



Legend	Taxonomy	Total	MGPLP1 3	MGPLP1 4	MGPLP1 7	MGPLP1 8	MGPLP2 0	MGPLP2 1
		%	%	%	%	%	%	%
	Unassigned;Other;Other;Other;Other;Other	1.3%	1.0%	1.4%	0.9%	2.2%	1.4%	1.1%
	k__Archaea;p__Crenarchaeota;c__MBGA;o__NRP-J;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Archaea;p__Crenarchaeota;c__MCG;o__pGrfC26;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Archaea;p__Crenarchaeota;c__Thaumarchaeota;o__AK31;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Archaea;p__Crenarchaeota;c__Thaumarchaeota;o__Cenarchaeales;f__SAGMA-X;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Archaea;p__Crenarchaeota;c__Thaumarchaeota;o__Nitrososphaerales;f__Nitrososphaeraeae;g__Candidatus Nitrososphaera	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Archaea;p__Crenarchaeota;c__Thermoprotei;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Archaea;p__Euryarchaeota;c__Methanobacteria;o__Methanobacteriales;f__Methanobacteriaceae;g__Methanobacterium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Archaea;p__Euryarchaeota;c__Methanomicrobia;o__Methanocellales;f__Methanocellaceae;g__Methanocella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Archaea;p__Euryarchaeota;c__Methanomicrobia;o__Methanosarcinales;f__Methanosarcinaceae;g__Methanosarcina	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Archaea;p__Euryarchaeota;c__Methanomicrobia;o__YC-E6;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Archaea;p__Euryarchaeota;c__Thermoplasmata;o__E2;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Archaea;p__Euryarchaeota;c__Thermoplasmata;o__E2;f__[Methanomassiliococcaceae];g__	0.0%	0.0%	0.0%	0.0%	0.2%	0.0%	0.0%
	k__Archaea;p__[Parvarchaeota];c__[Micrarchaeae];o__[Micrarchaeales];f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Archaea;p__[Parvarchaeota];c__[Parvarchaeae];o__WCHD3-30;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Archaea;p__[Parvarchaeota];c__[Parvarchaeae];o__YLA114;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Bacteria;Other;Other;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Bacteria;p__c__o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Bacteria;p__AD3;c__ABS-6;o__f__g__	0.2%	0.0%	0.0%	0.1%	1.3%	0.0%	0.0%
	k__Bacteria;p__AD3;c__JG37-AG-4;o__f__g__	0.1%	0.0%	0.0%	0.0%	0.4%	0.0%	0.0%
	k__Bacteria;p__Acidobacteria;Other;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Bacteria;p__Acidobacteria;c__o__f__g__	0.1%	0.0%	0.2%	0.1%	0.0%	0.5%	0.1%
	k__Bacteria;p__Acidobacteria;c__AT-s54;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Bacteria;p__Acidobacteria;c__Acidobacteria-5;o__f__g__	0.4%	0.2%	0.3%	0.8%	0.0%	0.2%	0.7%
	k__Bacteria;p__Acidobacteria;c__Acidobacteria-6;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Bacteria;p__Acidobacteria;c__Acidobacteria-6;o__BPC015;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Bacteria;p__Acidobacteria;c__Acidobacteria-6;o__CCU21;f__g__	0.1%	0.0%	0.1%	0.1%	0.0%	0.1%	0.1%
	k__Bacteria;p__Acidobacteria;c__Acidobacteria-6;o__iii1-15;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
	k__Bacteria;p__Acidobacteria;c__Acidobacteria-6;o__iii1-15;f__g__	3.1%	2.2%	3.3%	4.5%	0.1%	2.2%	5.9%
	k__Bacteria;p__Acidobacteria;c__Acidobacteria-6;o__iii1-15;f__RB40;g__	0.3%	0.4%	0.4%	0.5%	0.0%	0.1%	0.6%
	k__Bacteria;p__Acidobacteria;c__Acidobacteria-6;o__iii1-15;f__mb2424;g__	0.9%	1.0%	1.1%	0.7%	0.0%	1.5%	1.0%

k_Bacteria;p_Acidobacteria;c_Acidobacteriia;o_Acidobacteriales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_Acidobacteriia;o_Acidobacteriales;f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_Acidobacteriia;o_Acidobacteriales;f_Acidobacteriaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_Acidobacteriia;o_Acidobacteriales;f_Acidobacteriaceae;g	0.1%	0.0%	0.0%	0.1%	0.6%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_Acidobacteriia;o_Acidobacteriales;f_Acidobacteriaceae;g_Edaphobacter	0.1%	0.0%	0.0%	0.0%	0.7%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_Acidobacteriia;o_Acidobacteriales;f_Acidobacteriaceae;g_Terriglobus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_Acidobacteriia;o_Acidobacteriales;f_Koribacteraceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_Acidobacteriia;o_Acidobacteriales;f_Koribacteraceae;g	0.5%	0.0%	0.1%	0.7%	1.9%	0.1%	0.0%
k_Bacteria;p_Acidobacteria;c_Acidobacteriia;o_Acidobacteriales;f_Koribacteraceae;g_Candidatus Koribacter	0.4%	0.0%	0.4%	0.4%	1.3%	0.0%	0.1%
k_Bacteria;p_Acidobacteria;c_BPC102;o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_BPC102;o_B110;f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_BPC102;o_MVS-40;f_g	0.3%	0.0%	0.4%	0.0%	0.0%	0.3%	0.8%
k_Bacteria;p_Acidobacteria;c_DA052;o_E29;f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_DA052;o_Ellin6513;f_g	2.6%	0.0%	0.0%	0.9%	14.5%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_EC1113;o_f_g	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_GAL08;o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_Holophagae;o_Holophagales;f_Holophagaceae;g	0.0%	0.0%	0.0%	0.0%	0.2%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_Holophagae;o_Holophagales;f_Holophagaceae;g_Geothrix	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_PAUC37;o_f_g	0.2%	0.0%	0.2%	0.2%	0.0%	0.2%	0.7%
k_Bacteria;p_Acidobacteria;c_RB25;o_f_g	0.1%	0.0%	0.1%	0.2%	0.0%	0.1%	0.2%
k_Bacteria;p_Acidobacteria;c_S035;o_f_g	0.3%	0.0%	0.2%	0.2%	0.0%	0.3%	1.0%
k_Bacteria;p_Acidobacteria;c_Solibacteres;o_JH-WHS99;f_g	0.1%	0.0%	0.1%	0.0%	0.2%	0.1%	0.0%
k_Bacteria;p_Acidobacteria;c_Solibacteres;o_Solibacterales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.2%	0.0%
k_Bacteria;p_Acidobacteria;c_Solibacteres;o_Solibacterales;f_g	1.9%	0.8%	1.9%	2.2%	4.8%	0.6%	1.3%
k_Bacteria;p_Acidobacteria;c_Solibacteres;o_Solibacterales;f_AKIW659;g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_Solibacteres;o_Solibacterales;f_PAUC26f;g	0.1%	0.1%	0.0%	0.1%	0.2%	0.1%	0.0%
k_Bacteria;p_Acidobacteria;c_Solibacteres;o_Solibacterales;f_PK34;g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_Solibacteres;o_Solibacterales;f_Solibacteraceae;g	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_Solibacteres;o_Solibacterales;f_Solibacteraceae;g_Candidatus Solibacter	0.6%	0.1%	0.1%	1.3%	1.5%	0.2%	0.6%
k_Bacteria;p_Acidobacteria;c_Solibacteres;o_Solibacterales;f_Bryobacteraceae;g	0.1%	0.1%	0.3%	0.2%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_Sva0725;o_Sva0725;f_g	0.2%	0.3%	0.1%	0.1%	0.0%	0.1%	0.3%
