

<b>Species found in BIOTA water sample</b>	<b>Percentage of species</b>
cyanobacterium spp.	15.5552731893265
spirogyra maxima	10.7369758576874
luteolibacter sp.	10.3693350275307
microcystis spp.	5.44430326132994
cymatopleura solea	4.53960186361711
asterionellopsis glacialis	3.83650995340957
luteolibacter spp.	2.20076238881829
erysipelothrix spp.	1.78568403218975
fluvicola spp.	1.45870393900889
solitalea spp.	1.38924184667513
phaeodactylum tricornutum	1.33079203727234
geobacter spp.	1.30622617534942
rhodobacter spp.	1.15290131300296
rhodobacter capsulatus	0.929267259635747
lewinella spp.	0.864887759423972
anaerophaga spp.	0.808132147395171
prolixibacter spp.	0.765777213045319
dendrobium officinale	0.716645489199491
malikia spp.	0.670902160101651
flavobacterium spp.	0.657348581109699
leptothrix sp.	0.630241423125794
porticoccus spp.	0.621770436255823
alkanindiges spp.	0.57687420584498
ohtaekwangia spp.	0.573485811096992
pedosphaera parvula	0.56840321897501
methyllobacter spp.	0.55993223210504
pedobacter spp.	0.546378653113087
cytophaga spp.	0.524354087251164
rhodobacter sp.	0.521812791190173
candidatus magnospira bakii	0.518424396442185
thalassiosira pseudonana	0.507412113511224
prosthecobacter spp.	0.500635324015247
arcicella sp.	0.498094027954256
dechloromonas denitrificans	0.470986869970351
curvibacter spp.	0.452350698856416
caedibacter spp.	0.443032613299449
oscillatoria limosa	0.425243540872511
rhodocyclus tenuis	0.403218975010588
verrucomicrombium spp.	0.378653113087674
sphingobacterium spp.	0.37695891571368
haliscomenobacter hydrossis	0.360016941973739
opitutus spp.	0.352393053790766
terrimonas sp.	0.349004659042778
sediminibacterium spp.	0.343074968233799
candidatus nardonella endosymbiont of scyphophorus yuccae	0.340533672172808
acidovorax facilis	0.333756882676831
peziza succosa	0.332909783989834
terrimonas spp.	0.316814908936891
flexibacter spp.	0.306649724692926

phaeodactylum sp.	0.29987293519695
paucibacter toxinivorans	0.29987293519695
methylotenera versatilis	0.295637441761965
solitalea canadensis	0.293943244387971
sphingomonas spp.	0.281236764083015
acanthamoeba castellanii	0.275307073274036
dechloromonas spp.	0.270224481152054
aquimonas spp.	0.265141889030072
flavobacterium sp.	0.256670902160101
methylotenera spp.	0.255823803473104
novosphingobium spp.	0.253282507412113
sphingopyxis sp.	0.249047013977128
haliscomenobacter spp.	0.246505717916137
chitinophaga spp.	0.246505717916137
lacibacter spp.	0.243964421855146
bacteroides spp.	0.232952138924184
roseimicrobium gellanilyticum	0.23125794155019
novosphingobium subarcticum	0.229563744176196
peredibacter starrii	0.228716645489199
leptothrix spp.	0.223634053367217
sphaerotilus spp.	0.221939855993223
algoriphagus hongiella halophila	0.203303684879288
acidobacterium spp.	0.201609487505294
nitrosovibrio spp.	0.195679796696315
armatimonas rosea	0.193985599322321
rheinheimera texana	0.193138500635324
actibacter spp.	0.193138500635324
inhella inkyongensis	0.193138500635324
gemmaimonas sp.	0.184667513765353
luteolibacter yonseiensis	0.182126217704362
luteolibacter algae	0.181279119017365
phytophthora infestans	0.173655230834392
opitutus sp.	0.173655230834392
burkholderia spp.	0.168572638712409
