

Table S4. Microbial community composition of the water samples. Relative abundance (%) of taxa within 16S rRNA amplicon libraries for phase 1 (“ph1”) and phase 2 (“ph2”). R1, R2: petro-F76 fuel. R3, R4: FT-F76 fuel. R5, R6: mix of petro- and FT-F76 fuel. R7, R8: no fuel added.

A. Phylum level taxa, with the exception of Alphaproteobacteria and Gammaproteobacteria (Class level).

Taxon	R1ph1	R1ph2	R2ph1	R2ph2	R3ph1	R3ph2	R4ph1	R4ph2	R5ph1	R5ph2	R6ph1	R6ph2	R7ph1	R7ph2	R8ph1	R8ph2
Acetothermia	0.1	0.0	0.1	0.0	0.0	0.1	0.1	0.0	0.0	0.0	0.0	0.0	1.1	0.4	0.0	0.3
Actinobacteriota	0.1	0.3	0.1	0.0	0.3	0.0	0.4	0.1	0.0	0.0	0.0	0.1	0.0	0.1	0.0	0.2
Actinobacteriota	4.5	0.8	2.4	1.3	0.3	0.5	1.8	0.6	1.9	0.4	0.4	0.4	0.9	0.5	0.7	0.3
Bacteroidota	3.1	0.7	9.4	4.2	10.4	14.6	1.4	0.7	1.7	0.6	0.7	2.6	8.7	10.5	7.0	2.6
Campylobacterota	2.6	5.6	28.4	3.5	1.0	0.9	1.3	0.7	4.0	0.8	29.9	2.4	0.6	1.0	0.7	0.5
Chloroflexi	0.4	3.9	0.3	0.1	1.4	0.5	1.8	2.0	0.3	0.6	1.2	1.1	0.8	0.2	2.5	0.5
Cyanobacteria	3.0	2.4	1.0	7.2	0.1	0.4	0.7	3.1	1.3	0.1	0.4	3.0	0.1	0.7	0.3	1.5
Deferribacterota	0.0	0.9	0.1	0.0	0.0	0.1	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Desulfobacterota	34.1	30.7	27.0	52.7	56.8	71.4	22.9	44.5	60.8	42.1	39.8	40.2	20.0	61.3	14.2	47.4
Elusimicrobiota	0.0	0.0	0.1	0.0	0.0	0.1	0.0	0.0	0.1	0.0	0.0	0.1	0.0	0.0	4.1	0.0
Firmicutes	37.0	30.3	8.9	2.7	4.7	4.0	21.1	2.8	16.8	6.9	5.6	32.9	5.8	3.8	11.4	5.8
*Alpha Proteobacteria	7.9	7.3	12.3	23.6	21.3	3.1	38.3	31.9	7.4	5.1	7.6	6.9	57.2	6.8	57.7	31.3
*Gamma Proteobacteria	3.5	15.6	7.5	3.9	2.6	1.3	7.0	12.6	3.0	41.9	12.9	7.2	2.9	7.2	3.7	7.8
Spirochaetota	0.0	0.2	0.1	0.0	0.2	1.6	0.0	0.0	0.1	0.2	0.4	1.1	0.7	0.4	0.4	0.3
Synergistota	0.4	0.1	0.3	0.1	0.1	0.0	0.0	0.1	0.2	0.0	0.0	0.0	0.3	0.1	0.0	0.0
Thermotogota	1.6	0.4	0.9	0.2	0.2	0.2	1.0	0.4	0.7	0.5	0.3	0.3	0.4	0.2	0.8	0.4