k_Bacteria;p_Acidobacteria;c_TM1;o_f_g	0.4%	0.0%	0.0%	0.0%	2.3%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_[Chloracidobacteria];o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_[Chloracidobacteria];o_11-24;f_g	0.2%	0.0%	0.5%	0.2%	0.0%	0.2%	0.3%
k_Bacteria;p_Acidobacteria;c_[Chloracidobacteria];o_DS-100;f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p_Acidobacteria;c_[Chloracidobacteria];o_Ellin7246;f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_[Chloracidobacteria];o_PK29;f_g	0.4%	0.1%	1.4%	0.2%	0.0%	0.4%	0.4%
k_Bacteria;p_Acidobacteria;c_[Chloracidobacteria];o_RB41;f_g	1.4%	0.1%	3.5%	1.3%	0.0%	1.7%	1.7%
k_Bacteria;p_Acidobacteria;c_[Chloracidobacteria];o_RB41;f_Ellin6075;g	1.6%	0.5%	5.5%	1.4%	0.2%	1.3%	0.9%
k_Bacteria;p_Acidobacteria;c_iii1-8;o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Acidobacteria;c_iii1-8;o_32-20;f_g	0.2%	0.0%	0.9%	0.0%	0.0%	0.3%	0.2%
k_Bacteria;p_Acidobacteria;c_iii1-8;o_DS-18;f_g	0.5%	0.2%	1.3%	0.2%	0.0%	0.5%	0.7%
k_Bacteria;p_Acidobacteria;c_iii1-8;o_SJA-36;f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p_Actinobacteria;Other;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Acidimicrobiia;o_Acidimicrobiales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Acidimicrobiia;o_Acidimicrobiales;f_g	1.4%	3.2%	0.7%	0.7%	0.3%	1.8%	1.7%
k_Bacteria;p_Actinobacteria;c_Acidimicrobiia;o_Acidimicrobiales;f_AKIW874;g	0.3%	1.2%	0.0%	0.0%	0.0%	0.2%	0.2%
k_Bacteria;p_Actinobacteria;c_Acidimicrobiia;o_Acidimicrobiales;f_C111;g	0.7%	1.8%	0.2%	0.7%	0.0%	0.3%	1.1%
k_Bacteria;p_Actinobacteria;c_Acidimicrobiia;o_Acidimicrobiales;f_EB1017;g	0.6%	1.4%	0.2%	1.0%	0.1%	0.3%	0.8%
k_Bacteria;p_Actinobacteria;c_Acidimicrobiia;o_Acidimicrobiales;f_Iamiaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Acidimicrobiia;o_Acidimicrobiales;f_Iamiaceae;g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Acidimicrobiia;o_Acidimicrobiales;f_Iamiaceae;g_Iamia	0.1%	0.4%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Acidimicrobiia;o_Acidimicrobiales;f_Microthrixaceae;g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Acidimicrobiia;o_Acidimicrobiales;f_TK06;g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Acidimicrobiia;o_Acidimicrobiales;f_koll13;g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.2%
k_Bacteria;p_Actinobacteria;c_Acidimicrobiia;o_Acidimicrobiales;f_ntu14;g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales;Other;Other	0.0%	0.1%	0.0%	0.1%	0.0%	0.0%	0.1%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales;f_g	0.2%	0.3%	0.0%	0.3%	0.1%	0.1%	0.1%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales;f_ACK-M1;g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales;f_Actinomycetaceae;g_Actinomycetes	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales;f_Actinospicaceae;g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales;f_Actinosynnemataceae;Other	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales;f_Actinosynnemataceae;g	0.0%	0.3%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales;f_Actinosynnemataceae;g_Kibdelosporangium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales;f_Actinosynnemataceae;g_Lentzea	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales;f_Beutenbergiaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales;f_Beutenbergiaceae;g_Salana	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%

k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Bogoriellaceae;g_Georgenia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Brevibacteriaceae;g_Brevibacterium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Cellulomonadaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Cellulomonadaceae;g_Cellulomonas	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Cellulomonadaceae;g_Demequina	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Corynebacteriaceae;g_Corynebacterium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Cryptosporangiaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Cryptosporangiaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Dermabacteraceae;g_Brachybacterium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Dermacoccaceae;g_Dermacoccus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Dietziaceae;g_Dietzia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Frankiaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Frankiaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Frankiaceae;g_Frankia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Geodermatophilaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Geodermatophilaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Geodermatophilaceae;g_Geodermatophilus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Geodermatophilaceae;g_Modestobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Glycomycetaceae;g_Glycomyces	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Gordoniaceae;g_Gordonia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Intrasporangiaceae;Other	0.1%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Intrasporangiaceae;g_	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Intrasporangiaceae;g_Knoellia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Intrasporangiaceae;g_Phycococcus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Intrasporangiaceae;g_Terracoccus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Kinemosporiaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Kinemosporiaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Microbacteriaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Microbacteriaceae;g_	0.0%	0.1%	0.0%	0.1%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Microbacteriaceae;g_Agromyces	0.1%	0.4%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Microbacteriaceae;g_Cryocola	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Microbacteriaceae;g_Leucobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Microbacteriaceae;g_Microbacterium	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Microbacteriaceae;g_Mycetocola	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Microbacteriaceae;g_Salinibacterium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Micrococcaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Micrococcaceae;g_	0.5%	0.1%	0.7%	0.3%	0.1%	1.6%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Micrococcaceae;g_Arthrobacter	0.1%	0.0%	0.1%	0.1%	0.0%	0.1%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Micrococcaceae;g_Microbispora	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Micrococcaceae;g_Micrococcus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Micromonosporaceae;Other	0.0%	0.2%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Micromonosporaceae;g_	0.1%	0.3%	0.0%	0.1%	0.0%	0.1%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Micromonosporaceae;g_Actinoplanes	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Micromonosporaceae;g_Catellatospora	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Micromonosporaceae;g_Dactylosporangium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Micromonosporaceae;g_Micromonospora	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Micromonosporaceae;g_Pilimelia	0.0%	0.1%	0.0%	0.1%	0.0%	0

