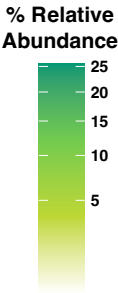


Most Abundant Taxa

Campylobacterota; Sulfuricurvum-Chloroflexi; SCGC-AB-539-J10	11	25.6	0.1
Campylobacterota; Sulfurimonas-Actinobacteriota; Mycobacterium-Halobacterota; Methanolinea-Chloroflexi; Pelolinea	0.1	1.3	11.2
Acidobacteriota; Aminicenantales ASV499	2.6	14.4	
Acidobacteriota; Aminicenantales ASV485	3.5	2.9	2.8
Halobacterota; Methanosaeta-Chloroflexi; GIF9 ASV5911	0.7	1.2	2.3
Acidobacteriota; Aminicenantales ASV410		0.5	1.3
Actinobacteriota; Actinomarinales ASV28	0.4	0.5	0.7
Chloroflexi; MSBL5 ASV6114	2.1	0.4	0.4
Acidobacteriota; Aminicenantales ASV516	1.4	0.3	0.5
Acidobacteriota; Aminicenantales ASV464	0.8	0.6	0.4
Actinobacteriota; RBG-16-55-12 ASV7757	0.8	0.5	0.5
Acidobacteriota; Aminicenantales ASV479	0.5	0.3	0.6
Acidobacteriota; Aminicenantales ASV423			0.7
Actinobacteriota; RBG-16-55-12 ASV7799		0.3	0.6
Chloroflexi; MSBL5 ASV6116	1.6	0.5	0.3
Actinobacteriota; Actinomarinales ASV37	1	0.2	0.4
Acidobacteriota; Aminicenantales ASV500	3.8	0.1	
Acidobacteriota; Aminicenantales ASV5629	0.4	0.3	0.5
Acidobacteriota; Aminicenantales ASV5622		0.1	0.7
Acidobacteriota; Aminicenantales ASV506		0.1	0.7
Acidobacteriota; Aminicenantales ASV711	1.1	0.3	0.4
Acidobacteriota; Aminicenantales ASV690	0.9	0.3	0.4
Crenarchaeota; Bathyarchaeia ASV2284		0.2	0.5
Campylobacterota; Pseudarcobacter		0.5	0.4
Acidobacteriota; Aminicenantales ASV401		0.2	0.5
Actinobacteriota; RBG-16-55-12 ASV7835		0.3	0.5
Acidobacteriota; Aminicenantales ASV484		0.3	0.5
Chloroflexi; MSBL5 ASV6104		0.2	0.5
Chloroflexi; KD4-96 ASV6942	0.3	0.1	0.4
Acidobacteriota; Aminicenantales ASV513		0.2	0.4
Chloroflexi; KD4-96 ASV6933	0.7	0.2	0.4
Chloroflexi; GIF9 ASV5901		0.3	0.4
Chloroflexi; GIF9 ASV5896	0.4	0.3	0.3
Chloroflexi; Thermoflexus			0.5



Zone I Zone II Zone III