

Supplementary Table S1. Class comparison of phylum Proteobacteria before and after air surging. The data are prevalence rates (%) in terms of the number of OTUs.

Proteobacteria	Before	After
Alphaproteobacteria	5.1	6.2
Betaproteobacteria	38.7	30.3
Deltaproteobacteria	0.4	7.8
Gammaproteobacteria	39.4	1.8
Other Proteobacteria	< 0.1	0.5
Total % of Proteobacteria	83.6	46.6

Supplementary Table S2. Comparison of major bacterial communities before and after air surging. The data are prevalence rate (%) in terms of number of OTUs.

OTUs number	Phylum	Class	Order	Family	Genus	Air surging work (%) Before After	
Phylum Acidobacteria-OTU01	Acidobacteria	11-24	uncultured_Acidobacteria_bacterium	-	-	0.0	0.1
Phylum Acidobacteria-OTU03	Acidobacteria	32-21	uncultured_bacterium	-	-	0.0	0.2
Phylum Acidobacteria-OTU06	Acidobacteria	Acidobacteriales	Acidobacteriaceae	uncultured	uncultured_bacterium	0.0	0.2
Phylum Acidobacteria-OTU13	Acidobacteria	DA023	uncultured_bacterium	-	-	0.1	2.0
Phylum Acidobacteria-OTU14	Acidobacteria	DA052	uncultured_bacterium	-	-	0.0	0.5
Phylum Acidobacteria-OTU19	Acidobacteria	Holophagae	Holophagales	Holophagaceae	Geothrix	0.1	0.5
Phylum Acidobacteria-OTU20	Acidobacteria	Holophagae	Holophagales	Holophagaceae	uncultured	0.2	0.3
Phylum Acidobacteria-OTU21	Acidobacteria	Holophagae	SJA-36	uncultured_bacterium	-	0.0	0.1
Phylum Acidobacteria-OTU31	Acidobacteria	SJA-149	-	-	-	0.0	0.1
Phylum Actinobacteria-OTU02	Actinobacteria	Acidimicrobia	Acidimicrobiales	Acidimicrobiaceae	CL500-29_marine_group	0.0	1.8

Phylum Actinobacteria-OTU09	Actinobacteria	Corynebacteriales	Nocardiaceae	Rhodococcus	-	0.5	0.1
Phylum Actinobacteria-OTU16	Actinobacteria	Frankiales	Sporichthyaceae	Candidatus_Planktophila	-	2.4	11.2
Phylum Actinobacteria-OTU25	Actinobacteria	Micrococcales	Microbacteriaceae	-	-	0.0	0.1
Phylum Actinobacteria-OTU27	Actinobacteria	Micrococcales	Micrococcaceae	Arthrobacter	-	0.1	0.1
Phylum Actinobacteria-OTU28	Actinobacteria	Micrococcales	Micrococcaceae	Arthrobacter	uncultured_bacterium	0.0	0.2
Phylum Actinobacteria-OTU29	Actinobacteria	Micrococcales	Micrococcaceae	Kocuria	-	0.2	0.0
Phylum Actinobacteria-OTU31	Actinobacteria	OPB41	uncultured_bacterium	-	-	0.0	0.1
Phylum Actinobacteria-OTU32	Actinobacteria	PeM15	uncultured_bacterium	-	-	0.0	0.1
Phylum Actinobacteria-OTU42	Actinobacteria	Thermoleophil	Gaiellales	Gaiellaceae	Gaiella	0.0	0.1
Phylum Bacteroidetes-OTU04	Bacteroidetes	Cytophagia	Cytophagales	Cytophagaceae	Arcicella	0.6	0.1
Phylum Bacteroidetes-OTU07	Bacteroidetes	Cytophagia	Cytophagales	Cytophagaceae	Flexibacter	0.1	0.4
Phylum Bacteroidetes-OTU08	Bacteroidetes	Cytophagia	Cytophagales	Cytophagaceae	Hymenobacter	0.3	0.0
Phylum Bacteroidetes-OTU17	Bacteroidetes	Flavobacteria	Flavobacteriales	Flavobacteriaceae	Flavobacterium	2.2	0.4
Phylum Bacteroidetes-OTU18	Bacteroidetes	Sphingobacteri	Sphingobacteriales	Chitinophagaceae	-	0.0	1.0
Phylum Bacteroidetes-OTU22	Bacteroidetes	Sphingobacteri	Sphingobacteriales	Chitinophagaceae	Hydrotalea	0.4	0.0
Phylum Bacteroidetes-OTU23	Bacteroidetes	Sphingobacteri	Sphingobacteriales	Chitinophagaceae	Lacibacter	0.0	0.1
Phylum Bacteroidetes-OTU24	Bacteroidetes	Sphingobacteri	Sphingobacteriales	Chitinophagaceae	Sediminibacterium	2.0	1.5
Phylum Bacteroidetes-OTU26	Bacteroidetes	Sphingobacteri	Sphingobacteriales	Chitinophagaceae	uncultured	0.0	0.1
Phylum Bacteroidetes-OTU29	Bacteroidetes	Sphingobacteri	Sphingobacteriales	KD3-93	uncultured_bacterium	0.0	0.1