k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Propionibacteriaceae;g_	0.3%	0.2%	0.1%	0.1%	0.0%	0.4%	0.8%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Propionibacteriaceae;g_Luteococcus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Propionibacteriaceae;g_Microlunatus	0.3%	0.2%	0.0%	0.2%	0.0%	0.1%	1.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Propionibacteriaceae;g_Propionibacterium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Pseudonocardiaceae;Other	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Pseudonocardiaceae;g_	0.6%	1.0%	0.0%	0.0%	0.0%	2.6%	0.1%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Pseudonocardiaceae;g_Actinomycetospora	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Pseudonocardiaceae;g_Amycolatopsis	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Pseudonocardiaceae;g_Jiangella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Pseudonocardiaceae;g_Pseudonocardia	0.1%	0.5%	0.0%	0.1%	0.0%	0.1%	0.1%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Pseudonocardiaceae;g_Saccharopolyspora	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Pseudonocardiaceae;g_Thermobispora	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Sanguibacteraceae;g_Sanguibacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Sporichthyaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Sporichthyaceae;g_Sporichthya	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Streptomycetaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Streptomycetaceae;g_	0.2%	0.4%	0.0%	0.4%	0.0%	0.1%	0.4%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Streptomycetaceae;g_Streptacidiphilus	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Streptomycetaceae;g_Streptomyces	0.3%	1.1%	0.0%	0.2%	0.0%	0.1%	0.2%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Streptosporangiaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Streptosporangiaceae;g_	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Streptosporangiaceae;g_Nonumurea	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Streptosporangiaceae;g_Sphaerisporangium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Streptosporangiaceae;g_Streptosporangium	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Thermomonosporaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Thermomonosporaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Thermomonosporaceae;g_Actinoallomurus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Thermomonosporaceae;g_Actinocorallia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Thermomonosporaceae;g_Actinomadura	0.1%	0.3%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales:f_Williamsiaceae;g_Williamsia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Bifidobacteriales:f_;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Micrococcales:f_;g_	0.1%	0.1%	0.0%	0.0%	0.0%	0.1%	0.2%
k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_WCHB1-81;f_At425_EubF1;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Coriobacteriia;o_Coriobacteriales:f_Coriobacteriaceae;g_Adlercreutzia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_KIST-JY010;o_;f_;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_MB-A2-108;o_;f_;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_MB-A2-108;o_0319-7L14;f_;g_	1.1%	0.7%	0.6%	0.6%	0.0%	2.8%	1.7%
k_Bacteria;p_Actinobacteria;c_Nitriliruptoria;o_Euzebyales;f_Euzebyaceae;g_Euzebya	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%
k_Bacteria;p_Actinobacteria;c_OPB41;o_;f_;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p_Actinobacteria;c_Rubrobacteria;o_Rubrobacterales;f_Rubrobacteraceae;g_	0.1%	0.1%	0.0%	0.0%	0.0%	0.1%	0.2%
k_Bacteria;p_Actinobacteria;c_Rubrobacteria;o_Rubrobacterales;f_Rubrobacteraceae;g_Rubrobacter	0.2%	0.1%	0.8%	0.0%	0.0%	0.2%	0.2%
k_Bacteria;p_Actinobacteria;c_Thermoleophilia;o_;f_;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p_Actinobacteria;c_Thermoleophilia;o_Gaiellales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Thermoleophilia;o_Gaiellales;f_;g_	0.0%	0.0%	0.0%	0.1%	0.0%	0.1%	0.1%
k_Bacteria;p_Actinobacteria;c_Thermoleophilia;o_Gaiellales;f_AK1AB1_02E;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Thermoleophilia;o_Gaiellales;f_Gaiellaceae;g_	1.7%	1.7%	0.8%	0.9%	0.1%	2.9%	3.8%
k_Bacteria;p_Actinobacteria;c_Thermoleophilia;o_Solirubrobacterales;Other;Other	0.0%	0.2%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p_Actinobacteria;c_Thermoleophilia;o_Solirubrobacterales;f_;g_	0.7%	2.1%	0.1%	0.5%	0.0%	0.5%	0.9%
k_Bacteria;p_Actinobacteria;c_Thermoleophilia;o_Solirubrobacterales;f_Conexibacteraceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Thermoleophilia;o_Solirubrobacterales;f_Conexibacteraceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Thermoleophilia;o_Solirubrobacterales;f_Conexibacteraceae;g_Conexibacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Thermoleophilia;o_Solirubrobacterales;f_Patulibacteraceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Thermoleophilia;o_Solirubrobacterales;f_Patulibacteraceae;g_Patulibacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Thermoleophilia;o_Solirubrobacterales;f_Solirubrobacteraceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Actinobacteria;c_Thermoleophilia;o_Solirubrobacterales;f_Solirubrobacteraceae;g_	0.1%	0.5%	0.0%	0.1%	0.0%	0.1%	0.1%
k_Bacteria;p_Actinobacteria;c_Thermoleophilia;o_Solirubrobacterales;f_Solirubrobacteraceae;g_Solirubrobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Armatimonadetes;c_0319-6E2;o_;f_;g_	0.1%	0.0%	0.2%	0.0%	0.0%	0.1%	0.1%