pedosphaera spp.	0.164337145277424
arenimonas sp.	0.155019059720457
bacillaria paxillifer	0.152477763659466
bdellovibrio spp.	0.152477763659466
pelobacter spp.	0.148242270224481
novosphingobium stygium	0.144006776789495
arthrobacter spp.	0.130453197797543
methylomonas methanica	0.127911901736552
geobacter pelophilus	0.126217704362558
clostridium spp.	0.124523506988564
aquabacterium spp.	0.123676408301567
emticicia sp.	0.11689961880559
methylibium spp.	0.114358322744599
xanthomonas spp.	0.114358322744599
nannocystis spp.	0.113511224057602
zhongshania spongiibacter sp.	0.105040237187632

pontibacter salisaro	0.0991105463786531
hydrogenophaga atypica	0.096569250317662
pseudospirillum spp.	0.0931808555696738
aeromonas sp.	0.0906395595086827
methylovulum miyakonense	0.0897924608216857
flavobacterium cheniae	0.0897924608216857
dehalococcoides spp.	0.0864040660736975
ochrobactrum grignonense	0.0864040660736975
aquimonas voraii	0.0855569673867005
spirochaeta spp.	0.0855569673867005
novosphingobium sp.	0.0838627700127064
ehrlichia spp.	0.0830156713257094
gemmatimonas spp.	0.0804743752647183
tolumonas spp.	0.0787801778907242
undibacterium sp.	0.0770859805167301
haliea spp.	0.0770859805167301
bdellovibrio sp.	0.0753917831427361
pseudomonas spp.	0.074544684455739
hirschia sp.	0.073697585768742
crenothrix polyspora	0.073697585768742
pseudomonas putida	0.072850487081745
hydrogenophaga spp.	0.072850487081745
ruberivivax gelatinosus	0.0720033883947479
candidatus zinderia insecticola	0.0711562897077509
haslea ostrearia	0.0694620923337568
phacus orbicularis	0.0686149936467598
bacteriovorax spp.	0.0677678949597628
diatoma tenuis	0.0669207962727657
algoriphagus aquatilis	0.0660736975857687
spirobacillus cienkowskii	0.0652265988987717
planctomyces spp.	0.0635324015247776
malawimonas jakobiformis	0.0635324015247776
porphyrobacter sp.	0.0635324015247776
melosira varians	0.0635324015247776
rubrobacter spp.	0.0635324015247776
desulfobulbus spp.	0.0635324015247776
methylophilus spp.	0.0626853028377806
sphingomonas sp.	0.0626853028377806
ochromonas distigma	0.0618382041507835
nitrosospira spp.	0.0618382041507835
adhaeribacter spp.	0.0618382041507835
amphora ovalis	0.0609911054637865
caulobacter vibrioides	0.0601440067767894
methylosoma sp.	0.0576027107157983
candidatus rhodoluna rhodoluna sp. kas9	0.0567556120288013
crocinitomix spp.	0.0559085133418043
steroidobacter spp.	0.0559085133418043
sulfurisoma sediminicola	0.0550614146548072
rickettsia canadensis	0.0550614146548072
bacteriovorax sp.	0.0550614146548072

balneola sp.	0.0533672172808132
candidatus solibacter uncultured solibacter sp.	0.0533672172808132
anaeromyxobacter spp.	0.0525201185938161
leadbetterella sp.	0.049978822532825
phacus granum	0.049978822532825
perlucidibaca piscinae	0.049978822532825
aquincola tertiaricarbonis	0.049131723845828
paludibacter spp.	0.049131723845828
ideonella sp.	0.049131723845828
ponticoccus sp.	0.048284625158831
rhizobium sp.	0.048284625158831
marinobacter spp.	0.048284625158831
arcicella spp.	0.0474375264718339
herbaspirillum spp.	0.0474375264718339
cosmarium botrytis	0.0457433290978398
sphingopyxis spp.	0.0457433290978398
halochromatium sp.	0.0448962304108428