Phylum Bacteroidetes-OTU31	Bacteroidetes	Sphingobacteriia	Sphingobacteriales	NS11-12_marine_group	uncultured_bacterium	0.7	2.4
Phylum Bacteroidetes-OTU32	Bacteroidetes	Sphingobacteriia	Sphingobacteriales	PHOS-HE51	uncultured_bacterium	0.0	0.2
Phylum Bacteroidetes-OTU35	Bacteroidetes	Sphingobacteriia	Sphingobacteriales	Sphingobacteriaceae	Mucilaginibacter	0.1	0.0
Phylum Bacteroidetes-OTU39	Bacteroidetes	Sphingobacteriia	Sphingobacteriales	Sphingobacteriaceae	Sphingobacterium	0.0	0.1
Phylum Bacteroidetes-OTU41	Bacteroidetes	Sphingobacteriia	Sphingobacteriales	env.OPS_17	-	0.0	0.1
Phylum Bacteroidetes-OTU44	Bacteroidetes	WCHB1-32	uncultured_bacterium	-	-	0.0	5.9
Phylum Chlorobi-OTU01	Chlorobi	Chlorobia	Chlorobiales	OPB56	uncultured_bacterium	0.0	0.2
Phylum Chlorobi-OTU02	Chlorobi	Ignavibacteria	Ignavibacteriales	BSV26	uncultured_bacterium	0.0	0.1
Phylum Chlorobi-OTU03	Chlorobi	Ignavibacteria	Ignavibacteriales	SR-FBR-L83	uncultured_bacterium	0.0	0.6
Phylum Chloroflexi-OTU09	Chloroflexi	KD4-96	uncultured_bacterium	-	-	0.0	2.2
Phylum Chloroflexi-OTU12	Chloroflexi	P2-11E	uncultured_bacterium	-	-	0.0	0.2
Phylum Chloroflexi-OTU13	Chloroflexi	S085	uncultured_bacterium	-	-	0.0	0.1

Supplementary Table S2. Continued

OTUs number	Phylum	Class	Order	Family	Genus	Air surging work (%)	Before	After
Phylum Chloroflexi-OTU14	Chloroflexi	SAR202_clade	uncultured_bacterium	-	-	0.0	0.1	
Phylum Chloroflexi-OTU15	Chloroflexi	TK10	-	-	-	0.0	0.2	