k_Bacteria;p_Armatimonadetes;c_Armatimonadia;o_Armatimonadales;f_Armatimonadaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Armatimonadetes;c_Armatimonadia;o_Armatimonadales;f_Armatimonadaceae;g_Armatimonas	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Armatimonadetes;c_Armatimonadia;o_FW68;f_;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Armatimonadetes;c_Chthonomonadetes;o_Chthonomonadales;f_Chthonomonadaceae;g_	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Armatimonadetes;c_Chthonomonadetes;o_Chthonomonadales;f_Chthonomonadaceae;g_Chthonomonas	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Armatimonadetes;c_Chthonomonadetes;o_SJA-22;f_;g_	0.1%	0.0%	0.7%	0.0%	0.0%	0.1%	0.0%
k_Bacteria;p_Armatimonadetes;c_OPB50;o_;f_;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Armatimonadetes;c_OS-L;o_;f_;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Armatimonadetes;c_SHA-37;o_;f_;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Armatimonadetes;c_SJA-176;o_GAB-B06;f_;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%

k_Bacteria;p_Armatimonadetes;c_SJA-176;o_RB046:f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Armatimonadetes;o_Fimbrimonadia;o_Fimbrimonadales:f_g	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p_Armatimonadetes;c_Fimbrimonadia;o_Fimbrimonadales:f_Fimbrimonadales:g	0.1%	0.0%	0.3%	0.0%	0.0%	0.0%
k_Bacteria;p_Armatimonadetes;c_Fimbrimonadia;o_Fimbrimonadales:f_Fimbrimonadales:g_Fimbrimonas	0.1%	0.0%	0.3%	0.1%	0.0%	0.0%
k_Bacteria;p_BHI80-139;c_o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_BRC1;c_NPL-UPA2;o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_BRC1;c_PRR-11;o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;Other;Other;Other	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_At12OctB3;o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p_Bacteroidetes;c_BME43;o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Bacteroidia;o_Bacteroidales:f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p_Bacteroidetes;c_Bacteroidia;o_Bacteroidales:f_BA008:g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Bacteroidia;o_Bacteroidales:f_Rikenellaceae:g_Blvii28	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Bacteroidia;o_Bacteroidales:f_S24-7:g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cyclobacteriaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cyclobacteriaceae:g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cyclobacteriaceae:g_Algoriphagus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cytophagaceae:g	0.8%	2.7%	0.5%	1.4%	0.0%	0.3%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cytophagaceae:g_Adhaeribacter	0.1%	0.1%	0.1%	0.1%	0.0%	0.1%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cytophagaceae:g_Cytophaga	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cytophagaceae:g_Dyadobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cytophagaceae:g_Enticicia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cytophagaceae:g_Flectobacillus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cytophagaceae:g_Hymenobacter	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cytophagaceae:g_Larkinella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cytophagaceae:g_Leadbetterella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cytophagaceae:g_Pontibacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cytophagaceae:g_Rhodocytaphaga	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cytophagaceae:g_Runella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cytophagaceae:g_Spirosoma	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Cytophagaceae:g_Sporocytaphaga	0.1%	0.1%	0.0%	0.2%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Flammeovirgaceae:g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Flammeovirgaceae:g_Fulvivirga	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Flammeovirgaceae:g_Roseivirga	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Amoebophilaceae:g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Amoebophilaceae:g_Candidatus Amoebophilus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Cytophagia;o_Cytophagales:f_Amoebophilaceae:g_SC3-56	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Flavobacteriia;o_Flavobacteriales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Flavobacteriia;o_Flavobacteriales:f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Flavobacteriia;o_Flavobacteriales:f_Cryomorphaceae:g	0.1%	0.1%	0.2%	0.1%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Flavobacteriia;o_Flavobacteriales:f_Cryomorphaceae:g_Crocinitomix	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Flavobacteriia;o_Flavobacteriales:f_Cryomorphaceae:g_Fluviicola	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Flavobacteriia;o_Flavobacteriales:f_Cryomorphaceae:g_Owenweeksia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Flavobacteriia;o_Flavobacteriales:f_Flavobacteriaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Flavobacteriia;o_Flavobacteriales:f_Flavobacteriaceae:g	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Flavobacteriia;o_Flavobacteriales:f_Flavobacteriaceae:g_Flavobacterium	0.8%	1.2%	0.7%	2.5%	0.0%	0.2%
k_Bacteria;p_Bacteroidetes;c_Flavobacteriia;o_Flavobacteriales:f_Flavobacteriaceae:g_Gillisia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Flavobacteriia;o_Flavobacteriales:f_Flavobacteriaceae:g_Myroides	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Flavobacteriia;o_Flavobacteriales:f_Weeksellaceae:g_Chryseobacterium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Flavobacteriia;o_Flavobacteriales:f_Weeksellaceae:g_Elizabethkingia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Sphingobacteriia;o_Sphingobacteriales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Sphingobacteriia;o_Sphingobacteriales:f_g	1.8%	0.8%	3.0%	4.2%	0.3%	0.7%
k_Bacteria;p_Bacteroidetes;c_Sphingobacteriia;o_Sphingobacteriales:f_Sphingobacteriaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Sphingobacteriia;o_Sphingobacteriales:f_Sphingobacteriaceae:g	0.4%	0.2%	0.8%	0.3%	0.3%	0.2%
k_Bacteria;p_Bacteroidetes;c_Sphingobacteriia;o_Sphingobacteriales:f_Sphingobacteriaceae:g_Olivibacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Sphingobacteriia;o_Sphingobacteriales:f_Sphingobacteriaceae:g_Pedobacter	0.1%	0.0%	0.4%	0.1%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Sphingobacteriia;o_Sphingobacteriales:f_Sphingobacteriaceae:g_Solitalea	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Sphingobacteriia;o_Sphingobacteriales:f_Sphingobacteriaceae:g_Sphingobacterium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_VC2_1_Bac22;o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Rhodothermi;o_Rhodothermales:f_Rhodothermaceae:g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Rhodothermi;o_Rhodothermales:f_Rhodothermaceae:g_Rubricoccus	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Saprospirae;o_Saprospirales:f_g	0.1%	0.1%	0.1%	0.3%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Saprospirae;o_Saprospirales:f_Chitinophagaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Bacteroidetes;c_Saprospirae;o_Saprospirales:f_Chitinophagaceae:g	2.6%	1.7%	3.4%	5.9%	1.0%	2.3%