uncultured candidatus pelagibacter sp.	0.0448962304108428
geobacter luticola	0.0448962304108428
ignavibacterium spp.	0.0448962304108428
lactuca sativa	0.0440491317238458
sporocytophaga spp.	0.0440491317238458
cymbella subturgidula	0.0432020330368487
chitinimonas koreensis	0.0423549343498517
rhodobacter blasticus	0.0423549343498517
runella slithyformis	0.0415078356628547
gemmata spp.	0.0406607369758576
legionella spp.	0.0406607369758576
pirellula spp.	0.0406607369758576
rheinheimera chironomi	0.0398136382888606
altererythrobacter spp.	0.0398136382888606
hyphomicrobium spp.	0.0398136382888606
xylella spp.	0.0398136382888606
smithella spp.	0.0389665396018636
candidatus thiopilula aggregata	0.0381194409148665
rhodomicrobiun spp.	0.0381194409148665
aquicella spp.	0.0381194409148665
spiribacter sp.	0.0381194409148665
oleiphilus spp.	0.0372723422278695
thiothrix spp.	0.0372723422278695
alistipes massiliensis	0.0372723422278695
parasegetibacter sp.	0.0372723422278695
candidatus neoehrlichia mikurensis	0.0364252435408725
orientia tsutsugamushi	0.0364252435408725
pelobacter carbinolicus	0.0364252435408725
candidatus regiella insecticola	0.0364252435408725
bradyrhizobium spp.	0.0355781448538754
acidovorax delafieldii	0.0347310461668784
thioalkalispira spp.	0.0347310461668784
polyangium sp.	0.0338839474798814

saccharophagus spp.	0.0338839474798814
lupinus luteus	0.0338839474798814
tissierella spp.	0.0338839474798814
luteolibacter cuticulihirudinis	0.0330368487928843
hymenobacter sp.	0.0330368487928843
leptothrix mobilis	0.0330368487928843
hydrogenophaga sp.	0.0330368487928843
limnobacter spp.	0.0321897501058873
cytophaga xylolyticus	0.0321897501058873
mesorhizobium spp.	0.0321897501058873
cystobacter miniatus	0.0321897501058873
rhodoplanes spp.	0.0313426514188903
bdellovibrio bacteriovorus	0.0304955527318932
arcobacter spp.	0.0304955527318932
cellvibrio fulvus	0.0304955527318932
pelotomaculum spp.	0.0304955527318932
pontibacter spp.	0.0304955527318932
fusibacter sp.	0.0296484540448962
monomorphina pyrum	0.0296484540448962
algoriphagus sp.	0.0296484540448962
longilinea spp.	0.0296484540448962
navicula sp.	0.0288013553578991
lewinella sp.	0.0288013553578991
marinoscillum furvescens	0.0288013553578991
sterolibacterium spp.	0.0288013553578991
phacus euglena limnophila	0.0288013553578991
gemmatimonas aurantiaca	0.0279542566709021
chitinimonas taiwanensis	0.0279542566709021
desulfobacterium spp.	0.0279542566709021
lewinella marina	0.0279542566709021
algoriphagus dokdonensis	0.0271071579839051
fusibacter spp.	0.0271071579839051
arenimonas spp.	0.0271071579839051
uncultured candidatus odyssella sp.	0.026260059296908
thiobacillus spp.	0.025412960609911
denitratisoma spp.	0.025412960609911
paludibacter propionicigenes	0.025412960609911
candidatus vidania fulgoroideae	0.025412960609911
mesotaenium caldariorum	0.024565861922914
nostoc sp.	0.024565861922914
syntrophomonas spp.	0.0237187632359169
anaerolinea spp.	0.0237187632359169
desulfocapsa spp.	0.0237187632359169
nitrospira spp.	0.0237187632359169
ferribacterium spp.	0.0237187632359169
turicibacter spp.	0.0237187632359169
pedobacter sp.	0.0228716645489199
hyphomonas spp.	0.0228716645489199
pontibacter korlensis	0.0228716645489199