Phylum Cyanobacteria-OTU04	Cyanobacteria	MLE1-12	uncultured_bacterium	-	-	0.8	2.4
Phylum Cyanobacteria-OTU05	Cyanobacteria	SHA-109	uncultured_bacterium	-	-	0.0	0.1
Phylum Elusimicrobia-OTU04	Elusimicrobia	Lineage_IV	uncultured_Termite_group_1_bacterium	-	-	0.0	0.1
Phylum Elusimicrobia-OTU05	Elusimicrobia	Lineage_IV	uncultured_bacterium	-	-	0.3	1.5
Phylum Firmicutes-OTU01	Firmicutes	Bacilli	Bacillales	Bacillaceae	Anoxybacillus	0.0	0.1
Phylum Firmicutes-OTU06	Firmicutes	Bacilli	Bacillales	Staphylococcaceae	Staphylococcus	0.1	0.0
Phylum Firmicutes-OTU10	Firmicutes	Clostridia	Clostridiales	Family_XIII_Incertae_Sedis	uncultured	0.0	0.3
Phylum Firmicutes-OTU11	Firmicutes	Clostridia	Clostridiales	Gracilibacteraceae	Lutispora	0.0	0.4
Phylum Firmicutes-OTU12	Firmicutes	Clostridia	Clostridiales	Gracilibacteraceae	uncultured	0.0	0.1
Phylum Firmicutes-OTU14	Firmicutes	Clostridia	Clostridiales	Peptococcaceae	Desulfosporosinus	0.0	0.9
Phylum Firmicutes-OTU15	Firmicutes	Clostridia	Clostridiales	Ruminococcaceae	Incertae_Sedis	0.0	0.1
Phylum Gemmatimonadetes-OTU03	Gemmatimonadetes	Gemmatimonadales	Gemmatimonadaceae	uncultured	-	0.0	0.1
Phylum Gemmatimonadetes-OTU05	Gemmatimonadetes	Gemmatimonadales	Gemmatimonadaceae	uncultured	uncultured_bacterium	0.2	1.5
Phylum Nitrospirae-OTU01	Nitrospirae	Nitrospira	Nitrospirales	-	-	0.0	0.6
Phylum Nitrospirae-OTU03	Nitrospirae	Nitrospira	Nitrospirales	0319-6A21	uncultured_bacterium	0.0	0.4
Phylum Nitrospirae-OTU06	Nitrospirae	Nitrospira	Nitrospirales	Nitrospiraceae	Nitrospira	0.2	0.7
Phylum Planctomycetes-OTU04	Planctomycetes	OM190	uncultured_bacterium	-	-	0.0	0.1

Phylum Planctomycetes-OTU07	Planctomycetes	Phycisphaerae	Phycisphaerales	Phycisphaeraeae	SM1A02	0.0 0.2
Phylum Planctomycetes-OTU17	Planctomycetes	Planctomycetacia	Planctomycetales	Planctomycetaceae	Planctomyces	0.0 0.1
Phylum Planctomycetes-OTU20	Planctomycetes	Planctomycetacia	Planctomycetales	Planctomycetaceae	uncultured	0.0 0.1
Phylum Planctomycetes-OTU21	Planctomycetes	vadinHA49	uncultured_bacterium	-	-	0.0 0.1
Class Alphaproteobacteria-OTU03	Proteobacteria	Alphaproteobacteria	Caulobacterales	Caulobacteraceae	Brevundimonas	0.1 0.1
Class Alphaproteobacteria-OTU04	Proteobacteria	Alphaproteobacteria	Caulobacterales	Caulobacteraceae	Caulobacter	0.7 0.2
Class Alphaproteobacteria-OTU07	Proteobacteria	Alphaproteobacteria	Caulobacterales	Hyphomonadaceae	Woodsholea	0.0 0.1
Class Alphaproteobacteria-OTU08	Proteobacteria	Alphaproteobacteria	DB1-14	-	-	0.0 0.4
Class Alphaproteobacteria-OTU12	Proteobacteria	Alphaproteobacteria	Rhizobiales	A0839	uncultured_bacterium	0.0 0.1
Class Alphaproteobacteria-OTU14	Proteobacteria	Alphaproteobacteria	Rhizobiales	Bradyrhizobiaceae	Bradyrhizobium	0.0 0.1
Class Alphaproteobacteria-OTU23	Proteobacteria	Alphaproteobacteria	Rhizobiales	Methylobacteriaceae	Methylobacterium	0.1 0.1
Class Alphaproteobacteria-OTU33	Proteobacteria	Alphaproteobacteria	Rhodobacterales	Rhodobacteraceae	Rhodovulum	1.5 0.2
Class Alphaproteobacteria-OTU47	Proteobacteria	Alphaproteobacteria	Rhodospirillales	KCM-B-60	uncultured_Acetobacteraceae_bacterium	0.0 0.1
Class Alphaproteobacteria-OTU48	Proteobacteria	Alphaproteobacteria	Rhodospirillales	KCM-B-60	uncultured_Rhodospirillales_bacterium	0.0 0.1

