

PARTICLE-ASSOCIATED

Upper 200m

No significant indicator taxa.

Lower 200m

ASV #	index	OP	West CC	Dotson	Eastern CC	Getz	p.value.bh	Order	Family	Genus
Seq235	2	0	0	0	1	0	0.047	Planctomycetales	Gimesiaceae	uncultured 8
Seq801	2	0	0	0	1	0	0.042	Victivallales	Arctic95B-14	uncultured Verrucomicrobia bacterium
Seq1526	3	0	0	0	0	1	0.027	Methylococcales	Methylomonadaceae	Marine Methylotrophic Group 2
Seq156	3	0	0	0	0	1	0.047	Defluviicoccales	uncultured	uncultured bacterium KM3-47-A6
Seq272	3	0	0	0	0	1	0.033	Phycisphaerales	Phycisphaeraceae	CL500-3
Seq380	3	0	0	0	0	1	0.009	Pirellulales	Pirellulaceae	Pirellulaceae Family
Seq407	3	0	0	0	0	1	0.009	Nannocystales	Nannocystaceae	Nannocystaceae Family
Seq452	3	0	0	0	0	1	0.022	Planctomycetales	Rubinisphaeraceae	uncultured 15
Seq482	3	0	0	0	0	1	0.009	Bradymonadales	uncultured marine bacterium	uncultured marine bacterium Family 2
Seq653	3	0	0	0	0	1	0.027	Oligoflexales	uncultured	uncultured sediment bacterium
Seq171	5	0	1	0	0	0	0.047	Lentisphaerales	Lentisphaeraceae	Lentisphaera
Seq3	5	0	1	0	0	0	0.027	Oceanospirillales	Nitrincolaceae	uncultured 1
Seq420	7	0	0	1	0	1	0.042	Pirellulales	Pirellulaceae	Blastopirellula
Seq95	10	0	0	0	1	1	0.028	Pirellulales	Pirellulaceae	Rhodopirellula
Seq6	15	1	1	0	0	0	0.047	Cellvibrionales	Porticoccaceae	SAR92 clade
Seq1	16	0	0	1	1	1	0.009	Pseudomonadales	Pseudomonadaceae	Pseudomonas
Seq155	16	0	0	1	1	1	0.042	Planctomycetales	uncultured	uncultured bacterium HF0500_40D21
Seq308	16	0	0	1	1	1	0.047	Pirellulales	Pirellulaceae	uncultured 11