bellilinea spp.	0.0228716645489199

alkaliflexus spp.	0.0228716645489199
oscillatoria sp.	0.0228716645489199
microvirga spp.	0.0228716645489199
cytophaga hutchinsonii	0.0228716645489199
rhodoferax spp.	0.0228716645489199
lishizhenia caseinilytica	0.0220245658619229
chlorobium spp.	0.0220245658619229
dechloromonas aromatica	0.0220245658619229
desulfuromonas spp.	0.0211774671749258
tistrella spp.	0.0211774671749258
brucella spp.	0.0211774671749258
candidatus carsonella primary endosymbiont of psylla floccosa	0.0211774671749258
bacillus spp.	0.0203303684879288
perlucidibaca sp.	0.0203303684879288
thermomonas spp.	0.0203303684879288
uncultured candidatus solibacter sp.	0.0203303684879288
candidatus carsonella ruddii	0.0203303684879288
lewinella cohaerens	0.0203303684879288
owenweeksia spp.	0.0203303684879288
thiobacillus sp.	0.0194832698009318
limnohabitans curvus	0.0194832698009318
thermomonas brevis	0.0194832698009318
sulfuritalea spp.	0.0194832698009318
sellaphora blackfordensis	0.0194832698009318
acidovorax temperans	0.0186361711139347
thiorhodospira spp.	0.0186361711139347
solirubrobacter spp.	0.0186361711139347
clostridium sp.	0.0186361711139347
allobaculum sp.	0.0186361711139347
francisella spp.	0.0177890724269377
sandarakinorhabdus spp.	0.0177890724269377
methylobacillus spp.	0.0177890724269377
acanthamoeba sp.	0.0177890724269377
fibrobacter spp.	0.0177890724269377
lepoctenoides ovum	0.0177890724269377
phormidium tergestinum	0.0177890724269377
hydrogenophaga taeniospiralis	0.0169419737399407
rhizobium spp.	0.0169419737399407
georgfuchsia toluolica	0.0169419737399407
brevinema spp.	0.0169419737399407
devosia spp.	0.0169419737399407
leptolyngbya spp.	0.0160948750529436
holophaga spp.	0.0160948750529436
beggiatoa sp.	0.0160948750529436
chthoniobacter flavus	0.0160948750529436
cytophaga aurantiaca	0.0160948750529436
conexibacter spp.	0.0160948750529436
fluviimonas pallidilutea	0.0152477763659466
azospirillum spp.	0.0152477763659466
rhodoferax albidiferax sp.	0.0152477763659466

methylsinus spp.	0.0152477763659466
acidisphaera spp.	0.0152477763659466
lepoctenoides euglena spirogyra	0.0152477763659466
rickettsia spp.	0.0152477763659466
lysobacter spp.	0.0152477763659466
campylobacter canadensis	0.0152477763659466
methermicoccus sp.	0.0152477763659466
alcanivorax sp.	0.0152477763659466
candidatus metachlamydia lacustris	0.0152477763659466
denitromonas spp.	0.0152477763659466
alteromonas sp.	0.0152477763659466
methylomonas sp.	0.0152477763659466
bradyrhizobium sp.	0.0144006776789495
beijerinckia spp.	0.0144006776789495
caedibacter macronucleorum	0.0144006776789495
methylobacter sp.	0.0144006776789495
geothrix spp.	0.0144006776789495
chitinimonas sp.	0.0144006776789495
saccharospirillum spp.	0.0144006776789495
bacillus sp.	0.0135535789919525
rhodocista spp.	0.0135535789919525
euglena stellata	0.0135535789919525
desulfomonile spp.	0.0135535789919525
skermanella sp.	0.0135535789919525
kaistobacter spp.	0.0135535789919525
sphingobacterium sp.	0.0135535789919525
candidatus midichloria sp. hymares3	0.0135535789919525
syntrophus spp.	0.0135535789919525
bacteroides sp.	0.0135535789919525
wolbachia spp.	0.0127064803049555
