SP74 Alcoscandavirus_[SR1_JG-1]:bacterium_HMT_351 (0.0%0.0%0.39%0.81%)
SP20 Alcoscandavirus_[SR1_JG-1]:bacterium_HMT_351 (0.0%0.0%0.24%0.45%0.2%)
SP12 Alcoscandavirus_[SR1_JG-1]:bacterium_HMT_351 (0.0%0.0%0.0%0.0%)
SP63 Gracilibacteria_[GN02_JG-2]:bacterium_HMT_871 (0.0%0.0%0.0%0.0%)
SP53 Gracilibacteria_[GN02_JG-2]:bacterium_HMT_871 (0.0%0.0%0.0%0.0%)
SP60 Saccharibacteria_[TM7_JG-1]:bacterium_HMT_348 (0.0%0.0%0.0%0.0%0.0%0.0%)
SP52 Saccharibacteria_[TM7_JG-1]:bacterium_HMT_348 (0.0%0.0%0.0%0.0%0.0%0.0%)
SP28 Saccharibacteria_[TM7_JG-1]:bacterium_HMT_351 (0.0%0.0%0.0%0.0%0.0%0.0%)
SP12 Saccharibacteria_[TM7_JG-1]:bacterium_HMT_351 (0.0%0.0%0.0%0.0%0.0%0.0%)
SP62 Saccharibacteria_[TM7_JG-1]:bacterium_HMT_348 (0.0%0.0%0.0%0.0%0.0%0.0%)
SP61 Saccharibacteria_[TM7_JG-1]:bacterium_HMT_348 (0.0%0.0%0.0%0.0%0.0%0.0%)
SP51 Saccharibacteria_[TM7_JG-1]:bacterium_HMT_348 (0.0%0.0%0.0%0.0%0.0%0.0%)
SP50 Saccharibacteria_[TM7_JG-1]:bacterium_HMT_348 (0.0%0.0%0.0%0.0%0.0%0.0%)
SP10 Saccharibacteria_[TM7_JG-1]:bacterium_HMT_352 (0%0.0%0.0%0.0%)
SP60 Saccharibacteria_[TM7_JG-1]:bacterium_HMT_351 (0.0%0.0%0.0%0.0%)
SP18 Spirosoma_rigui:bacterium_HMT_990 (0.0%0.0%0.0%0.0%)
SP19 Bergeriella_sp._HMT_290 (0.0%0.0%0.0%0.0%0.4%)
SP18 Bergeriella_sp._HMT_351 (0.0%0.0%0.0%0.0%0.0%)
SP17 Bergeriella_sp._HMT_351 (0.0%0.0%0.0%0.0%0.0%)
SP16 Bergeriella_sp._HMT_351 (0.0%0.0%0.0%0.0%0.0%)
SP48 Bergeriella_sp._HMT_351 (0.0%0.0%0.0%0.0%0.0%)
SP08 Capnocytophaga_sp._HMT_338 (0.0%0.0%0.0%0.0%0.0%)
SP13 Bacteroides_[G-1]:bacterium_HMT_803 (0.0%0.0%0.0%0.0%)
SP14 Capnocytophaga granulosa (0.1%0.0%0.1%0.1%0.5%)
SP34 Capnocytophaga gingivalis (0.1%0.0%0.0%0.0%0.0%)
SP620 Capnocytophaga haemolytica (0.0%0.0%0.0%0.0%0.0%)
SP61 Capnocytophaga leadbetter (0.0%0.0%0.0%0.0%0.0%)
SP53 Capnocytophaga sp._HMT_380 (0.0%0.0%0.0%0.0%0.0%)
SP686 Capnocytophaga sp._HMT_903 (0.0%0.0%0.0%0.0%0.0%)
SP788 Capnocytophaga sp._HMT_332 (0.0%0.0%0.0%0.0%0.0%)
SP54 Capnocytophaga leadbetter (0.0%0.0%0.0%0.0%0.0%)
SP54 Capnocytophaga sp._HMT_901 (0%0.1%0.0%0.0%)
SP52 Capnocytophaga ochracea (0.0%0.0%0.0%0.0%0.0%)
SP62 Bacteroides_[G-1]:bacterium_HMT_507 (0.0%0.0%0.0%0.0%0.0%)
SP15 Bacteroides_[G-1]:bacterium_HMT_803 (0.0%0.0%0.0%0.0%0.0%)
SP11 Bacteroides_[G-1]:bacterium_HMT_511 (0.0%0.0%0.0%0.0%0.0%)
SP63 Bacteroides_[G-1]:bacterium_HMT_803 (0.0%0.0%0.0%0.0%0.0%)
SP38 Alistipes putredinis (0.0%0.0%0.0%0.0%0.0%)
SP687 Bacteroides_[G-1]:bacterium_HMT_803 (0.0%0.0%0.0%0.0%0.0%)
SP29 Bacteroides_[G-1]:bacterium_HMT_803 (0.0%0.0%0.0%0.0%0.0%)
SP32 Bacteroides_[G-1]:bacterium_HMT_803 (0.0%0.0%0.0%0.0%0.0%)
SP189 Bacteroides_[G-1]:bacterium_HMT_281 (0.0%0.0%0.0%0.0%0.0%)