B. Genus level taxa.

Phylum	Family	Genus	R1ph1	R1ph2	R2ph1	R2ph2	R3ph1	R3ph2	R4ph1	R4ph2	R5ph1	R5ph2	R6ph1	R6ph2	R7ph1	R7ph2	R8ph1	R8ph2
Acetothermia	Acetothermiia	Acetothermiia	0.1	0.0	0.1	0.0	0.0	0.1	0.1	0.0	0.0	0.0	0.0	0.0	1.1	0.5	0.0	0.3
Actinobacteriota	Actinomarinales	Actinomarinales	0.1	0.0	0.0	0.0	0.3	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.2
Actinobacteriota	Dietziaceae	Dietzia	0.7	0.0	0.2	0.1	0.0	0.1	0.2	0.1	0.3	0.1	0.1	0.0	0.1	0.0	0.1	0.1
Actinobacteriota	Cellulomonadaceae	Actinotalea	0.0	0.1	0.3	0.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Actinobacteriota	Intrasporangiaceae	Ornithinococcus	0.5	0.0	0.4	0.2	0.1	0.1	0.4	0.1	0.2	0.0	0.1	0.1	0.2	0.1	0.2	0.0
Actinobacteriota	Micrococcaceae	Nesterenkonia	2.8	0.7	0.5	0.5	0.2	0.2	1.3	0.5	1.1	0.3	0.2	0.3	0.4	0.2	0.4	0.2
Bacteroidota	Bacteroidetes BD2-2	Bacteroidetes BD2-2	0.0	0.0	0.5	0.1	0.0	0.2	0.7	0.0	0.2	0.0	0.0	0.1	6.0	0.8	4.8	0.7
Bacteroidota	Prolixibacteraceae	Draconibacterium	0.2	0.0	0.1	0.4	0.0	0.1	0.2	0.2	0.3	0.0	0.2	1.5	0.0	0.2	0.0	0.0
Bacteroidota	Prolixibacteraceae	Prolixibacteraceae	0.1	0.3	0.1	0.0	0.4	1.1	0.1	0.0	0.0	0.2	0.3	0.7	0.3	0.7	0.3	0.4
Bacteroidota	Flavobacteriaceae	Lutibacter	0.7	0.0	8.3	3.6	9.9	0.1	0.1	0.3	0.2	0.1	0.0	0.0	0.6	0.3	0.4	0.9
Bacteroidota	Lentimicrobiaceae	Lentimicrobiaceae	1.5	0.1	0.3	0.1	0.2	13.1	0.1	0.1	0.4	0.2	0.1	0.1	1.5	8.8	0.8	0.6
Campylobacterota	Arcobacteraceae	Arcobacteraceae	0.5	0.5	5.0	0.7	0.3	0.3	0.3	0.0	0.1	0.0	6.1	0.4	0.1	0.4	0.2	0.2
Campylobacterota	Arcobacteraceae	Halarcobacter	0.2	1.5	2.4	1.0	0.0	0.0	0.1	0.0	1.1	0.0	4.2	0.5	0.1	0.2	0.2	0.0
Campylobacterota	Arcobacteraceae	Poseidonibacter	1.7	3.7	22.2	1.6	0.7	0.5	1.0	0.5	2.6	0.8	21.0	1.4	0.4	0.4	0.4	0.2
Campylobacterota	Sulfurimonadaceae	Sulfurimonas	0.0	0.0	0.2	0.2	0.0	0.0	0.0	0.0	0.2	0.0	0.1	0.1	0.0	0.0	0.0	0.0
Campylobacterota	Sulfurovaceae	Sulfurovum	0.3	0.2	0.2	0.0	0.1	0.0	0.0	0.1	0.2	0.0	0.1	0.1	0.0	0.0	0.0	0.1
Chloroflexi	Anaerolineaceae	ADurb.Bin120	0.4	4.0	0.2	0.2	0.1	0.5	1.9	1.9	0.3	0.6	0.0	0.1	0.2	0.0	0.9	0.3
Chloroflexi	Anaerolineaceae	Ornatilinea	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.9	0.9	0.3	0.1	0.6	0.1
Chloroflexi	Anaerolineaceae	Pelolinea	0.0	0.0	0.0	0.0	1.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.3	0.0	0.6	0.1