parasegetibacter luojiensis	0.0127064803049555
pseudonocardia alaniniphila	0.0127064803049555
cellvibrio ostraviensis	0.0127064803049555
lactobacillus alimentarius	0.0127064803049555
luteolibacter luojiensis	0.0127064803049555
acetobacter spp.	0.0127064803049555
enchelyodon sp.	0.0127064803049555
parvularcula spp.	0.0127064803049555
acidovorax spp.	0.0127064803049555
cytophaga sp.	0.0127064803049555
allochromatium vinosum	0.0127064803049555
cryptomonas pyrenoidifera	0.0118593816179584
methylotilis spp.	0.0118593816179584
bradyrhizobium lupini	0.0118593816179584
rheinheimera sp.	0.0118593816179584
rhodopseudomonas spp.	0.0118593816179584
pseudomonas sp.	0.0118593816179584
azotobacter spp.	0.0118593816179584
sneathiella spp.	0.0118593816179584
xylella fastidiosa	0.0118593816179584

arthrobacter ramosus	0.0118593816179584
gluconobacter oxydans	0.0118593816179584
prosthecobacter debontii	0.0118593816179584
woodsholea maritima	0.0118593816179584
thiohalomonas denitrificans	0.0118593816179584
gyrosigma limosum	0.0118593816179584
sterolibacterium sp.	0.0110122829309614
spongiibacter spp.	0.0110122829309614
jahnella thaxteri	0.0110122829309614
discoplastis euglena spathirhyncha	0.0110122829309614
thiocystis violacea	0.0110122829309614
bdellovibrio exovorus	0.0110122829309614
desulfomicrobium spp.	0.0110122829309614
candidatus xiphinematobacter brevicolli	0.0110122829309614
porticoccus hydrocarbonoclasticus	0.0110122829309614
verrucomicrobium sp.	0.0110122829309614
borrelia carolinensis	0.0101651842439644
herminiimonas sp.	0.0101651842439644
prosthecobacter vanneervenii	0.0101651842439644
treponema spp.	0.0101651842439644
levilinea spp.	0.0101651842439644
serratia spp.	0.0101651842439644
euglenaria euglena anabaena	0.0101651842439644
saccharothrix spp.	0.0101651842439644
marichromatium sp.	0.0101651842439644
candidatus nasuia deltocephalincola	0.0101651842439644
mycobacterium pulveris	0.0101651842439644
euglena cuneata	0.0101651842439644
euglenaformis euglena proxima	0.0101651842439644
flexibacter sp.	0.0101651842439644
sandarakinorhabdus sp.	0.0101651842439644
haliscomenobacter sp.	0.00931808555696738
rhodopirellula spp.	0.00931808555696738
guillardia theta	0.00931808555696738
halomonas spp.	0.00931808555696738
nevskia ramosa	0.00931808555696738
strombomonas acuminata	0.00931808555696738
stenotrophomonas maltophilia	0.00931808555696738
riftia sp.	0.00931808555696738
desulforhopalus spp.	0.00931808555696738
lysobacter daejeonensis	0.00931808555696738
stigmatella spp.	0.00931808555696738
leporinclus euglena tripteris	0.00931808555696738
nitrosococcus spp.	0.00931808555696738
methanobrevibacter sp.	0.00931808555696738
aquabacterium sp.	0.00931808555696738
monomorphina striata	0.00931808555696738
cloacibacterium spp.	0.00931808555696738
tolumonas auensis	0.00931808555696738
ignavibacterium sp.	0.00847098686997035

mitsuaria sp.	0.00847098686997035
ralstonia spp.	0.00847098686997035
roseomonas lacus	0.00847098686997035
syntrophobacter spp.	0.00847098686997035
ichthyobodo sp.	0.00847098686997035
thiorhodovibrio spp.	0.00847098686997035
beggiatoa spp.	0.00847098686997035
georgfuchsia spp.	0.00847098686997035
methylocapsa aurea	0.00847098686997035