k_Bacteria:p_Bacteroidetes;c_[Saprosirae];o_[Saprosirales];f_Chitinophagaceae;g_Chitinophaga	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria:p_Bacteroidetes;c_[Saprosirae];o_[Saprosirales];f_Chitinophagaceae;g_Flavihumibacter	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Bacteroidetes;c_[Saprosirae];o_[Saprosirales];f_Chitinophagaceae;g_Flavisolibacter	0.3%	0.1%	1.3%	0.1%	0.0%	0.1%	0.5%
k_Bacteria:p_Bacteroidetes;c_[Saprosirae];o_[Saprosirales];f_Chitinophagaceae;g_Niabella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Bacteroidetes;c_[Saprosirae];o_[Saprosirales];f_Chitinophagaceae;g_Niastella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Bacteroidetes;c_[Saprosirae];o_[Saprosirales];f_Chitinophagaceae;g_Sediminibacterium	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria:p_Bacteroidetes;c_[Saprosirae];o_[Saprosirales];f_Chitinophagaceae;g_Segetibacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Bacteroidetes;c_[Saprosirae];o_[Saprosirales];f_Saprosiraceae;g_	0.5%	0.8%	0.6%	0.9%	0.0%	0.0%	0.8%
k_Bacteria:p_Bacteroidetes;c_[Saprosirae];o_[Saprosirales];f_Saprosiraceae;g_Haliscomenobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Bacteroidetes;c_[Saprosirae];o_[Saprosirales];f_Saprosiraceae;g_Saprosira	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_f_:g_	0.1%	0.1%	0.0%	0.2%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_Chlamydiales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_Chlamydiales;f_:g_	0.0%	0.0%	0.1%	0.1%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_Chlamydiales;f_Chlamydiaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_Chlamydiales;f_Chlamydiaceae;g_Chlamydia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_Chlamydiales;f_Criblamydiaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_Chlamydiales;f_Criblamydiaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_Chlamydiales;f_Parachlamydiaceae;Other	0.2%	0.2%	0.1%	0.5%	0.1%	0.1%	0.3%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_Chlamydiales;f_Parachlamydiaceae;g_	0.0%	0.0%	0.0%	0.1%	0.1%	0.0%	0.0%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_Chlamydiales;f_Parachlamydiaceae;g_Candidatus Protochlamydia	0.1%	0.1%	0.0%	0.1%	0.0%	0.0%	0.1%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_Chlamydiales;f_Parachlamydiaceae;g_Parachlamydia	0.1%	0.1%	0.0%	0.1%	0.3%	0.0%	0.1%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_Chlamydiales;f_Rhabdochlamydiaceae;g_Candidatus Rhabdochlamydia	0.1%	0.0%	0.0%	0.1%	0.2%	0.0%	0.0%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_Chlamydiales;f_Simkaniaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_Chlamydiales;f_Simkaniaceae;g_Simkania	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_Chlamydiales;f_Waddiaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_Chlamydiales;f_Waddiaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlamydiae;c_Chlamydiia;o_Chlamydiales;f_Waddiaceae;g_Waddlia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlorobi;c_:o_:f_:g_	0.1%	0.0%	0.2%	0.1%	0.1%	0.0%	0.1%
k_Bacteria:p_Chlorobi;c_BSV26;o_A89;f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlorobi;c_BSV26;o_C20;f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlorobi;c_BSV26;o_PK329;f_:g_	0.0%	0.0%	0.1%	0.1%	0.0%	0.0%	0.1%
k_Bacteria:p_Chlorobi;c_BSV26;o_VC38;f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlorobi;c_Ignavibacteria;o_Ignavibacteriales;f_Ignavibacteriaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chlorobi;c_OPB56;o_:f_:g_	0.1%	0.1%	0.1%	0.0%	0.0%	0.0%	0.1%
k_Bacteria:p_Chlorobi;c_SJA-28;o_:f_:g_	0.2%	0.0%	0.3%	0.2%	0.0%	0.1%	0.6%
k_Bacteria:p_Chloroflexi;Other;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_:o_:f_:g_	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_:f_:g_	0.1%	0.0%	0.2%	0.0%	0.0%	0.1%	0.0%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_A31;f_:g_	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_A31;f_S47;g_	0.2%	0.0%	0.1%	0.2%	0.0%	0.0%	1.0%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_Anaerolineales;f_Anaerolinaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_Anaerolineales;f_Anaerolinaceae;g_Anaerolinea	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.2%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_Ardenscatenales;f_Ardenscatenaceae;g_Ardenscatena	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_CFB-26;f_:g_	0.1%	0.1%	0.1%	0.0%	0.0%	0.1%	0.0%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_Caldilineales;f_Caldilineaceae;g_	0.2%	0.3%	0.3%	0.1%	0.0%	0.1%	0.5%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_Caldilineales;f_Caldilineaceae;g_Caldilinea	0.1%	0.0%	0.0%	0.0%	0.0%	0.0%	0.5%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_Caldilineales;f_Caldilineaceae;g_Litorilinea	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_GCA004;f_:g_	0.2%	0.0%	0.0%	0.0%	0.0%	0.0%	1.1%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_H39;f_:g_	0.1%	0.1%	0.1%	0.0%	0.0%	0.1%	0.0%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_S0208;f_:g_	0.3%	0.1%	0.4%	0.2%	0.0%	0.4%	0.8%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_SB-34;f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_SBR1031;f_A4b;g_	0.2%	0.3%	0.2%	0.3%	0.0%	0.1%	0.2%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_SBR1031;f_SJA-101;g_	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_SBR1031;f_oc28;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_SHA-20;f_:g_	0.5%	0.0%	2.8%	0.0%	0.0%	0.2%	0.0%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_SJA-15;f_:g_	0.1%	0.0%	0.0%	0.0%	0.0%	0.0%	0.5%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_WCHB1-50;f_:g_	0.2%	0.0%	0.9%	0.1%	0.0%	0.2%	0.2%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_envOPS12;f_:g_	0.1%	0.0%	0.1%	0.1%	0.0%	0.0%	0.3%
k_Bacteria:p_Chloroflexi;c_Anaerolineae;o_pLW-97;f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_C0119;o_:f_:g_	0.1%	0.0%	0.2%	0.1%	0.0%	0.0%	0.1%