Chloroflexi	JG30-KF-CM45	JG30-KF-CM45	0.3	0.0	0.3	0.1	0.0	0.1	0.0	0.1	0.1	0.0	0.0	0.1	0.1	0.0	0.1	0.0
Cyanobacteria	Chloroplast	Chloroplast	3.2	2.4	1.0	7.3	0.1	0.4	0.6	3.1	1.3	0.2	0.4	3.1	0.1	0.7	0.3	1.6
Deferribacterota	Deferribacteraceae	Denitrovibrio	0.0	0.7	0.1	0.0	0.0	0.1	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Desulfobacterota	Desulfobacteraceae	Desulfobacter	7.4	2.6	2.6	0.7	30.7	0.8	4.7	29.0	17.4	1.1	1.6	1.5	4.6	0.7	0.9	0.7
Desulfobacterota	Desulfobacteraceae	Desulfobacteraceae	0.7	0.1	0.4	0.1	0.2	0.4	0.2	0.2	0.2	0.0	7.3	5.0	0.0	0.0	0.8	0.1
Desulfobacterota	Desulfobacteraceae	Desulfoconvexum	0.0	0.0	0.5	0.0	2.8	0.0	0.0	0.0	0.0	0.0	0.5	0.0	0.0	0.0	0.0	0.0
Desulfobacterota	Desulfobacteraceae	Desulfospira	0.0	0.0	0.0	0.0	0.9	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.7	0.0	2.4	0.2
Desulfobacterota	Desulfobacteraceae	Desulfotignum	0.0	14.7	0.1	0.0	0.0	0.4	0.0	0.0	0.0	2.7	0.0	0.0	0.0	0.2	0.2	0.2
Desulfobacterota	Desulfolunaceae	Desulfoluna	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.1	0.9	7.8	0.0	0.0	0.0	0.0
Desulfobacterota	Desulfosarcinaceae	Desulfatitalea	0.0	6.6	0.1	0.0	0.0	0.0	0.0	0.0	0.1	1.9	0.0	0.1	0.0	0.0	0.0	0.0
Desulfobacterota	Desulfosarcinaceae	Desulfonatronobacter	1.0	0.0	0.7	0.0	0.1	12.4	0.2	0.1	0.4	3.6	0.0	0.0	0.0	3.1	0.0	0.0
Desulfobacterota	Desulfosarcinaceae	Desulfosarcina	0.0	0.0	0.1	0.0	2.1	0.0	0.0	0.2	0.0	0.0	0.9	0.2	0.0	0.0	0.2	0.0
Desulfobacterota	Desulfosarcinaceae	Desulfosarcinaceae	1.1	0.2	1.5	33.2	0.3	0.3	0.4	0.3	0.5	0.1	0.6	3.8	0.2	0.5	0.4	0.4
Desulfobacterota	Desulfosarcinaceae	Sva0081 sediment group	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.6	0.5	0.0	0.0	0.0	0.0
Desulfobacterota	Desulfocapsaceae	Desulfocapsaceae	0.4	0.0	0.2	0.4	0.3	0.0	0.4	2.5	0.6	0.3	0.0	0.2	3.1	1.5	0.0	0.3
Desulfobacterota	Desulfocapsaceae	Desulfopila	0.4	1.1	0.7	0.2	6.7	0.4	1.7	0.4	1.6	0.6	0.3	0.2	0.0	0.2	0.1	0.2
Desulfobacterota	Desulfocapsaceae	Desulfotalea	1.0	0.5	1.0	0.6	0.2	0.3	1.2	0.3	30.0	0.5	3.8	0.6	0.5	0.3	0.7	0.4
Desulfobacterota	Desulfurivibrionaceae	MSBL7	0.2	0.2	0.2	0.0	0.0	0.2	0.0	0.0	0.1	0.0	0.0	1.5	0.0	1.3	0.0	0.0
Desulfobacterota	Desulfomicrobiaceae	Desulfomicrobium	1.0	0.2	0.5	0.2	0.3	0.3	0.5	0.1	0.7	0.2	0.2	0.3	0.1	0.3	0.2	0.1
Desulfobacterota	Desulfonatronaceae	Desulfonatronum	3.6	0.3	1.4	0.7	0.5	43.2	1.2	0.5	0.9	2.6	0.9	0.5	0.6	8.9	0.5	0.6
