)
SP219 Devosia luckhowensis (0%|0.05%)
SP284 Devosia chihaiensis (0%|0.03%)
SP37 Cheilarovirus sp._str._JA6 (0.01%|0%)
SP152 Phyllobacterium myrsinacearum (0%|0.01%)
SP321 Pseudaminobacter defluvii (0%|0.01%)
SP37 Aquamicrobium terrae (0%|0.01%)
SP164 Defluvibacter lusatiensis (0.04%|0%)
SP252 Paenochrobactrum glaciei (0.01%|0%)
SP336 Pseudochrobactrum asaccharolyticum (0.01%|0%)
SP39 Agrobacterium tumefaciens (0.01%|0.14%)
SP5 Rhizobium larrymoorei_Oral_Taxon_045 (0.01%|0.01%)
SP313 Auremonas glacistagni (0%|0.01%)
SP75 Aurantimonas jatophae (0.01%|0%)
SP25 Rhizobium tropici (0.02%|0%)
SP246 Rhizobium cellulosilytica_Oral_Taxon_A34 (0.04%|0%)
SP114 Neorhizobium galegae (0.01%|0.01%)
SP307 Rhizobium soli (0%|0.01%)
SP131 Alphaproteobacteria_[G]_sp._Oral_Taxon_D00 (0%|0.01%)
SP112 Alphaproteobacteria_[G]_sp._Oral_Taxon_A73 (0.03%|0.09%)
SP340 Rhizobium rhizoryzae (0.01%|0%)
SP350 Dokdonella koreensis (0%|0.01%)
SP96 Lysobacter spongicola (0%|0.05%)
SP9 Pseudoxanthomonas indica (0%|0.04%)
SP235 Pseudoxanthomonas suwonensis (0%|0.01%)
SP231 Stenotrophomonas rhizophila (0%|0.04%)
SP92 Stenotrophomonas chelatiphaga (0%|0.03%)
SP312 Stenotrophomonas koreensis (0%|0.01%)
SP111 Stenotrophomonas nitritireducens (0%|0.05%)
SP51 Stenotrophomonas pavanii (0%|0.08%)
SP142 Stenotrophomonas hibiscicola (0.01%|0.03%)
SP2 Sporotrichomonas maltophilia (0.01%|0.30%)
SP131 Vogesella mureinivorus (0%|0.04%)
SP57 Rhodoferax ferrireducens (0%|0.01%)
SP37 Hydrogenophaga electricum (0%|0.01%)
SP240 Comamonadaceae_[F]_sp._Oral_Taxon_B87 (0%|0.04%)
SP273 Comamonas testosteroni (0%|0.03%)
SP225 Cupriavidus gilardii (0%|0.04%)
SP16 Parapusillomonas granuli (0.01%|0%)
SP207 Paraburkholderia tropica (0%|0.03%)
SP277 Massilia consociata (0%|0.03%)
SP24 Massilia timonae (0%|0.01%)
SP255 Massilia oculi (0%|0.01%)
SP229 Acinetobacter calcoaceticus (0%|0.01%)
SP322 Acinetobacter ieoffii (0%|0.01%)
SP314 Acinetobacter bouvetii (0%|0.01%)
SP149 Acinetobacter sp._str_DR1 (0%|0.04%)
SP241 Acinetobacter sp._Oral_Taxon_C99 (0%|0.01%)
SP178 Acinetobacter baumannii (0%|0.13%)
SP83 Acinetobacter sp._Oral_Taxon_229 (0%|0.08%)
SP205 Acinetobacter sp._Oral_Taxon_258 (0%|0.01%)
SP143 Pseudomonas baucanensis (0%|0.03%)
SP66 Pseudomonas xiamenensis (0%|0.04%)
SP90 Pseudomonas stutzeri_Oral_Taxon_A68 (0%|0.01%)
SP141 Pseudomonas mallusiae (0%|0.23%)
SP139 Pseudomonas otitidis (0%|0.01%)
SP172 Pseudomonas sp._str._M1 (0%|0.05%)
SP65 Pseudomonas pseudoalcaligenes (0.01%|0.32%)
SP29 Pseudomonas stutzeri (0.01%|0.22%)
SP46 Pseudomonas flavescens (0%|0.15%)
SP99 Pseudomonas punonensis (0.01%|0.46%)
SP33 Pseudomonas straminea (0%|0.44%)
SP69 Pseudomonas argentinensis (0%|0.55%)
SP15 Pseudomonas mosselii_Oral_Taxon_A88 (0.04%|0.69%)
SP35 Pseudomonas putida (0%|0.08%)
SP23 Pseudomonas lutea (0%|0.22%)
SP110 Pseudomonas gingeri (0.01%|0.06%)
SP30 Pseudomonas fluorescens (0.04%|2.91%)
SP126 Pseudomonas sp._Oral_Taxon_899 (0%|0.04%)
SP298 Salinicola peshiumi (0%|0.01%)
SP32 Halomonas nanhaiensis (0%|0.30%)
SP123 Halomonas johnsoniae (0%|0.01%)
SP159 Halomonas aquamarina (0%|0.03%)
SP279 Shewanella sp._str_M4_4 (0%|0.01%)
SP104 Pragia fontium (0.01%|0.08%)
SP74 Serratia proteamaculans (0.01%|0.23%)
SP14 Yersinia mollaretii (0.01%|2.18%)
SP311 Sodalis glossinidius (0%|0.01%)
SP188 Pantoea ananatis (0%|0.01%)
SP125 Scibacillus turicensis (0%|0.03%)
SP19 Crossiella sabuleti (0%|0.04%)
SP254 Mangrovibacter yiangensis (0%|0.01%)
SP232 Citrobacter koseri (0%|0.06%)
SP108 Escherichia coli (0%|0.10%)
SP325 Pantoea dispersa (0%|0.03%)
SP332 Erwinia persicina (0%|0.01%)
SP13 Erwinia billingiae (0%|2.23%)
SP24 Enterobacter cancerogenus (0%|0.46%)
SP77 Enterobacter hormaechei (0%|0.17%)
SP20 Klebsiella pneumoniae (0%|61.55%)
SP28 Klebsiella sp._Oral_Taxon_A36 (0.01%|0.04%)
SP251 Pectobacterium carotovorum (0.02%|0%)
SP109 Enterobacter cloacae (0%|0.08%)
SP265 Citrobacter sp._str._AMG_1 (0%|0.01%)
SP224 Enterobacter sp._str._638 (0.02%|0.03%)
SP264 Klebsiella aerogenes (0%|0.01%)
SP363 Citrobacter youngae (0%|0.08%)
Constitutions/site
0.1
0.3
0.4
0.5
0.6
0.7
American
Indian