k_Bacteria:p_Chloroflexi;c_Chloroflexi;o_AKIW781;f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Chloroflexi;o_Chloroflexales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Chloroflexi;o_Chloroflexales;f_01D2Z48;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Chloroflexi;o_Chloroflexales;f_Chloroflexaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Chloroflexi;o_Chloroflexales;f_Chloroflexaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Chloroflexi;o_Chloroflexales;f_Chloroflexaceae;g_Chloroflexus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%

k_Bacteria:p_Chloroflexi;c_Chloroflexi;o_Chloroflexales:f_Chloroflexaceae:g_Chloronema	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Chloroflexi;o_Chloroflexales:f_FFCH7168:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Chloroflexi;o_Chloroflexales:f_Oscillochloridaceae:g_Oscillochloris	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Chloroflexi;o_Chloroflexales:f_[Chlorothrixaceae]:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Chloroflexi;o_Herpetosiphonales:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Chloroflexi;o_[Roseiflexales];Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Chloroflexi;o_[Roseiflexales];f_:g_	0.1%	0.0%	0.0%	0.1%	0.0%	0.0%	0.2%
k_Bacteria:p_Chloroflexi;c_Chloroflexi;o_[Roseiflexales];f_[Kouleothrixaceae]:g_	0.1%	0.0%	0.1%	0.2%	0.0%	0.1%	0.3%
k_Bacteria:p_Chloroflexi;c_Chloroflexi;o_[Roseiflexales];f_[Roseiflexaceae]:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Dehalococcoidetes;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Dehalococcoidetes;o_Dehalococcoidales:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Dehalococcoidetes;o_Dehalococcoidales:f_Dehalococcoidaceae:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Dehalococcoidetes;o_Dehalococcoidales:f_Dehalococcoidaceae:g_Dehalogenimonas	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Ellin6529;o_:f_:g_	0.6%	0.9%	0.3%	0.6%	0.0%	0.3%	1.3%
k_Bacteria:p_Chloroflexi;c_Gitt-GS-136;o_:f_:g_	0.4%	1.5%	0.1%	0.1%	0.0%	0.4%	0.2%
k_Bacteria:p_Chloroflexi;c_Ktedonobacteria;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Ktedonobacteria;o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Ktedonobacteria;o_B10-SB3A:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Ktedonobacteria;o_B12-WMSP1:f_:g_	0.0%	0.0%	0.0%	0.0%	0.2%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Ktedonobacteria;o_Elev-1554:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Ktedonobacteria;o_JG30-KF-AS9:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Ktedonobacteria;o_Ktedonobacterales:f_Ktedonobacteraceae:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Ktedonobacteria;o_Thermogemmatisporales:f_Thermogemmatisporaceae:g_	2.5%	0.0%	0.0%	0.1%	15.1%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_MVP-21;o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_P2-1IE;o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%
k_Bacteria:p_Chloroflexi;c_S085;o_:f_:g_	0.8%	1.1%	0.6%	0.5%	0.1%	1.4%	0.9%
k_Bacteria:p_Chloroflexi;c_SAR202;o_:f_:g_	0.1%	0.2%	0.0%	0.0%	0.0%	0.1%	0.0%
k_Bacteria:p_Chloroflexi;c_SHA-26;o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_TK10;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_TK10;o_:f_:g_	0.0%	0.1%	0.1%	0.0%	0.0%	0.1%	0.0%
k_Bacteria:p_Chloroflexi;c_TK10;o_AKYG885;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_TK10;o_AKYG885:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_TK10;o_AKYG885:f_5B-12:g_	0.1%	0.0%	0.1%	0.0%	0.0%	0.2%	0.1%
k_Bacteria:p_Chloroflexi;c_TK10;o_AKYG885:f_Dolo_23:g_	0.0%	0.1%	0.0%	0.1%	0.0%	0.0%	0.1%
k_Bacteria:p_Chloroflexi;c_TK10;o_B07_WMSP1:f_:g_	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_TK10;o_B07_WMSP1:f_FFCH4570:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_TK17;o_:f_:g_	0.1%	0.2%	0.1%	0.0%	0.2%	0.1%	0.0%
k_Bacteria:p_Chloroflexi;c_TK17;o_mle1-48:f_:g_	0.2%	0.4%	0.4%	0.2%	0.0%	0.2%	0.0%
k_Bacteria:p_Chloroflexi;c_Thermomicrobia;o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Thermomicrobia;o_AKYG1722:f_:g_	0.1%	0.5%	0.1%	0.0%	0.0%	0.1%	0.0%
k_Bacteria:p_Chloroflexi;c_Thermomicrobia;o_Ellin6537:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_Thermomicrobia;o_JG30-KF-CM45:f_:g_	0.3%	0.7%	0.2%	0.2%	0.0%	0.2%	0.3%
k_Bacteria:p_Chloroflexi;c_Thermomicrobia;o_Sphaerobacterales:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Chloroflexi;c_[Thermobacula];o_[Thermobaculales];f_[Thermobaculaceae]:g_Thermobaculum	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;Other;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_:o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_4C0d-2;o_MLE1-12:f_:g_	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%	0.1%
k_Bacteria:p_Cyanobacteria;c_4C0d-2;o_SM1D11:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_4C0d-2;o_SM2F09:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_4C0d-2;o_YS2:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_Chloroplast;o_Chlorophyta:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_Chloroplast;o_Chlorophyta:f_Chlamydomonadaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_Chloroplast;o_Chlorophyta:f_Chlamydomonadaceae:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_Chloroplast;o_Chlorophyta:f_Trebouxiophyceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_Chloroplast;o_Chlorophyta:f_Trebouxiophyceae:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_Chloroplast;o_Euglenozoa:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_Chloroplast;o_Stramenopiles:f_:g_	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.2%
k_Bacteria:p_Cyanobacteria;c_Chloroplast;o_Streptophyta:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_ML635J-21;o_:f_:g_	0.1%	0.0%	0.1%	0.1%	0.0%	0.1%	0.1%
k_Bacteria:p_Cyanobacteria;c_Nostocophycideae;o_Nostocales:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_Nostocophycideae;o_Nostocales:f_Nostocaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_Nostocophycideae;o_Nostocales:f_Nostocaceae:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_Nostocophycideae;o_Nostocales:f_Nostocaceae:g_Anabaena	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_Nostocophycideae;o_Nostocales:f_Nostocaceae:g_Nodularia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_Nostocophycideae;o_Nostocales:f_Nostocaceae:g_Nostoc	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_Nostocophycideae;o_Nostocales:f_Scytonemataceae:g_Scytonema	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Cyanobacteria;c_Oscillatoriophycideae;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%