Desulfobacterota	Desulfonauticaceae	Desulfonauticus	2.2	0.4	0.7	0.4	0.2	0.2	0.7	0.6	0.9	0.3	0.3	0.2	0.3	0.4	0.3	0.2
Desulfobacterota	Desulfovibrionaceae	Desulfovibrio	14.5	3.0	12.6	14.9	13.4	13.4	12.4	10.9	8.7	28.0	12.6	17.4	7.7	45.0	8.0	45.6
Desulfobacterota	Desulfovibrionaceae	Halodesulfovibrio	0.0	0.2	1.5	0.3	0.0	0.0	0.0	0.0	0.0	0.1	0.2	0.0	0.0	0.0	0.0	0.0
Desulfobacterota	Sva1033	Sva1033	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Desulfobacterota	Syntrophotaleaceae	Syntrophotalea	2.0	1.3	3.0	1.6	0.3	0.2	0.6	0.4	1.3	0.3	9.8	1.1	1.9	0.7	0.0	0.2
Elusimicrobiota	Lineage IV	Lineage IV	0.0	0.0	0.1	0.0	0.0	0.1	0.0	0.0	0.1	0.0	0.0	0.1	0.0	4.3	0.0	0.0

Phylum	Family	Genus	R1ph1	R1ph2	R2ph1	R2ph2	R3ph1	R3ph2	R4ph1	R4ph2	R5ph1	R5ph2	R6ph1	R6ph2	R7ph1	R7ph2	R8ph1	R8ph2
Firmicutes	Acholeplasmataceae	Acholeplasma	0.0	4.7	0.0	0.0	0.2	0.0	0.6	0.2	0.3	0.5	0.8	0.2	1.1	0.1	0.2	0.0
Firmicutes	Bacillaceae	Bacillaceae	5.9	0.7	1.4	0.7	0.5	0.4	1.1	0.4	1.6	0.5	0.5	0.6	0.4	0.2	0.5	0.5
Firmicutes	Bacillaceae	Bacillus	3.6	0.9	0.7	0.3	0.2	0.2	0.5	0.3	1.6	0.2	0.1	0.4	0.3	0.3	0.3	0.0
Firmicutes	Caldalkalibacillaceae	Caldalkalibacillus	4.4	2.2	0.4	0.4	0.6	0.5	2.8	0.2	2.0	0.1	0.3	0.3	1.0	0.6	0.6	0.3
Firmicutes	Erysipelothricaceae	Erysipelothrix	0.2	2.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.3	0.0	0.0	0.0	0.0	0.0	0.0
Firmicutes	Izemoplasmataceae	Izemoplasmataceae	0.3	0.1	0.2	0.0	0.2	0.0	0.7	0.1	0.4	0.0	0.8	1.9	0.0	0.2	2.4	1.6
Firmicutes	Izemoplasmataceae	Izimaplasma	0.5	1.7	0.7	0.0	1.3	0.9	13.5	0.2	4.0	1.3	0.3	0.2	1.0	0.1	4.5	2.1
Firmicutes	Clostridia vadinBB60 group	Clostridia vadinBB60 group	0.0	0.1	0.0	0.0	0.2	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.1	0.0	1.1	0.1
Firmicutes	Clostridiaceae	Clostridium sensu stricto 18	2.2	0.3	0.8	0.1	0.2	0.2	0.4	0.2	1.2	0.3	0.4	0.1	0.3	0.2	0.4	0.2
Firmicutes	Garciellaceae	Garciella	4.8	0.9	2.2	0.5	0.8	0.9	0.9	0.8	2.5	0.7	1.0	0.8	0.7	0.6	0.7	0.8
Firmicutes	Hungateiclostridiaceae	Ercella	0.0	4.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Firmicutes	Dethiosulfatibacteraceae	Dethiosulfatibacter	0.1	9.2	0.1	0.0	0.0	0.0	0.0	0.0	0.1	1.9	0.0	0.0	0.0	0.1	0.0	0.0
Firmicutes	JTB215	JTB215	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.7	0.1	0.0	0.0	0.0	0.0
Firmicutes	Thermincolaceae	Thermincola	0.5	0.0	0.3	0.1	0.0	0.1	0.1	0.0	0.2	0.0	0.1	0.1	0.1	0.0	0.0	0.1
