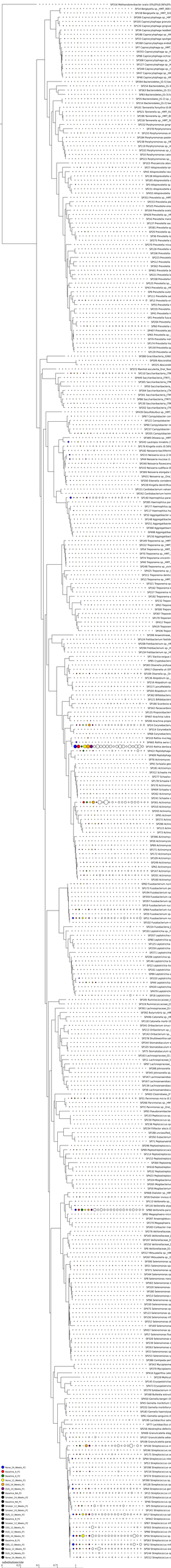
SP116 Methanobrevibacter oralis DSZ02b20 DENZ01
SP14 Bergeyella sp_HMT_950
SP158 Bergeyella sp_HMT_322
SP209 Capnocytophaga sp_HMT
SP320 Capnocytophaga granulosa
SP320 Capnocytophaga gingivalis
SP34 Capnocytophaga leadbetteri
SP186 Capnocytophaga sp_HMT
SP315 Capnocytophaga ochracea
SP308 Capnocytophaga endosymbiont
SP7 Capnocytophaga sp_HMT
SP255 Capnocytophaga sp_HMT
SP98 Capnocytophaga ochracea
SP308 Capnocytophaga sp_HMT
SP127 Capnocytophaga sp_HMT
SP349 Capnocytophaga sp_HMT
SP47 Capnocytophaga sp_HMT
SP48 Capnocytophaga sp_HMT
SP264 Bacteroides_ZQ-D1 bacteriotaxon
SP14 Bacteroides_ZQ-D1
SP63 Bacteroides_ZQ-D1 bacteriotaxon
SP53 Bacteroides_ZQ-D1 bacteriotaxon
SP14 Bacteroides_ZQ-D1 bacteriotaxon
SP101 Tannerella forsythia DI-B
SP521 Tannerella sp_HMT_911
SP136 Tannerella sp_HMT_917
SP118 Tannerella sp_HMT_21
SP52 Porphyromonas gingivalis
SP39 Porphyromonas
SP153 Porphyromonas endodontalis
SP184 Porphyromonas pasteri
SP238 Porphyromonas sp_HMT
SP10 Porphyromonas sp_HMT
SP133 Porphyromonas cangingivalis
SP10 Porphyromonas cangingivalis
SP133 Phocaeicola dorei
SP27 Alloprevotella tannerae
SP42 Alloprevotella rava
SP138 Alloprevotella sp_HMT
SP185 Alloprevotella sp_HMT
SP3 Alloprevotella sp_HMT
SP231 Alloprevotella sp_HMT
SP65 Alloprevotella sp_HMT
SP51 Prevotella sp_HMT
SP233 Prevotella pallens
SP125 Prevotella enoeca
SP154 Prevotella oralis
SP428 Prevotella sp_HMT
SP56 Prevotella marshii
SP137 Prevotella saccharolytica
SP261 Prevotella sp_HMT
SP5 Prevotella sp_HMT
SP6 Prevotella sp_HMT
SP275 Prevotella sp_HMT
SP370 Prevotella intermedia
SP126 Prevotella sp_HMT
SP136 Prevotella sp_HMT
SP223 Prevotella sp_HMT
SP112 Prevotella sp_HMT
SP302 Prevotella sp_HMT
SP461 Prevotella sp_HMT
SP221 Prevotella sp_HMT
SP138 Prevotella sp_HMT
SP125 Prevotella sp_HMT
SP43 Prevotella sp_HMT
SP6 Prevotella oulorum
SP111 Prevotella sp_HMT
SP2 Prevotella sp_HMT
SP15 Prevotella sp_HMT
SP235 Prevotella sp_HMT
SP41 Prevotella sp_HMT
SP2 Prevotella fusca
SP254 Prevotella sp_HMT
SP60 Prevotella sp_HMT
SP407 Prevotella sp_HMT
SP65 Prevotella sp_HMT
SP79 Prevotella nanceiensis
SP134 Prevotella sp_HMT
SP159 Prevotella sp_HMT
SP129 Prevotella sp_HMT
SP56 Bacteroides_CIM7
SP59 Alkacoldibacter
SP225 Alcanivorans
SP32 Manihot esculenta_Oral_Taxon
SP110 Saccharibacteria_TM7
P449 Saccharibacteria_(TM7)
SP35 Saccharibacteria_TM7
SP50 Saccharibacteria_TM7
SP384 Saccharibacteria_TM7
SP391 Saccharibacteria_TM7
SP96 Saccharibacteria_(TM7)
SP130 Saccharibacteria_TM7
SP429 Desulfobulbus sp_HMT
SP47 Campylobacter
SP122 Campylobacter
SP96 Campylobacter
SP237 Campylobacter
SP105 Campylobacter
SP389 Ottowia sp_HMT
SP61 Lautropia mirabilis
SP176 Kingella oralis DLS65
SP61 Neisseria bacilliformis
SP115 Neisseria sicca
