SHIME_Ma_et_al_2022

Phylum	SRR18730896	SRR18730897	SRR18730898	SRR18730899	SRR18730900	SRR18730901	SRR18730902	SRR18730903	SRR18730904	SRR18730905	SRR18730906	SRR18730907	SRR18730908	SRR18730909	SRR18730910	SRR18730911	SRR18730912	SRR18730913	SRR18730914	SRR18730915	SRR18730916	SRR18730917	SRR18730918	SRR18730919	SRR18730920	SRR18730921	SRR18730922	SRR18730923	SRR18730924	SRR18730925	SRR18730926	SRR18730927	SRR18730929	SRR18730930	SRR18730931	SRR18730932	SRR18730933	SRR18730934	SRR18730935	SRR18730936	SRR18730937	SRR18730938	SRR18730939
Synergistota	60.7		54	63.8	61	68.7		47.5		51.9		32.8		27.6		44.8		58.8		40.9	39	74.3		34.5		58.7		58.8		62.4	66	60.3		55.6		32.8		51.2		56.6		68.6	
Fusobacteriota	18.5		17	15	14	20.6		36.1		27.4		42.6		30.2		44.1		30.8		48.9		8.8	58.8		42.3		36.8		21.1		21.1		20.4		66.2		32.4		40.1		22.5		8.4
Bacillota	15.8		24.1		17.9	6.8	8.1	11.5		16	17.8	19	23.6		24.2	8.8	12.2	8.4	8.3	7.5	11.5		13.4		11.2		11.5		11.5	9.6	9.5	12.6		14.8	13	12	12.3		13	13.5		12.9	23
Actinomycetota	4.2	1.9	3	2.7	5.2	2.2	1.9	2.9	5.2	4.1	5.2	4.7	7.4	11.8		1.6	1.2	1.4	2.2	1.9	3	3.5	3.6	3.3	3.1	2.3	2.2	1.4	3	2.7	2.4	1.8	2.6	4.7	1.9	1.9	2	4.9	4.5	4.4	4.2	2.9	4.9
k__Unclassified_Unknown	0	1.7	0.9	1.5	0.4	0.9	1	1.1	0.4	0.1	0.1	0.2	1.3	0.4	0.3	0.2	0.4	0.3	0.3	0.3	0.6	0.4	0.6	0.3	0.4	1.3	1.3	1.2	0.2	0.1	0.1	0.5	0.2	0.6	1.5	1	0.8	0.9	0.5	0.3	0.1	0.1	0.2
Verrucomicrobiota	0.5	0.5	0.6	0.6	0.6	0.2	0.2	0.5	0.7	0.2	1.1	0.4	0.2	0.4	0.5	0.2	0.1	0.1	0.1	0.2	0.3	0.8	0.2	0.2	0.3	0.1	0.1	0.2	0.5	0.4	0.2	0.3	0.3	1	0.4	0.4	0.2	0.3	0.3	1	0.8	1.4	0.5
Pseudomonadota	0.2	0.2	0.2	0.4	0.2	0.1	0.1	0.2	0.1	0.1	0.2	0.3	0.3	0.1	0.4	0.2	0.2	0.1	0.1	0.1	0.6	0.1	0.2	0.1	0.1	0.2	0.2	0	0.4	0.4	0.3	0.2	0.1	0.2	0.3	0.2	0.6	0.2	0.4	0.6	0.4	0.2	
Bacteroidota	0	0	0	0.1	0.4	0.2	0	0.1	0.3	0	0.1	0	0.2	0	0.3	0	0.1	0	0.1	0.1	0	0	0.4	0.2	0	0	0.1	0.1	0.2	0.2	0.1	0.3	0.2	0.1	0	0.1	0	0.1	0.2	0.2	0	0.1	
Lentisphaerota	0	0.2	0	0	0.2	0	0.1	0	0.1	0	0	0	0	0	0	0.1	0.1	0	0.1	0	0	0	0.1	0	0	0.1	0.1	0.1	0	0	0.1	0	0	0	0.1	0.3	0.2	0.2	0	0	0	0	0.1
Thermodesulfobacteriota	0	0.1	0	0	0.1	0.1	0	0.1	0.2	0	0	0	0.2	0	0	0	0.1	0	0	0	0.1	0.1	0	0.1	0.1	0	0	0	0	0	0.1	0	0	0	0	0.1	0	0	0.1	0.3	0	0	0
Acidobacteriota	0	0.1	0.1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0.1	0
Campylobacterota	0	0	0	0	0	0	0	0	0.1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Cyanobacteriota	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Aquificota	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Chloroflexota	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0

Legend: % Read Abundance (color scale: 60 / 40 / 20)