Firmicutes	Desulfotomaculales Incertae Sedis	Desulfallax-Sporotomaculum	2.3	0.2	0.3	0.1	0.1	0.2	0.4	0.1	0.8	0.3	0.1	0.3	0.1	0.1	0.3	0.1
Firmicutes	Firmicutes	Firmicutes	0.3	0.1	0.3	0.1	0.0	0.1	0.1	0.0	0.1	0.0	0.1	1.1	0.0	0.0	0.0	0.0
Firmicutes	Thermolithobacteraceae	Thermolithobacter	9.0	0.0	0.1	0.0	0.0	0.0	0.2	0.0	0.2	0.0	0.1	0.1	0.1	0.1	0.0	0.0
Firmicutes	Desulfitibacteraceae	Desulfitibacter	1.0	0.2	0.6	0.2	0.1	0.2	0.2	0.0	0.8	0.4	0.3	28.6	0.3	0.2	0.2	0.1
Firmicutes	Thermacetogeniaceae	Thermacetogeniaceae	2.4	0.0	0.3	0.0	0.0	0.1	0.1	0.1	0.7	0.0	0.1	0.1	0.1	0.1	0.3	0.0
Patescibacteria	Absconditabacteriales (SR1)	Absconditabacteriales (SR1)	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.3	0.0	0.0
Proteobacteria	Hyphomonadaceae	Hyphomonadaceae	0.0	0.0	0.1	0.0	0.0	0.0	3.8	1.0	0.3	0.0	0.0	0.0	0.0	0.0	0.2	0.0
Proteobacteria	Hyphomonadaceae	Hyphomonas	0.1	0.2	0.2	0.4	2.2	0.0	2.1	0.0	0.2	0.1	0.6	0.7	0.8	0.0	0.1	0.1
Proteobacteria	Parvibaculaceae	Mf105b01	0.0	2.8	0.0	0.0	0.1	0.0	0.0	1.1	0.0	0.0	0.0	0.0	0.0	0.1	0.1	0.3
Proteobacteria	Parvibaculaceae	Parvibaculum	0.3	0.1	4.2	16.5	0.2	0.0	0.2	0.0	0.2	0.1	0.0	0.1	0.3	0.3	0.1	0.1
Proteobacteria	Rhizobiaceae	Cohaesibacter	0.0	0.3	0.3	0.0	1.1	0.1	1.6	0.7	0.0	0.0	1.6	0.4	0.3	0.1	0.4	0.0
Proteobacteria	Rhodobacteraceae	Paracoccus	1.7	0.3	1.8	0.4	0.1	0.1	0.3	0.2	1.3	0.3	0.2	0.2	0.0	0.1	0.3	0.2
Proteobacteria	Rhodobacteraceae	Rhodobacteraceae	2.1	1.4	1.3	2.0	12.7	0.9	6.3	4.1	1.7	3.5	2.3	0.7	5.3	0.7	4.1	2.2
Proteobacteria	Rhodobacteraceae	Ruegeria	0.0	0.2	0.2	3.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.6	0.0	0.0	0.0
Proteobacteria	AEGEAN-169 marine group	AEGEAN-169 marine group	0.3	0.0	0.1	0.1	0.1	0.0	0.7	0.1	0.2	0.2	0.0	0.0	0.1	0.0	0.1	0.0
Proteobacteria	Magnetospiraceae	Magnetovibrio	3.0	0.9	2.0	0.9	2.9	1.7	22.0	23.2	2.0	0.9	1.3	0.6	46.1	4.6	49.6	25.2
Proteobacteria	Terasakiellaceae	Terasakiella	0.1	0.5	0.8	0.2	0.0	0.0	0.0	0.0	0.2	0.0	0.6	0.7	0.0	0.0	0.0	0.0
Proteobacteria	Thalassospiraceae	Thalassospira	0.1	0.6	0.5	0.1	0.2	0.0	0.2	0.0	0.4	0.1	0.3	0.2	0.1	0.0	0.0	0.0
Proteobacteria	Sphingomonadaceae	Erythrobacter	0.5	0.0	0.5	0.1	0.2	0.1	0.1	0.1	0.3	0.1	0.2	0.2	3.5	0.7	1.9	0.7
Proteobacteria	Thalassobaculales	Thalassobaculales	0.1	0.0	0.0	0.0	0.1	0.0	0.4	0.2	0.1	0.0	0.0	0.3	0.2	0.1	0.2	0.1
