

FREE-LIVING										
ASV #	index	Getz	West OP	OP	Dotson	East CC	p.value.bh	Order	Family	Genus
Seq260	1	0	0	0	1	0	0.014	SAR202 clade	uncultured Chloroflexi bacterium HF4000_28F02	uncultured Chloroflexi bacterium HF4000_28F02 Family
Seq134	2	0	0	0	0	1	0.031	Thiomicrospirales	Thioglobaceae	Thioglobaceae Family
Seq1405	3	1	0	0	0	0	0.045	Anaerolineales	Anaerolineaceae	uncultured 24
Seq1671	3	1	0	0	0	0	0.045	Coxiellales	Coxiellaceae	Coxiella
Seq210	3	1	0	0	0	0	0.037	Burkholderiales	Nitrosomonadaceae	Nitrosomonas
Seq228	3	1	0	0	0	0	0.046	Cellvibrionales	Halieaceae	Halieaceae Family
Seq326	3	1	0	0	0	0	0.045	Pseudomonadales	Pseudomonadaceae	Pseudomonadaceae Family
Seq447	3	1	0	0	0	0	0.024	Opitutales	Puniceicoccaceae	Pelagicoccus
Seq661	3	1	0	0	0	0	0.024	Actinomarinales	Actinomarinaceae	Candidatus Actinomarina
Seq17	5	0	1	0	0	0	0.014	Flavobacteriales	Flavobacteriaceae	uncultured 2
Seq171	5	0	1	0	0	0	0.031	Lentisphaerales	Lentisphaeraceae	Lentisphaera
Seq188	5	0	1	0	0	0	0.045	Oceanospirillales	Nitrincolaceae	Profundimonas
Seq215	5	0	1	0	0	0	0.033	Alteromonadales	Pseudoalteromonadaceae	Pseudoalteromonadaceae Family
Seq3	5	0	1	0	0	0	0.029	Oceanospirillales	Nitrincolaceae	uncultured 1
Seq359	5	0	1	0	0	0	0.037	Fibrobacterales	Fibrobacteraceae	uncultured 13
Seq43	5	0	1	0	0	0	0.031	Alteromonadales	Alteromonadaceae	Paraglaciecola
Seq5	5	0	1	0	0	0	0.024	Flavobacteriales	Flavobacteriaceae	Polaribacter
Seq6	5	0	1	0	0	0	0.033	Cellvibrionales	Porticoccaceae	SAR92 clade
Seq775	5	0	1	0	0	0	0.045	Campylobacterales	Arcobacteraceae	uncultured 18
Seq81	5	0	1	0	0	0	0.045	Chitinophagales	Saprospiraceae	Saprospiraceae Family
Seq834	5	0	1	0	0	0	0.045	Campylobacterales	Arcobacteraceae	Halarcobacter
Seq117	6	0	0	0	1	1	0.029	Sphingobacteriales	Lentimicrobiaceae	uncultured deep-sea archaeon
Seq155	6	0	0	0	1	1	0.014	Planctomycetales	uncultured	uncultured bacterium HF0500_40D21
Seq195	6	0	0	0	1	1	0.029	Rhodospirillales	Magnetospiraceae	uncultured 7
Seq58	6	0	0	0	1	1	0.011	Verrucomicrobiales	Rubritaleaceae	Roseibacillus
Seq563	7	1	0	0	1	0	0.013	SAR202 clade	uncultured bacterium HF0500_03M05	uncultured bacterium HF0500_03M05 Family
Seq157	10	1	0	0	0	1	0.014	SAR11 clade	Clade II	uncultured marine bacterium 440
Seq161	10	1	0	0	0	1	0.011	Nitrospinales	Nitrospinaceae	LS-NOB
Seq265	10	1	0	0	0	1	0.024	Rhodospirillales	AEGEAN-169 marine group	unidentified marine bacterioplankton 2
Seq110	16	1	0	0	1	1	0.031	Flavobacteriales	Flavobacteriaceae	NS4 marine group
Seq132	16	1	0	0	1	1	0.011	Steroidobacterales	Woeseiaceae	Woeseia
Seq199	16	1	0	0	1	1	0.013	SAR202 clade	hydrothermal vent metagenome	hydrothermal vent metagenome Family 3
Seq220	16	1	0	0	1	1	0.011	Phycisphaerales	Phycisphaeraceae	Phycisphaeraceae Family
Seq38	16	1	0	0	1	1	0.013	SAR11 clade	Clade I	Clade I Family
Seq95	16	1	0	0	1	1	0.031	Pirellulales	Pirellulaceae	Rhodopirellula
Seq97	16	1	0	0	1	1	0.024	Burkholderiales	Methylophilaceae	OM43 clade

\* no sig indicator taxa for particle-associated

\* no western CC has pure CDW