SP34 Neisseria mucosa
SP240 Neisseria flavescens
SP141 Neisseria subflava
SP88 Neisseria elongata
SP61 Neisseria sp_CIM
SP100 Eikenella corrodens
SP208 Kingella denitrificans
SP151 Cardiobacterium valvarum
SP62 Cardiobacterium hominis
SP140 Haemophilus parainfluenzae
SP385 Haemophilus sp_HMT
SP177 Haemophilus sp_HMT
SP117 Haemophilus sp_HMT
SP32 Aggregatibacter sp_HMT
SP148 Aggregatibacter
SP251 Aggregatibacter
SP168 Aggregatibacter
SP469 Aggregatibacter
SP130 Aggregatibacter
SP18 Treponema sp_HMT
SP522 Treponema sp_HMT
SP54 Treponema sp_HMT
SP78 Treponema sp_HMT
SP74 Treponema vincentii
SP46 Treponema sp_HMT
SP148 Treponema sp_HMT
SP420 Treponema sp_HMT
SP111 Treponema denticola
SP13 Treponema sp_HMT
SP183 Treponema
SP162 Treponema
SP227 Treponema
SP192 Treponema
SP63 Treponema
SP180 Treponema
SP367 Treponema
SP179 Treponema
SP412 Treponema
SP419 Treponema
SP428 Treponema
SP398 Anaerolineae
SP124 Fretibacterium fastidiosum
SP208 Fretibacterium sp_HMT
SP204 Fretibacterium sp_HMT
SP124 Fretibacterium sp_HMT
SP1 Slackia exigua
SP85 Cryptobacterium
SP383 Olsenella profusa
SP417 Olsenella sp_oral_taxon
SP189 Olsenella sp_Oral_Taxon
SP136 Atopobium sp_HMT
SP116 Atopobium sp_HMT
SP217 Lancefieldella
SP104 Atopobium rimae
SP362 Bifidobacterium
SP121 Bifidobacterium
SP186 Scardovia sp_HMT
SP367 Parascardovia
SP135 Propionibacterium
SP467 Actinomyces radicidentis
SP288 Arachnia propionica
SP14 Corynebacterium
SP414 Corynebacterium
SP68 Corynebacterium
SP118 Rothia mucilaginosa
SP465 Rothia aeria
SP19 Rothia dentocariosa
SP422 Peptostreptococcaceae
SP469 Peptococcus
SP76 Actinomyces
SP91 Schaalia gerencseriae
SP181 Actinomyces
SP212 Schaalia meyeri
SP277 Schaalia
SP139 Schaalia
SP170 Actinomyces
SP404 Schaalia
SP262 Actinomyces
SP421 Schaalia
SP361 Actinomyces
SP110 Actinomyces
SP233 Actinomyces
SP5 Actinomyces
SP172 Actinomyces
SP98 Actinomyces
SP108 Actinomyces
SP171 Actinomyces
SP172 Actinomyces
SP105 Actinomyces
SP249 Actinomyces
SP62 Actinomyces
SP147 Actinomyces
SP101 Actinomyces
SP160 Actinomyces
SP63 Fusobacterium nucleatum
SP173 Fusobacterium sp_HMT
SP184 Fusobacterium sp_HMT
SP359 Fusobacterium sp_HMT
SP297 Fusobacterium sp_HMT
SP19 Fusobacterium nucleatum
SP84 Fusobacterium sp_HMT
SP20 Fusobacterium sp_HMT
SP13 Fusobacterium sp_HMT
SP102 Fusobacterium sp_HMT
SP124 Fusobacterium sp_HMT
SP191 Leptotrichia sp_HMT
SP207 Leptotrichia sp_HMT
P86 Leptotrichia sp_HMT
SP125 Leptotrichia
SP158 Leptotrichia
SP171 Leptotrichia
SP26 Leptotrichia sp_HMT
SP148 Leptotrichia
SP22 Leptotrichia hofstadii
SP161 Leptotrichia
SP8 Leptotrichia
SP229 Leptotrichia
SP9 Leptotrichia
SP420 Leptotrichia
SP79 Leptotrichia
SP18 Leptotrichia
SP185 Rumminococcaceae_Oral_Taxon
SP188 Ruminococcaceae_Oral_Taxon
SP81 Lachnospiraceae_ZQ-D1
