

ASV #	index	Size Fract.*	Flow **	p	Relative Abundance	Order	Family	Genus
Seq58	2	FL	Outflow	0.027	5.6E-04	Verrucomicrobiales	Rubritaleaceae	Roseibacillus
Seq71	2	FL	Outflow	0.015	2.8E-04	Flavobacteriales	Cryomorphaceae	Vicingus
Seq78	2	FL	Outflow	0.009	4.2E-06	Cellvibrionales	Cellvibrionaceae	uncultured 5
Seq91	2	FL	Outflow	0.005	1.0E-05	Alteromonadales	Colwelliaceae	Colwellia
Seq93	2	FL	Outflow	0.009	4.9E-03	SAR86 clade	uncultured Oceanospirillales bacterium	uncultured Oceanospirillales bacterium Family
Seq94	2	FL	Outflow	0.023	1.1E-03	Phycisphaerales	Phycisphaeraceae	JL-ETNP-F27
Seq97	2	FL	Outflow	0.044	5.8E-03	Burkholderiales	Methylophilaceae	OM43 clade

\* No sig. taxa in outflow for particle-associated.

\*\* Sig. taxa in only outflow.