

Family	% Contribution	Cumulative %	Mean time 1	Mean time 2	Change in abundance with time
k__Bacteria;p__Firmicutes;c__Clostridia;o__Clostridiales; f__Ruminococcaceae	11.52	11.52	21.00%	22.70%	1.70%
k__Bacteria;p__Firmicutes;c__Clostridia;o__Clostridiales; f__Lachnospiraceae	10.63	22.15	18.40%	13.00%	-5.40%
k__Bacteria;p__Firmicutes;c__Clostridia;o__Clostridiales; f__	7.833	29.98	14.20%	15.50%	1.30%
k__Archaea;p__Euryarchaeota;c__Methanobacteria;o__Methanobacteriales; f__Methanobacteriaceae	6.694	36.68	2.76%	7.18%	4.42%
k__Bacteria;p__Bacteroidetes;c__Bacteroidia;o__Bacteroidales; f__	6.124	42.8	1.55%	6.00%	4.45%
k__Bacteria;p__Firmicutes;c__Bacilli;o__Bacillales; f__Planococcaceae	5.518	48.32	3.32%	2.65%	-0.67%
k__Bacteria;p__Firmicutes;c__Bacilli;o__Lactobacillales; f__Lactobacillaceae	4.825	53.15	4.98%	1.45%	-3.53%
k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Pseudomonadales; f__Moraxellaceae	4.316	57.46	2.85%	1.32%	-1.53%
k__Bacteria;p__Bacteroidetes;c__Bacteroidia;o__Bacteroidales; f__Bacteroidaceae	3.806	61.27	3.75%	1.59%	-2.16%
k__Bacteria;p__Firmicutes;c__Clostridia;o__Clostridiales; f__Christensenellaceae	3.483	64.75	3.40%	2.71%	-0.69%
k__Bacteria;p__Firmicutes;c__Clostridia;o__Clostridiales; f__[Mogibacteriaceae]	2.996	67.75	2.64%	4.11%	1.47%
k__Bacteria;p__Firmicutes;c__Clostridia;o__Clostridiales; f__Clostridiaceae	2.574	70.32	2.02%	3.31%	1.29%
k__Bacteria;p__Verrucomicrobia;c__Verrucomicrobiae;o__Verrucomicrobiales; f__Verrucomicrobiaceae	2.489	72.81	1.87%	1.55%	-0.32%
k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Enterobacteriales; f__Enterobacteriaceae	2.251	75.06	1.19%	1.16%	-0.03%
k__Bacteria;p__Verrucomicrobia;c__Verrucomicrobiae;o__WCHB1-41; f__RFP12	2.06	77.12	0.46%	1.97%	1.51%
k__Bacteria;p__Spirochaetes;c__Spirochaetes;o__Spirochaetales; f__Spirochaetaceae	1.877	79	0.40%	1.91%	1.51%
k__Bacteria;p__Firmicute	1.831	80.83	1.75%	0.87%	-0.88%

s;c__Bacilli;o__Lactobacillales;f__Streptococcaceae					
k__Bacteria;p__Bacteroidetes;c__Bacteroidia;o__Bacteroidales;f__Parapre	1.342	82.17	0.65%	1.20%	0.55%
k__Bacteria;p__Bacteroidetes;c__Bacteroidia;o__Bacteroidales;f__BS11	1.306	83.48	0.51%	1.07%	0.56%
k__Bacteria;p__Firmicutes;c__Clostridia;o__Clostridiales;f__Veillonellaceae	1.298	84.77	1.24%	0.74%	-0.50%
k__Bacteria;p__Firmicutes;c__Bacilli;o__Bacillales;f__Bacillaceae	1.27	86.04	0.86%	0.56%	-0.30%
k__Bacteria;p__Bacteroidetes;c__Bacteroidia;o__Bacteroidales;f__Porphyromonadaceae	0.9022	86.95	0.84%	0.24%	-0.60%
k__Bacteria;p__Actinobacteria;c__Coriobacteriia;o__Coriobacteriales;f__Coriobacteriaceae	0.8914	87.84	1.19%	0.85%	-0.34%
k__Bacteria;p__Firmicutes;c__Bacilli;o__Lactobacillales;f__Carnobacteriaceae	0.7014	88.54	0.60%	0.06%	-0.54%
k__Bacteria;p__Chlamydiae;c__Chlamydia;o__Chlamydiales;f__Chlamydiaceae	0.6959	89.24	0.21%	0.50%	0.29%
k__Bacteria;p__Firmicutes;c__Clostridia;o__Clostridiales;f__Dehalobacteriaceae	0.6345	89.87	0.52%	0.63%	0.12%
k__Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Desulfo	0.6073	90.48	0.58%	0.12%	-0.45%
k__Bacteria;p__WPS-2;c__o__;f__	0.54	91.02	0.43%	0.09%	-0.34%
k__Bacteria;p__Bacteroidetes;c__Bacteroidia;o__Bacteroidales;f__[Odoribacteraceae]	0.5115	91.53	0.46%	0.09%	-0.37%
k__Bacteria;p__Actinobacteria;c__Actinobacteria;o__Actinomycetales;f__Micrococcaceae	0.5029	92.03	0.42%	0.09%	-0.33%
k__Bacteria;p__Firmicutes;c__Clostridia;o__Clostridiales;Other	0.4865	92.52	0.55%	0.27%	-0.29%
k__Bacteria;p__Fusobacteria;c__Fusobacteriia;o__Fusobacteriales;f__Fusobacteriaceae	0.4355	92.95	0.31%	0.21%	-0.10%
k__Bacteria;p__Bacteroidetes;c__Bacteroidia;o__Bacteroidales;f__S24-7	0.4073	93.36	0.12%	0.33%	0.21%
k__Bacteria;p__Firmicutes;c__Bacilli;o__Lactobacillales;f__Enterococcaceae	0.3939	93.75	0.22%	0.19%	-0.03%
k__Bacteria;p__Firmicutes;c__Clostridia;o__Clostridiales;f__Eubacteriaceae	0.3889	94.14	0.15%	0.36%	0.22%
k__Bacteria;p__Firmicutes;c__Clostridia;o__Clostri	0.2873	94.43	0.14%	0.14%	0.01%