k_Bacteria;p_Cyanobacteria;c_Oscillatoriophycideae;o_Chroococcales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Cyanobacteria;c_Oscillatoriophycideae;o_Chroococcales;f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Cyanobacteria;c_Oscillatoriophycideae;o_Chroococcales;f_Chroococcaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Cyanobacteria;c_Oscillatoriophycideae;o_Chroococcales;f_Xenococcaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Cyanobacteria;c_Oscillatoriophycideae;o_Chroococcales;f_Xenococcaceae;g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Cyanobacteria;c_Oscillatoriophycideae;o_Oscillatoriales;f_Phormidiaceae;g_Microcoleus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Cyanobacteria;c_Oscillatoriophycideae;o_Oscillatoriales;f_Phormidiaceae;g_Phormidium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Cyanobacteria;c_Oscillatoriophycideae;o_Oscillatoriales;f_Phormidiaceae;g_Planktothrix	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Cyanobacteria;c_Synechococophycideae;o_Pseudanabaenales;f_Pseudanabaenaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Cyanobacteria;c_Synechococophycideae;o_Pseudanabaenales;f_Pseudanabaenaceae;g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p_Cyanobacteria;c_Synechococophycideae;o_Pseudanabaenales;f_Pseudanabaenaceae;g_Arthonema	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Cyanobacteria;c_Synechococophycideae;o_Pseudanabaenales;f_Pseudanabaenaceae;g_Halomicronema	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Cyanobacteria;c_Synechococophycideae;o_Pseudanabaenales;f_Pseudanabaenaceae;g_Leptolyngbya	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p_Cyanobacteria;c_Synechococophycideae;o_Pseudanabaenales;f_Pseudanabaenaceae;g_Pseudanabaena	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Cyanobacteria;c_Synechococophycideae;o_Synechococcales;f_Acaryochloridaceae;g_Acarochloris	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Elusimicrobia;c_o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Elusimicrobia;c_Elusimicrobia;o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Elusimicrobia;c_Elusimicrobia;o_Elusimicrobiales;f_g	0.1%	0.0%	0.1%	0.0%	0.2%	0.0%	0.0%
k_Bacteria;p_Elusimicrobia;c_Elusimicrobia;o_FAC88;f_g	0.2%	0.0%	0.2%	0.2%	0.4%	0.0%	0.2%
k_Bacteria;p_Elusimicrobia;c_Elusimicrobia;o_Iibf_g	0.1%	0.2%	0.2%	0.1%	0.0%	0.0%	0.1%
k_Bacteria;p_Elusimicrobia;c_Elusimicrobia;o_MVP-88;f_g	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Elusimicrobia;c_Endomicrobia;o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_FBP;c_o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_FCPU426;c_o_f_g	0.0%	0.0%	0.0%	0.0%	0.2%	0.0%	0.0%
k_Bacteria;p_Fibrobaeteres;c_o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Fibrobaeteres;c_B5-096;o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Fibrobaeteres;c_Fibrobaeteria;o_258ds10;f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Fibrobaeteres;c_Fibrobaeteria;o_Ucp1540;f_g	0.0%	0.0%	0.2%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Fibrobaeteres;c_SBZC_2415;o_f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Fibrobaeteres;c_TG3;o_TG3-1;f_g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Fibrobaeteres;c_TG3;o_TG3-1;f_TSCOR003-O20;g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;Other;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;Other;Other	0.3%	1.3%	0.0%	0.3%	0.1%	0.2%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_g	0.1%	0.1%	0.0%	0.0%	0.0%	0.3%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Alicyclobacillaceae;g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Alicyclobacillaceae;g_Alicyclobacillus	0.1%	0.0%	0.0%	0.1%	0.0%	0.0%	0.6%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Bacillaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Bacillaceae;g	0.1%	0.2%	0.0%	0.0%	0.0%	0.2%	0.1%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Bacillaceae;g_Anaerobacillus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Bacillaceae;g_Anoxybacillus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Bacillaceae;g_Bacillus	5.6%	4.9%	0.7%	1.4%	3.7%	21.4%	1.7%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Bacillaceae;g_Geobacillus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Bacillaceae;g_Natronobacillus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Bacillaceae;g_Oceanobacillus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Bacillaceae;g_Pontibacillus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Bacillaceae;g_Thalassobacillus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Bacillaceae;g_Virgibacillus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Listeriaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Paenibacillaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Paenibacillaceae;g	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Paenibacillaceae;g_Ammoniphilus	0.0%	0.0%	0.0%	0.0%	0.0%	0.2%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Paenibacillaceae;g_Aneurinibacillus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Paenibacillaceae;g_Brevibacillus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Paenibacillaceae;g_Cohnella	0.1%	0.1%	0.0%	0.1%	0.2%	0.1%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Paenibacillaceae;g_Paenibacillus	0.4%	0.4%	0.1%	0.5%	0.3%	0.9%	0.2%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Pasteuriaceae;g_Pasteuria	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Planococcaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Planococcaceae;g	0.2%	0.5%	0.0%	0.1%	0.0%	0.3%	0.1%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Planococcaceae;g_Lysinibacillus	0.0%	0.1%	0.0%	0.0%	0.0%	0.1%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Planococcaceae;g_Paenisporosarcina	0.1%	0.2%	0.0%	0.1%	0.0%	0.4%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Planococcaceae;g_Planococcus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Planococcaceae;g_Planomicrobium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Planococcaceae;g_Rummeliibacillus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Planococcaceae;g_Solibacillus	0.1%	0.1%	0.0%	0.0%	0.0%	0.3%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Planococcaceae;g_Sporosarcina	0.2%	0.4%	0.1%	0.3%	0.0%	0.2%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales;f_Planococcaceae;g_Ureibacillus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%