SP190 Bacteroides_[G-1]:bacterium_HMT_281 (0.0%0.0%0.0%0.0%0.0%)
SP191 Bacteroides_[G-1]:bacterium_HMT_514 (0.0%0.0%0.0%0.0%0.0%)
SP212 Porphyromonas gingivalis (0.0%0.0%0.0%0.2%)
SP618 Porphyromonas_sp._HMT_930 (1.5%0.9%0.0%0.2%)
SP279 Tannerella serpentiformis (0.0%0.0%0.0%0.0%0.0%)
SP188 Tannerella forsythia (0.0%0.0%0.1%0.1%0.0%)
SP18 Porphyromonas_sp._HMT_930 (0.0%0.0%0.0%0.0%)
SP61 Parabacteroides gordonii (0.0%0.0%0.0%0.0%)
SP58 Bacteroides uniformis (0.0%0.0%0.0%0.0%0.0%)
SP68 Prevotella vulgata (0.0%0.0%0.0%0.0%0.0%)
SP17 Prevotella dorei (0.0%0.0%0.0%0.0%0.0%)
SP703 Bacteroides ovatus/eggerthii (0.0%0.0%0.0%)
SP22 Bacteroides hepatocolens (0.0%0.0%0.0%0.0%0.0%)
SP55 Bacteroides stercoris (0.0%0.0%0.0%0.0%0.0%)
SP229 Bacteroides thetaiotaomicron (0.0%0.0%0.0%0.0%0.0%)
SP125 Bacteroides fragilis (0.0%0.0%0.0%0.0%0.0%)
SP18 Bacteroides xylanisolvens (0.0%0.0%0.0%0.0%0.0%)
SP680 Bacteroides caccae (0.0%0.0%0.0%0.0%0.0%)
SP52 Alloprevotella tannerae (0.0%0.0%0.0%0.0%0.0%)
SP617 Alloprevotella rava (0.0%0.0%0.0%0.0%0.0%)
SP62 Alloprevotella_sp._HMT_912 (0.0%0.0%0.0%0.0%0.0%)
SP448 Alloprevotella_sp._HMT_473 (0.8%0.1%0.3%0.2%0.4%)
SP140 Alloprevotella_sp._HMT_473 (0.0%0.0%0.0%0.0%0.0%)
SP134 Alloprevotella_sp._HMT_308 (0.2%0.1%0.3%0.0%0.0%)
SP62 Prevotella_sp._HMT_300 (0.0%0.0%0.0%0.0%0.0%)
SP52 Prevotella micans (0.0%0.0%0.0%0.0%0.0%)
SP48 Prevotella nanceiensis (0.0%0.0%0.0%0.0%0.0%)
SP628 Prevotella_sp._HMT_912 (0.0%0.0%0.0%0.0%0.0%)
SP39 Prevotella oris (0.0%0.0%0.0%0.0%)
SP58 Prevotella_sp._HMT_301 (0.0%0.0%0.0%0.0%0.0%)
SP48 Prevotella marshii (0.0%0.0%0.0%0.0%0.0%)
SP65 Prevotella saccharolytica (0.0%0.0%0.0%0.0%0.0%)
SP78 Prevotella oris (0.0%0.0%0.0%0.0%0.0%)
SP299 Prevotella shahii (0.0%0.0%0.0%0.0%0.0%)
SP14 Prevotella_sp._HMT_317 (0.1%0.0%0.0%0.1%0.0%)
SP465 Prevotella_sp._HMT_472 (0.0%0.0%0.0%0.0%0.0%)
SP57 Prevotella pleuritis (0.1%0.0%0.0%0.0%0.0%)
SP58 Prevotella enoeca (0.0%0.0%0.0%0.0%0.0%)
SP80 Prevotella_sp._HMT_313 (0.0%0.0%0.0%0.0%0.0%)
SP317 Prevotella buccae (0.0%0.0%0.0%0.0%0.0%)
SP60 Prevotella heparinolytica (0.0%0.0%0.0%0.0%0.0%)
SP125 Prevotella_sp._HMT_313 (0.0%0.0%0.0%0.0%0.0%)
SP51 Prevotella_sp._HMT_376 (0.0%0.0%0.0%0.0%0.0%)
SP175 Prevotella_sp._HMT_526 (0.0%0.0%0.0%0.0%0.0%)
SP68 Prevotella tannerae (0.0%0.0%0.0%0.0%0.0%)
SP52 Prevotella oulorum (0.0%0.0%0.0%0.1%0.1%)
SP58 Prevotella_sp._HMT_299 (0.0%0.0%0.0%0.0%0.0%)
SP628 Prevotella salivae (0.0%0.0%0.0%0.0%0.0%)
SP54 Prevotella melaninogenica (0.0%0.0%0.0%0.0%0.0%)
SP54 Prevotella_sp._HMT_300 (0.0%0.0%0.0%0.0%0.0%)
SP60 Prevotella_sp._HMT_306 (0.0%0.0%0.0%0.0%0.0%)
SP58 Prevotella nanceiensis (0.0%0.0%0.0%0.0%0.0%)
