

Relative abundance of each class	Sample1-30c	Sample1-25c	Sample1-20c	Sample1-15c	Sample2-30c	Sample2-25c	Sample2-20c	Sample2-15c
Alphaproteobacteria	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
Betaproteobacteria	3.0%	3.1%	3.0%	3.0%	3.1%	3.5%	2.9%	2.0%
Gammaproteobacteria	0.7%	1.1%	0.4%	0.6%	0.7%	0.8%	0.8%	0.6%
Deltaproteobacteria	0.7%	0.9%	0.5%	0.7%	0.7%	1.0%	0.6%	0.4%
Epsilonproteobacteria	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
Clostridia	66.0%	64.4%	67.5%	65.6%	71.4%	67.7%	70.2%	73.2%
Mollicutes	0.7%	1.0%	0.3%	0.3%	1.3%	2.2%	1.3%	0.4%
Bacilli	2.2%	2.8%	1.4%	1.2%	9.0%	8.2%	9.7%	12.2%
Actinobacteria	14.4%	13.3%	16.1%	18.1%	1.1%	2.2%	1.6%	0.6%
Spirochaetes	0.3%	0.1%	0.5%	0.4%	0.3%	0.1%	0.4%	0.8%
Bacteroidetes	10.2%	11.6%	8.6%	8.1%	11.6%	13.7%	11.8%	9.4%
Verrucomicrobiae	1.4%	1.3%	1.5%	1.8%	0.7%	0.7%	0.6%	0.4%
Lentisphaerae	0.3%	0.5%	0.1%	0.1%	0.0%	0.0%	0.0%	0.0%