kitasatospora spp.	0.00847098686997035
desulfovibrio spp.	0.00847098686997035
sediminibacterium sp.	0.00847098686997035
ferruginibacter sp.	0.00847098686997035
sphingorhabdus marina	0.00847098686997035
nitrobacter spp.	0.00847098686997035
fucus vesiculosus	0.00762388818297331
candidatus haliscomenobacter calcifugiens	0.00762388818297331
alkaliflexus imshenetskii	0.00762388818297331
ideonella spp.	0.00762388818297331
blastopirellula spp.	0.00762388818297331
flavihumibacter sp.	0.00762388818297331
blastococcus spp.	0.00762388818297331
euglena deses	0.00762388818297331
moorella spp.	0.00762388818297331
lacibacterium sp.	0.00762388818297331
thalassomonas sp.	0.00762388818297331
aetherobacter sp.	0.00762388818297331
monomorphina ovata	0.00762388818297331
chlamydia spp.	0.00762388818297331
pinnularia acrosphaeria	0.00762388818297331
propionivibrio spp.	0.00762388818297331
vogesella indigofera	0.00762388818297331
rickettsia sp.	0.00762388818297331
flexibacter flexilis	0.00762388818297331
altererythrobacter alterierythrobacter sp.	0.00762388818297331
nitratireductor spp.	0.00677678949597628
alkaliphilus transvaalensis	0.00677678949597628
paracraurococcus spp.	0.00677678949597628
leptospira spp.	0.00677678949597628
candidatus anadelfobacter veles	0.00677678949597628
uncultured candidatus competitibacter sp.	0.00677678949597628
parabacteroides distasonis	0.00677678949597628
saprospira grandis	0.00677678949597628
oxalobacter spp.	0.00677678949597628
kribbella spp.	0.00677678949597628
cellvibrio gandavensis	0.00677678949597628
achromobacter spp.	0.00677678949597628
haliangium spp.	0.00677678949597628
siluania monomastiga	0.00677678949597628
dyella spp.	0.00677678949597628

pedomicrobium spp.	0.00677678949597628
chondromyces apiculatus	0.00677678949597628
chelatococcus sp.	0.00677678949597628
acidaminobacter sp.	0.00677678949597628
tetrasphaera sp.	0.00677678949597628
stigeoclonium helveticum	0.00677678949597628
gordonibacter spp.	0.00677678949597628
youngiibacter multivorans	0.00677678949597628
leptospira interrogans	0.00592969080897924
thalassiosira sp.	0.00592969080897924
pseudochrobactrum sp.	0.00592969080897924
bacillus amyloliquefaciens	0.00592969080897924
legionella pneumophila	0.00592969080897924
sulfuricella spp.	0.00592969080897924
thioalkalivibrio spp.	0.00592969080897924
thioalkalibacter halophilus	0.00592969080897924
chitinophaga sp.	0.00592969080897924
desulfococcus spp.	0.00592969080897924
candidatus accumulibacter sp. jj007	0.00592969080897924
limnothrix redekei	0.00592969080897924
actinoplanes friuliensis	0.00592969080897924
leptolyngbya sp.	0.00592969080897924
ferruginibacter spp.	0.00592969080897924
dongia spp.	0.00592969080897924
uroglena sp.	0.00592969080897924
thermoanaerobacter sulfurigignens	0.00592969080897924
niastella spp.	0.00592969080897924
flavisolibacter sp.	0.00592969080897924
acidovorax sp.	0.00592969080897924
petrimonas spp.	0.00592969080897924
roseateles aquatilis	0.00508259212198221
desulfitobacterium spp.	0.00508259212198221
euglena schmitzii	0.00508259212198221
massilia timonae	0.00508259212198221
terrigraphus spp.	0.00508259212198221
turneriella parva	0.00508259212198221
phormidium tenuie	0.00508259212198221
rhodovastum spp.	0.00508259212198221