SP65 Prevotella pallens (0.0%0.0%0.0%0.0%0.0%)
SP58 Prevotella intermedia (0.0%0.0%0.0%0.0%0.0%)
SP183 Prevotella aurantiaca (0.0%0.0%0.0%0.0%0.0%)
SP54 Prevotella fusca (0.0%0.0%0.0%0.0%0.0%)
SP57 Prevotella multiformis (0.1%0.0%0.0%0.0%0.0%)
SP53 Prevotella dentalis (0.0%0.0%0.4%0.2%0.3%)
SP60 Prevotella melaninogenica (2.1%0.0%0.7%0.2%0.2%)
SP67 Prevotella_sp._HMT_313 (0.5%0.0%0.0%0.0%0.4%)
SP68 Prevotella nanceiensis (0.0%0.0%0.0%0.0%0.0%)
SP67 Prevotella_sp._HMT_306 (0.0%0.0%0.0%0.0%0.0%)
SP68 Prevotella oris (0.0%0.0%0.0%0.0%0.0%)
SP53 Campylobacter_[G-2]:bacterium_HMT_044 (0.0%0.0%0.0%0.0%0.0%)
SP52 Campylobacter_[G-2]:bacterium_HMT_044 (0.0%0.0%0.0%0.0%0.0%)
SP24 Campylobacter concisus (0.0%0.0%0.0%0.0%0.0%)
SP6 Campylobacter curvus (0.0%0.0%0.0%0.0%0.0%)
SP38 Anaerovibrio_[G-1]:bacterium_HMT_439 (0.0%0.0%0.0%0.0%0.0%)
SP53 Helicobacter_sp._HMT_050 (0.0%0.0%0.0%0.0%0.0%)
SP15 Flexilabrum balbiodonta (0.0%0.0%0.0%0.0%0.0%)
SP61 Synergistes jonesii (0.0%0.0%0.0%0.0%)
SP17 Pyramidobacter piscolens (0.0%0.0%0.0%0.0%)
SP56 Treponema_sp._HMT_247 (0.0%0.0%0.0%0.0%0.0%)
SP135 Treponema_sp._HMT_246 (0.0%0.0%0.0%0.0%0.0%)
SP54 Treponema denticola (0.0%0.0%0.0%0.0%0.0%)
SP63 Treponema_sp._HMT_249 (0.0%0.0%0.0%0.0%0.0%)
SP154 Treponema_sp._HMT_251 (0.0%0.0%0.0%0.0%0.0%)
SP247 Treponema_sp._HMT_256 (0.0%0.0%0.0%0.0%0.0%)
SP172 Treponema_sp._HMT_508 (0.0%0.0%0.0%0.0%0.0%)
SP172 Treponema_sp._HMT_518 (0.0%0.0%0.0%0.0%0.0%)
SP61 Treponema_sp._HMT_254 (0.0%0.0%0.0%0.0%0.0%)
SP19 Treponema_sp._HMT_252 (0.0%0.0%0.0%0.0%)
SP38 Treponema_sp._HMT_224 (0.0%0.0%0.0%0.0%0.0%)
SP86 Treponema_sp._HMT_239 (0.0%0.0%0.0%0.0%0.0%)
SP88 Treponema_sp._HMT_235 (0.0%0.0%0.0%0.0%0.0%)
SP86 Treponema_sp._HMT_242 (0.0%0.0%0.0%)
SP63 Treponema_sp._HMT_238 (0.0%0.0%0.0%0.0%0.0%)
SP119 Treponema_sp._HMT_226 (0.0%0.0%0.0%0.0%0.0%)
SP119 Treponema_sp._HMT_230 (0.0%0.0%0.0%0.0%0.0%)
SP467 Treponema_sp._HMT_237 (0.0%0.0%0.0%0.0%0.0%)
SP117 Treponema_sp._HMT_231 (0.0%0.0%0.0%0.0%0.0%)
SP58 Treponema vincentii (0.0%0.0%0.0%0.0%0.0%)
SP16 Treponema_sp._HMT_486 (0.0%0.0%0.0%0.0%)
SP32 Treponema_sp._HMT_258 (0.0%0.0%0.0%0.0%0.0%)
SP422 Treponema lecithinolyticum (0.0%0.0%0.0%0.0%0.0%)
SP33 Treponema maltophilum (0.0%0.0%0.0%0.0%0.0%)
SP32 Treponema_sp._HMT_272 (0.0%0.0%0.0%0.0%0.0%)
SP663 Treponema pectinovorum (0.0%0.0%0.0%0.0%0.0%)
SP55 Treponema_sp._HMT_250 (0.0%0.0%0.0%0.0%0.0%)
SP53 Treponema_sp._HMT_288 (0.0%0.0%0.0%0.0%0.0%)
SP52 Treponema_sp._HMT_289 (0.0%0.0%0.0%0.0%0.0%)
SP51 Treponema_sp._HMT_517 (0%0.0%0.0%0.0%)
SP38 Treponema_sp._HMT_262 (0.0%0.0%0.0%0.0%0.0%)
SP125 Treponema amylovorum (0.0%0.0%0.0%0.0%0.0%)