Proteobacteria	CH2b56	CH2b56	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.0	0.0	0.0	0.9
Proteobacteria	Sedimenticolaceae	Candidatus Thiodiazotropha	0.0	0.0	0.9	0.9	0.2	0.1	0.0	0.1	0.1	0.2	0.6	3.5	0.3	0.5	0.1	0.0
Proteobacteria	Thioalkalispiraceae	Thiohalophilus	0.1	0.0	0.1	0.0	0.1	0.0	0.5	5.8	0.1	0.2	0.0	0.1	0.0	0.0	0.0	0.1
Proteobacteria	Pseudoalteromonadaceae	Pseudoalteromonas	0.0	1.7	0.1	0.0	0.1	0.0	0.3	0.0	0.1	0.2	0.0	0.0	0.0	0.0	0.0	0.0
Proteobacteria	Shewanellaceae	Shewanella	0.0	1.1	0.2	0.0	0.0	0.0	0.5	0.0	0.2	0.0	0.3	0.2	0.0	0.0	0.2	0.0
Proteobacteria	Vibrionaceae	Vibrio	0.1	0.0	0.3	0.1	0.1	0.1	0.0	0.1	0.2	0.0	0.8	0.2	0.0	0.0	0.0	0.0
Proteobacteria	Unknown Family	Unknown Family	0.5	0.1	0.4	0.0	1.1	0.5	0.3	2.9	0.6	0.1	0.3	0.7	0.6	6.2	0.2	4.8
Proteobacteria	Immundisolibacteraceae	Immundisolibacter	0.0	4.4	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Proteobacteria	Halomonadaceae	Halomonas	0.3	0.2	0.4	0.0	0.1	0.0	4.6	0.0	0.7	0.0	0.1	0.0	0.1	0.0	2.5	0.2
Proteobacteria	Marinobacteraceae	Marinobacter	1.4	4.0	1.8	2.3	0.2	0.6	0.1	3.2	0.7	38.2	3.1	1.7	0.4	0.4	0.0	0.2
Proteobacteria	Moraxellaceae	Acinetobacter	0.0	0.0	0.1	0.0	0.0	0.1	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.2
Proteobacteria	Nitrincolaceae	Amphritea	0.2	0.0	0.5	0.0	0.2	0.0	0.5	0.4	0.2	0.1	2.3	0.3	0.3	0.1	0.2	0.0
Proteobacteria	Nitrincolaceae	Marinobacterium	0.4	0.8	0.6	0.2	0.0	0.0	0.0	0.0	0.0	0.0	1.5	0.1	0.0	0.0	0.0	0.0
Proteobacteria	Nitrincolaceae	Neptunomonas	0.0	0.2	0.3	0.1	0.0	0.0	0.0	0.0	0.2	0.0	2.9	0.1	0.0	0.0	0.0	0.0
Proteobacteria	Nitrincolaceae	Nitrincola	0.0	1.7	0.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.7	0.0	0.0	0.0	0.0	0.0
Proteobacteria	Woeseiaceae	Woeseia	0.0	0.3	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Proteobacteria	Thiotrichaceae	Thiotrichaceae	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
Spirochaetota	Spirochaetaceae	Alkalispirochaeta	0.0	0.0	0.0	0.0	0.0	1.5	0.0	0.0	0.0	0.2	0.0	0.0	0.7	0.4	0.0	0.0
Spirochaetota	Spirochaetaceae	Sediminispirochaeta	0.0	0.2	0.0	0.0	0.2	0.1	0.0	0.0	0.1	0.0	0.5	1.2	0.1	0.0	0.4	0.2
Spirochaetota	Spirochaetaceae	Spirochaeta 2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1
Thermotogota	Kosmotogaceae	Kosmotoga	0.8	0.4	0.5	0.1	0.2	0.1	0.3	0.4	0.4	0.2	0.3	0.2	0.2	0.2	0.2	0.1