Class							
Alphaproteobacteria-OTU51	Proteobacteria	Alphaproteobacteria	Rhodospirillales	Rhodospirillaceae	Magnetospirillum	0.0	0.2
Class							
Alphaproteobacteria-OTU53	Proteobacteria	Alphaproteobacteria	Rhodospirillales	Rhodospirillaceae	uncultured	0.0	0.1
Class							
Alphaproteobacteria-OTU54	Proteobacteria	Alphaproteobacteria	Rhodospirillales	wr0007	-	0.1	0.1
Class							
Alphaproteobacteria-OTU55	Proteobacteria	Alphaproteobacteria	Rhodospirillales	wr0007	uncultured_alpha_proteobacterium	0.0	0.1
Class							
Alphaproteobacteria-OTU67	Proteobacteria	Alphaproteobacteria	Rickettsiales	Rickettsiaceae	uncultured	0.0	0.5
Class							
Alphaproteobacteria-OTU73	Proteobacteria	Alphaproteobacteria	Sphingomonadales	Sphingomonadaceae	Novosphingobium	1.7	3.2
Class							
Alphaproteobacteria-OTU74	Proteobacteria	Alphaproteobacteria	Sphingomonadales	Sphingomonadaceae	Sphingobium	0.1	0.1
Class							
Alphaproteobacteria-OTU75	Proteobacteria	Alphaproteobacteria	Sphingomonadales	Sphingomonadaceae	Sphingomonas	0.3	0.2

Supplementary Table S2. Continued

OTUs number	Phylum	Class	Order	Family	Genus	Air surging work (%)	
						Before	After
Class							
Alphaproteobacteria-OTU76	Proteobacteria	Alphaproteobacteria	Sphingomonadales	Sphingomonadaceae	Sphingopyxis	0.3	0.0
Class							
Betaproteobacteria-OTU01	Proteobacteria	Betaproteobacteria	-	-	-	0.1	0.3
Class							
Betaproteobacteria-OTU02	Proteobacteria	Betaproteobacteria	Burkholderiales	Alcaligenaceae	Achromobacter	0.2	0.0

Class Betaproteobacteria-OTU04	Proteobacteria	Betaproteobacteria	Burkholderiales	Burkholderiaceae	Cupriavidus	0.2	0.0
Class Betaproteobacteria-OTU05	Proteobacteria	Betaproteobacteria	Burkholderiales	Comamonadaceae	-	3.9	6.4
Class Betaproteobacteria-OTU06	Proteobacteria	Betaproteobacteria	Burkholderiales	Comamonadaceae	Aquabacterium	0.1	0.1
Class Betaproteobacteria-OTU07	Proteobacteria	Betaproteobacteria	Burkholderiales	Comamonadaceae	Comamonas	0.2	0.0
Class Betaproteobacteria-OTU09	Proteobacteria	Betaproteobacteria	Burkholderiales	Comamonadaceae	Paucibacter	0.5	0.0
Class Betaproteobacteria-OTU11	Proteobacteria	Betaproteobacteria	Burkholderiales	Comamonadaceae	Polaromonas	0.4	0.1
Class Betaproteobacteria-OTU14	Proteobacteria	Betaproteobacteria	Burkholderiales	Comamonadaceae	Variovorax	0.3	0.0
Class Betaproteobacteria-OTU15	Proteobacteria	Betaproteobacteria	Burkholderiales	Comamonadaceae	uncultured	0.3	0.2
Class Betaproteobacteria-OTU16	Proteobacteria	Betaproteobacteria	Burkholderiales	Oxalobacteraceae	-	0.4	0.3
Class Betaproteobacteria-OTU18	Proteobacteria	Betaproteobacteria	Burkholderiales	Oxalobacteraceae	Massilia	0.2	0.0
Class Betaproteobacteria-OTU19	Proteobacteria	Betaproteobacteria	Burkholderiales	Oxalobacteraceae	Paucimonas	0.0	0.1
Class Betaproteobacteria-OTU20	Proteobacteria	Betaproteobacteria	Burkholderiales	Oxalobacteraceae	Undibacterium	0.3	0.3
Class Betaproteobacteria-OTU22	Proteobacteria	Betaproteobacteria	Hydrogenophiles	Hydrogenophilaceae	Sulfuricella	0.4	0.6
Class Betaproteobacteria-OTU25	Proteobacteria	Betaproteobacteria	Nitrosomonadales	Gallionellaceae	-	10.2	5.2
Class Betaproteobacteria-OTU26	Proteobacteria	Betaproteobacteria	Nitrosomonadales	Gallionellaceae	Candidatus_Nitrospirogaster	0.1	0.5
Class Betaproteobacteria-OTU27	Proteobacteria	Betaproteobacteria	Nitrosomonadales	Gallionellaceae	Gallionella	17.4	1.7
Class Betaproteobacteria-OTU28	Proteobacteria	Betaproteobacteria	Nitrosomonadales	Gallionellaceae	Sideroxydans	0.2	1.0