SP60 Treponema_sp._HMT_257 (0.1%0.1%0.2%0.0%0.0%)
SP679 Tritrichomonas foetus (0.0%0.0%0.0%0.0%0.0%)
SP51 Pseudomonas_[G-5]:bacterium_HMT_934 (0.0%0.0%0.0%0.0%0.0%)
SP50 Pseudomonas_[G-5]:bacterium_HMT_934 (0.0%0.0%0.0%0.0%0.0%)
SP35 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP34 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP33 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP32 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP31 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP30 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP29 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP28 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP27 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP26 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP25 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP24 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP23 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP22 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP21 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP20 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP19 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP18 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP17 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP16 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP15 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP14 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP13 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP12 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP11 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP10 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP9 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP8 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP7 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP6 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP5 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP4 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP3 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP2 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP1 Neisseria_[G-1]:bacterium_HMT_932 (0.0%0.0%0.0%0.0%0.0%)
SP7 Pseudomonas_[G-5]:bacterium_HMT_934 (0.0%0.0%0.0%0.0%0.0%)
SP6 Pseudomonas_[G-5]:bacterium_HMT_934 (0.0%0.0%0.0%0.0%0.0%)
SP5 Pseudomonas_[G-5]:bacterium_HMT_934 (0.0%0.0%0.0%0.0%0.0%)
SP4 Pseudomonas_[G-5]:bacterium_HMT_934 (0.0%0.0%0.0%0.0%0.0%)
SP3 Pseudomonas_[G-5]:bacterium_HMT_934 (0.0%0.0%0.0%0.0%0.0%)
SP2 Pseudomonas_[G-5]:bacterium_HMT_934 (0.0%0.0%0.0%0.0%0.0%)
SP1 Pseudomonas_[G-5]:bacterium_HMT_934