SP392 Butyrivibrio sp_HMT
SP466 Catonella sp_HMT
SP118 Catonella morbi
SP541 Oribacterium sinus
SP213 Oribacterium sp_HMT
SP142 Oribacterium sp_HMT
SP378 Shuttleworthia satiella
SP544 Stomatobaculum longum
SP105 Stomatobaculum longum
SP75 Stomatobaculum longum
SP183 Lachnospiraceae_ZQ-D1
SP11 Lachnospiraceae_ZQ-D1
SP91 Lachnospiraceae_ZQ-D1
SP288 Johnsonella
SP165 Johnsonella sp_HMT
SP347 Lachnoanaerobacter
SP347 Lachnoanaerobacter
SP139 Lachnoanaerobacter
SP18 Lachnoanaerobacter
SP403 Clostridiales_ZQ-D1
SP31 Peptoniphilus
SP158 Peptoniphilus sp_HMT
SP53 Peptoniphilus sp_HMT
SP53 Peptoniphilus sp_HMT
SP230 Peptococcus sp_HMT
SP236 Peptococcus sp_HMT
SP234 Filifactor alocis
SP188 uncultured
SP262 Eubacterium
SP71 Peptoniphilus
SP296 Peptostreptococcaceae
SP98 Peptostreptococcaceae
SP114 Peptostreptococcaceae
SP153 Peptostreptococcaceae
SP64 Peptostreptococcaceae
SP418 Peptostreptococcaceae
SP152 Peptostreptococcaceae
SP423 Peptostreptococcaceae
SP28 Mogibacterium
SP281 Mogibacterium
SP58 Mogibacterium
SP468 Dialister sp_HMT
SP28 Dialister invisus
SP12 Veillonella sp_HMT
SP116 Veillonella atypica
SP98 Veillonella parvula
SP82 Megasphaera
SP287 Anaerogloboides
SP129 Megasphaera
SP283 Collinsella
SP176 Veillonella
SP180 Veillonella
SP197 Veillonella
SP150 Veillonella
SP6 Veillonella
SP1 Selenomonas
SP128 Selenomonas
SP129 Selenomonas
SP153 Selenomonas
SP153 Selenomonas
SP386 Centipeda
SP343 Mycoplasma
SP416 Eggerthella
SP29 Rothia
SP145 Erysipelotrichaceae
SP473 Erysipelotrichaceae
SP279 Solobacterium
SP168 Bulleidia extructa
SP50 Gemella bergeri
SP63 Gemella morbillorum
SP132 Gemella morbillorum
SP130 Gemella haemolysans
SP61 Gemella sanguinis
SP180 Lactobacillus salivarius
SP77 Lactobacillus
SP236 Atopobiaceae defecta
SP66 Granulicatella elegans
SP187 Granulicatella adiacens
SP108 Granulicatella paraadiacens
SP126 Streptococcus
SP246 Streptococcus
SP175 Streptococcus
SP84 Streptococcus cristatus
SP53 Streptococcus
SP198 Streptococcus
SP54 Streptococcus
SP174 Streptococcus
SP390 Streptococcus
SP59 Streptococcus
SP64 Streptococcus
SP15 Streptococcus
SP19 Streptococcus
SP46 Streptococcus
SP5 Streptococcus
SP141 Streptococcus
SP12 Streptococcus
SP40 Streptococcus
SP83 Streptococcus
SP183 Streptococcus
SP96 Streptococcus
SP56 Streptococcus
SP53 Streptococcus
SP12 Streptococcus
SP14 Streptococcus
SP12 Streptococcus
Nova_16_Metaviro_F1
Novatine_A_F1
Novatine_A_F2
Nova_12_Metaviro_F1
DSS_24_Metaviro_F1
DSS_24_Metaviro_F2
Novatine_Sm_F1
Novatine_Sm_F2
Strother_21_Metaviro_F1
Strother_21_Metaviro_F2
DSS_12_Metaviro_F1
Novatine_B_F2
Strother_12_Metaviro_F2
DSS_12_Metaviro_F2
Nova_12_Metaviro_F2
DSS_24_Metaviro_F2
Novatine_B_F1
DSS_12_Metaviro_F1
Nova_24_Metaviro_F1
Nova_24_Metaviro_F2
0.2
0.1
0.5
0.75
0.9
0.95
1
Classification/site