Class Betaproteobacteria-OTU29	Proteobacteria	Betaproteobacteria	Nitrosomonadales	Gallionellaceae	uncultured	2.1	3.9
Class Betaproteobacteria-OTU30	Proteobacteria	Betaproteobacteria	Nitrosomonadales	Nitrosomonadaceae	-	0.0	0.4
Class Betaproteobacteria-OTU32	Proteobacteria	Betaproteobacteria	Nitrosomonadales	Nitrosomonadaceae	uncultured	0.0	0.8
Class Betaproteobacteria-OTU33	Proteobacteria	Betaproteobacteria	Rhodocyclales	Rhodocyclaceae	-	0.5	3.4
Class Betaproteobacteria-OTU34	Proteobacteria	Betaproteobacteria	Rhodocyclales	Rhodocyclaceae	Azospira	0.1	0.0
Class Betaproteobacteria-OTU35	Proteobacteria	Betaproteobacteria	Rhodocyclales	Rhodocyclaceae	Dechloromonas	0.3	1.3
Class Betaproteobacteria-OTU38	Proteobacteria	Betaproteobacteria	Rhodocyclales	Rhodocyclaceae	Propionivibrio	0.0	0.2
Class Betaproteobacteria-OTU39	Proteobacteria	Betaproteobacteria	Rhodocyclales	Rhodocyclaceae	Sterolibacterium	0.0	0.1
Class Betaproteobacteria-OTU40	Proteobacteria	Betaproteobacteria	Rhodocyclales	Rhodocyclaceae	Sulfuritalea	0.1	2.4
Class Betaproteobacteria-OTU41	Proteobacteria	Betaproteobacteria	Rhodocyclales	Rhodocyclaceae	Zoogloea	0.1	0.1
Class Betaproteobacteria-OTU42	Proteobacteria	Betaproteobacteria	Rhodocyclales	Rhodocyclaceae	uncultured	0.0	0.8
Class Deltaproteobacteria-OTU08	Proteobacteria	Deltaproteobacteria	Desulfovibrionales	Desulfovibrionaceae	Desulfovibrio	0.0	0.3
Class Deltaproteobacteria-OTU10	Proteobacteria	Deltaproteobacteria	Desulfuromonadales	-	-	0.0	0.6
Class Deltaproteobacteria-OTU11	Proteobacteria	Deltaproteobacteria	Desulfuromonadales	BVA18	uncultured_Geobacter_sp.	0.0	1.9
Class Deltaproteobacteria-OTU12	Proteobacteria	Deltaproteobacteria	Desulfuromonadales	BVA59	uncultured_bacterium	0.0	0.5
Class Deltaproteobacteria-OTU13	Proteobacteria	Deltaproteobacteria	Desulfuromonadales	Desulfuromonadaceae	-	0.0	0.1
Class Deltaproteobacteria-OTU14	Proteobacteria	Deltaproteobacteria	Desulfuromonadales	Desulfuromonadaceae	Pelobacter	0.0	0.2

Class Deltaproteobacteria-OTU15	Proteobacteria	Deltaproteobacteria	Desulfuromonadales	Geobacteraceae	Geobacter	0.1	3.5
Class Deltaproteobacteria-OTU17	Proteobacteria	Deltaproteobacteria	GR-WP33-30	uncultured_bacterium	-	0.0	0.2
Class Deltaproteobacteria-OTU20	Proteobacteria	Deltaproteobacteria	Myxococcales	0319-6G20	uncultured_bacterium	0.1	0.1
Class Deltaproteobacteria-OTU23	Proteobacteria	Deltaproteobacteria	Myxococcales	Cystobacterineae	uncultured	0.0	0.1