clostridium ruminantium	0.00508259212198221
uncultured candidatus ruthia sp.	0.00508259212198221
zoogloea spp.	0.00508259212198221
achromobacter sp.	0.00508259212198221
pseudoxanthomonas koreensis	0.00508259212198221
navicula cryptocephala	0.00508259212198221
nordella spp.	0.00508259212198221
deinococcus spp.	0.00508259212198221
streptomyces glaucescens	0.00508259212198221
rhodocista sp.	0.00508259212198221
phenyllobacterium sp.	0.00508259212198221
fulvivirga sp.	0.00508259212198221

rickettsia japonica	0.00508259212198221
aquaspirillum sp.	0.00508259212198221
sinorhizobium sp.	0.00508259212198221
pseudonocardia spp.	0.00508259212198221
saccharofermentans acetigenes	0.00508259212198221
segetibacter spp.	0.00423549343498517
myriangium duriaeae	0.00423549343498517
hellea balneolensis	0.00423549343498517
spirogyra sp.	0.00423549343498517
tabrizicola aquatica	0.00423549343498517
lacibacterium rhodospirillaceae bacterium	0.00423549343498517
legionella sp.	0.00423549343498517
algoriphagus marincola	0.00423549343498517
acidiphilum spp.	0.00423549343498517
flectobacillus spp.	0.00423549343498517
clostridium bowmanii	0.00423549343498517
methylococcus spp.	0.00423549343498517
lutibacter spp.	0.00423549343498517
rubellimicrobium spp.	0.00423549343498517
jeotgalicoccus psychrophilus	0.00423549343498517
anaerovorax spp.	0.00423549343498517
butyrivibrio spp.	0.00423549343498517
niastella sp.	0.00423549343498517
euglena cantabrica	0.00423549343498517
rhodococcus spp.	0.00423549343498517
saccharophagus degradans	0.00423549343498517
lewinella antarctica	0.00423549343498517
nitritalea halalkaliphila	0.00423549343498517
desulfovaba gelida	0.00423549343498517
acinetobacter radioresistens	0.00423549343498517
ilumatobacter spp.	0.00423549343498517
caldilinea spp.	0.00423549343498517
rhodoferax antarcticus	0.00423549343498517
halothiobacillus kellyi	0.00423549343498517
mycoplasma sp.	0.00423549343498517
sporocytophaga myxococcoides	0.00338839474798814
marivirga tractuosa	0.00338839474798814
syntrophus sp.	0.00338839474798814
actinomycetospora sp.	0.00338839474798814
desulfatitalea tepidiphila	0.00338839474798814
rhodomicrobium sp.	0.00338839474798814
gallionella spp.	0.00338839474798814
phyllobacterium sp.	0.00338839474798814
gembmobacter spp.	0.00338839474798814
streptomyces xanthophaeus	0.00338839474798814
oscillatoria spp.	0.00338839474798814
ekhidna sp.	0.00338839474798814
anoxybacillus spp.	0.00338839474798814
desulfovirga spp.	0.00338839474798814
vulgatibacter incomptus	0.00338839474798814

gemmata sp.	0.00338839474798814
mycoplana sp.	0.00338839474798814
desulfuromusa spp.	0.00338839474798814
preussia typharum	0.00338839474798814
alishewanella sp.	0.00338839474798814
hyphomicrobium denitrificans	0.00338839474798814
luminiphilus syltensis	0.00338839474798814
hirschia spp.	0.00338839474798814
subdoligranulum spp.	0.00338839474798814
flexibacter ruber	0.00338839474798814
rhizomicrobium spp.	0.00338839474798814
alterococcus agarolyticus	0.00338839474798814
thiohalophilus spp.	0.00338839474798814
nocardoides aquiterrae	0.00338839474798814
caenispirillum sp.	0.00338839474798814
geobacter metallireducens	0.00338839474798814
candidatus nitrososphaera	0.00338839474798814