diales;f__[Tissierellaceae]					
k__Bacteria;p__Firmicutes;c__Erysipelotrichi;o__Erysipelotrichales;f__Erysipelotrichaceae	0.2737	94.7	0.30%	0.19%	-0.11%
k__Bacteria;p__Firmicutes;c__Bacilli;o__Lactobacillales;f__Leuconostocaceae	0.2686	94.97	0.10%	0.16%	0.06%
k__Bacteria;p__Bacteroidetes;c__Bacteroidia;o__Bacteroidales;f__Rikenellaceae	0.2671	95.24	0.24%	0.04%	-0.19%
Unassigned;Other;Other;Other;Other	0.2656	95.51	0.23%	0.33%	0.10%
k__Bacteria;p__Spirochaetes;c__MVP-15;o__PL-11B10;f__	0.2652	95.77	0.09%	0.18%	0.09%
k__Bacteria;p__Proteobacteria;c__Epsilonproteobacteria;o__Campylobacterales;f__Campylobacteraceae	0.2623	96.03	0.22%	0.08%	-0.14%
k__Bacteria;p__Fibrobacteres;c__Fibrobacteria;o__Fibrobacterales;f__Fibrobacteraceae	0.2264	96.26	0.02%	0.21%	0.19%
k__Bacteria;p__Proteobacteria;c__Betaproteobacteria;o__Tremblayales;f__	0.2188	96.48	0.11%	0.12%	0.01%
k__Bacteria;p__Firmicutes;c__Bacilli;o__Bacillales;f__Staphylococcaceae	0.2187	96.7	0.20%	0.01%	-0.19%
k__Bacteria;p__Tenericutes;c__Mollicutes;o__RF39;f__	0.2133	96.91	0.23%	0.13%	-0.11%
k__Bacteria;p__Actinobacteria;c__Actinobacteria;o__Actinomycetales;f__Actinomycetaceae	0.2113	97.12	0.07%	0.19%	0.13%
k__Bacteria;p__Firmicutes;c__Clostridia;o__Clostridiales;f__Peptostreptococcaceae	0.1853	97.31	0.08%	0.11%	0.03%
k__Bacteria;p__Armatimonadetes;c__SJA-176;o__RB046;f__	0.1843	97.49	0.08%	0.15%	0.08%
k__Bacteria;p__Synergistetes;c__Synergistia;o__Synergistales;f__Synergistaceae	0.1799	97.67	0.09%	0.12%	0.03%
k__Archaea;p__Euryarchaeota;c__Methanomicrobia;o__Methanomicrobiales;f__Methanocorpusculaceae	0.1774	97.85	0.06%	0.14%	0.08%
k__Bacteria;p__Bacteroidetes;c__Bacteroidia;o__Bacteroidales;f__Pre	0.1415	97.99	0.07%	0.11%	0.04%
k__Bacteria;p__Firmicutes;c__Clostridia;o__Clostridiales;f__Peptococcaceae	0.1397	98.13	0.13%	0.08%	-0.05%

k__Bacteria;p__Bacteroidetes;c__Bacteroidia;o__Bacteroidales;f__RF16	0.129	98.26	0.05%	0.10%	0.06%
k__Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__RF32;f__	0.1204	98.38	0.10%	0.03%	-0.07%
k__Bacteria;p__Firmicutes;c__Bacilli;o__Lactobacillales;Other	0.1127	98.49	0.07%	0.05%	-0.01%
k__Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__f__	0.1101	98.6	0.10%	0.01%	-0.08%
k__Bacteria;p__Proteobacteria;c__Epsilonproteobacteria;o__Campylobacterales;f__Helicobacteraceae	0.1076	98.71	0.05%	0.08%	0.03%
k__Bacteria;p__Planctomycetes;c__Planctomycetia;o__Pirellulales;f__Pirellulaceae	0.09007	98.8	0.06%	0.09%	0.03%
k__Bacteria;p__Tenericutes;c__RF3;o__ML615J-28;f__	0.08261	98.88	0.07%	0.02%	-0.04%
k__Bacteria;p__Actinobacteria;c__Actinobacteria;o__Actinomycetales;f__Corynebacteriaceae	0.06411	98.95	0.06%	0.01%	-0.04%