Supplementary Table S2. Continued

OTUs number	Phylum	Class	Order	Family	Genus	Air surging work (%)	
						Bef	Aft
Class Gammaproteobacteria-OTU05	Proteobacteria	Gammaproteobacteria	Chromatiales	Ectothiorhodospiraceae	Acidiferrobacter	0.0	0.2
Class Gammaproteobacteria-OTU06	Proteobacteria	Gammaproteobacteria	Enterobacteriales	Enterobacteriaceae	Enterobacter	2.6	0.0
Class Gammaproteobacteria-OTU07	Proteobacteria	Gammaproteobacteria	Enterobacteriales	Enterobacteriaceae	Escherichia-Shigella	0.1	0.0
Class Gammaproteobacteria-OTU08	Proteobacteria	Gammaproteobacteria	Enterobacteriales	Enterobacteriaceae	Hafnia-Obesumbacterium	0.9	0.0
Class Gammaproteobacteria-OTU09	Proteobacteria	Gammaproteobacteria	Enterobacteriales	Enterobacteriaceae	Serratia	0.1	0.0
Class Gammaproteobacteria-OTU14	Proteobacteria	Gammaproteobacteria	Legionellales	Legionellaceae	Legionella	0.4	0.6
Class Gammaproteobacteria-OTU18	Proteobacteria	Gammaproteobacteria	Oceanospirillales	Oceanospirillaceae	Pseudospirillum	0.0	0.2
Class Gammaproteobacteria-OTU19	Proteobacteria	Gammaproteobacteria	PYR10d3	uncultured_sulfur-oxidizing_symbiont_bacterium	-	0.0	0.1

Class							
Gammaproteobacteria-OTU22	Proteobacteria	Gammaproteobacteria	Pseudomonadales	Moraxellaceae	Acinetobacter	15.8	0.2
Class							
Gammaproteobacteria-OTU23	Proteobacteria	Gammaproteobacteria	Pseudomonadales	Moraxellaceae	Alkanindiges	0.5	0.0
Class							
Gammaproteobacteria-OTU24	Proteobacteria	Gammaproteobacteria	Pseudomonadales	Moraxellaceae	Enhydrobacter	0.2	0.0
Class							
Gammaproteobacteria-OTU27	Proteobacteria	Gammaproteobacteria	Pseudomonadales	Pseudomonadaceae	Pseudomonas	11.7	0.0
Class							
Gammaproteobacteria-OTU30	Proteobacteria	Gammaproteobacteria	Xanthomonadales	Sinobacteraceae	Hydrocarboniphaga	1.1	0.0
Class							
Gammaproteobacteria-OTU34	Proteobacteria	Gammaproteobacteria	Xanthomonadales	Sinobacteraceae	uncultured	0.0	0.3
Class							
Gammaproteobacteria-OTU38	Proteobacteria	Gammaproteobacteria	Xanthomonadales	Xanthomonadaceae	Stenotrophomonas	5.7	0.0
Phylum							
Proteobacteria-OTU03	Proteobacteria	SPOTSOCT00m83	uncultured_bacterium	-	-	0.0	0.4
Phylum							
Proteobacteria-OTU04	Proteobacteria	TA18	-	-	-	0.0	0.1
Phylum							
Verrucomicrobia-OTU01	Verrucomicrobia	Candidatus_Methylocandidiphilum	uncultured_bacterium	-	-	0.0	0.1
Phylum							
Verrucomicrobia-OTU04	Verrucomicrobia	OPB35_soil_group	uncultured_bacterium	-	-	2.1	0.8
Phylum							
Verrucomicrobia-OTU05	Verrucomicrobia	Opitutae	Opitales	Opitutaceae	Opitutus	0.1	1.8
Phylum							
Verrucomicrobia-OTU06	Verrucomicrobia	Verrucomicrobiae	Verrucomicrobiales	Verrucomicrobiaceae	Prostheco bacter	0.1	0.0
Unknown	-	-	-	-	-	1.5	3.6
Ungrouped	BHI80-139	uncultured_bacterium	-	-	-	0.0	0.1
Ungrouped	Candidate_division_OD1	-	-	-	-	0.0	0.1
Ungrouped	Candidate_division_OD1	uncultured_bacterium	-	-	-	0.3	1.3

Ungrouped	Candidate_division_OP3	uncultured_bacterium	-	-	-	0.0	0.1
Ungrouped	Candidate_division_WS3	uncultured_bacterium	-	-	-	0.0	0.2
Ungrouped	SM2F11	uncultured_bacterium	-	-	-	0.1	0.1
Ungrouped	TA06	uncultured_bacterium	-	-	-	0.0	0.5
Ungrouped	TM6	uncultured_bacterium	-	-	-	0.1	0.2