mycoplasma canadense	0.00338839474798814
herminiiimonas saxobsidens	0.00338839474798814
algibacter lectus	0.00338839474798814
microbulbifer donghaiensis	0.00338839474798814
enterobacter spp.	0.00338839474798814
azospirillum sp.	0.00338839474798814
sporomusa spp.	0.00338839474798814
roseomonas stagni	0.0025412960609911
pontibacter sp.	0.0025412960609911
roseomonas sp.	0.0025412960609911
caulobacter spp.	0.0025412960609911
flavisolibacter spp.	0.0025412960609911
caulobacter fusiformis	0.0025412960609911
ornatilinea apprima	0.0025412960609911
ilumatobacter fluminis	0.0025412960609911
herbaspirillum sp.	0.0025412960609911
chloridium chlorella saccharophila	0.0025412960609911
chelatococcus spp.	0.0025412960609911
echinicola spp.	0.0025412960609911
rheinheimera soli	0.0025412960609911
nitrospina spp.	0.0025412960609911
coxiella spp.	0.0025412960609911
candidatus tremblaya phenacola	0.0025412960609911
fusobacterium spp.	0.0025412960609911
zavarzinella spp.	0.0025412960609911
mortierella microzygospora	0.0025412960609911
acetivibrio spp.	0.0025412960609911
thauera sp.	0.0025412960609911
mucilaginibacter spp.	0.0025412960609911
chloroflexus spp.	0.0025412960609911
methanosaeta sp.	0.0025412960609911
nordella oligomobilis	0.0025412960609911
candidatus alysiosphaera europeae	0.0025412960609911

aquiflexum spp.	0.0025412960609911
altererythrobacter ishigakiensis	0.0025412960609911
pelagibius spp.	0.0025412960609911
sulfuricurvum kujiense	0.0025412960609911
geothermobacter spp.	0.0025412960609911
kocuria rosea	0.0025412960609911
candidatus aquirestis calciphila	0.0025412960609911
plesiocystis spp.	0.0025412960609911
haliea sp.	0.0025412960609911
actinomadura spp.	0.0025412960609911
chitinophaga pinensis	0.0025412960609911
rhodospirillum sp.	0.0025412960609911
formivibrio citricus	0.0025412960609911
sulfurovum spp.	0.0025412960609911
borrelia garinii	0.0025412960609911
candidatus trichorickettsia mobilis	0.0025412960609911
flavisolibacter ginsengisoli	0.0025412960609911
methyllobacterium spp.	0.0025412960609911
pseudomonas pohangensis	0.0025412960609911
methanomicrobium sp.	0.0025412960609911
catenulispore spp.	0.0025412960609911
ectothiorhodosinus spp.	0.0025412960609911
holospora spp.	0.0025412960609911
ulvibacter spp.	0.0025412960609911
campylobacter spp.	0.0025412960609911
spirochaeta aurantia	0.0025412960609911
aciditerrimonas spp.	0.0025412960609911
phycisphaera spp.	0.0025412960609911
enterococcus spp.	0.0025412960609911
phaeobacter gallaeiensis	0.0025412960609911
shewanella spp.	0.0025412960609911
desulfobacula spp.	0.0025412960609911
candidatus procabacter sp. oew1	0.0025412960609911
roseiflexus spp.	0.0025412960609911
geoalkalibacter spp.	0.0025412960609911
exiguobacterium undae	0.0025412960609911
proteiniphilum spp.	0.0025412960609911
candidatus phytoplasma solani	0.0025412960609911
desulfovofaba hansenii	0.0025412960609911
spirillum winogradskyi	0.0025412960609911
halothiobacillus sp.	0.0025412960609911
ohtaekwangia koreensis	0.0025412960609911
sulfurospirillum thermobaculum sp.	0.0025412960609911
lewinella nigricans	0.0025412960609911
thermoanaerobacter spp.	0.0025412960609911
sphingomonas wittichii	0.00169419737399407