k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales:f_Sporolactobacillaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales:f_Sporolactobacillaceae:g_Sporolactobacillus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales:f_Staphylococcaceae:g_Staphylococcus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales:f_Thermoactinomycetaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales:f_Thermoactinomycetaceae:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales:f_Thermoactinomycetaceae:g_Planifilum	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales:f_Thermoactinomycetaceae:g_Shimazuella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales:f_[Exiguobacteraceae];g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales:f_[Exiguobacteraceae];g_Exiguobacterium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales:f_[Thermicanaceae];g_Thermicanus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Haloplasmatales:f_Haloplasmataceae:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Lactobacillales:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Lactobacillales:f_Aerococcaceae:g_Facklamia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Lactobacillales:f_Carnobacteriaceae:g_Carnobacterium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Lactobacillales:f_Carnobacteriaceae:g_Deszenvia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Lactobacillales:f_Enterococcaceae:g_Enterococcus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Lactobacillales:f_Lactobacillaceae:g_Lactobacillus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Lactobacillales:f_Leuconostocaceae:g_Leuconostoc	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Bacilli;o_Turicibacterales:f_Turicibacteraceae:g_Turicibacter	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Caldicoprobacteraceae:g_Caldicoprobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Christensenellaceae:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Clostridiaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Clostridiaceae:g_	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Clostridiaceae:g_Alkaliphilus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Clostridiaceae:g_Caloramator	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Clostridiaceae:g_Candidatus Arthromitus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Clostridiaceae:g_Clostridium	0.1%	0.1%	0.1%	0.1%	0.0%	0.1%	0.3%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Clostridiaceae:g_Geoposporobacter Thermotalea	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Clostridiaceae:g_Oxobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Clostridiaceae:g_SMB53	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Clostridiaceae:g_Sarcina	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Clostridiaceae:g_Thermoanaerobacterium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Dehalobacteriaceae:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Dehalobacteriaceae:g_Dehalobacterium	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_EtOH8:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Eubacteriaceae:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Gracilibacteraceae:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Gracilibacteraceae:g_Gracilibacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Gracilibacteraceae:g_Lutispora	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Lachnospiraceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Lachnospiraceae:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Lachnospiraceae:g_Anaerostipes	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Lachnospiraceae:g_Blautia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Lachnospiraceae:g_Coproccoccus	0.1%	0.0%	0.0%	0.1%	0.0%	0.5%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Lachnospiraceae:g_Dorea	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Lachnospiraceae:g_Epulopiscium	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Lachnospiraceae:g_Lachnospira	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Lachnospiraceae:g_Roseburia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Peptococcaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Peptococcaceae:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Peptococcaceae:g_Dehalobacter Syntrophobotulus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Peptococcaceae:g_Desulfonipora	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Peptococcaceae:g_Desulfosporosinus	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Peptococcaceae:g_Desulfotomaculum	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Peptococcaceae:g_Desulfurispora	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Peptococcaceae:g_Pelotomaculum	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Peptococcaceae:g_Sporotomaculum	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Peptococcaceae:g_WCHB1-84	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Peptostreptococcaceae;Other	0.0%	0.1%	0.0%	0.1%	0.0%	0.0%	0.1%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Peptostreptococcaceae:g_	0.1%	0.1%	0.0%	0.1%	0.0%	0.1%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Peptostreptococcaceae:g_Clostridium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Peptostreptococcaceae:g_Tepidibacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Ruminococcaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales:f_Ruminococcaceae:g_	0.1%	0.1%	0.0%	0.1%	0.0%	0.1%	0.3%

k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_Ruminococcaceae;g_Anaerotruncus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_Ruminococcaceae;g_Ethanoligenens	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_Ruminococcaceae;g_Oscillospira	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_Ruminococcaceae;g_Ruminococcus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_Symbiobacteriaceae;g_Symbiobacterium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_Syntrophomonadaceae;g_Syntrophomonas	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_Veillonellaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_Veillonellaceae;g_BSV43	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_Veillonellaceae;g_G07	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_Veillonellaceae;g_Pelosinus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_Veillonellaceae;g_Sporomusa	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_[Acidaminobacteraceae];g_Acidaminobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_[Acidaminobacteraceae];g_Fusibacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_[Mogibacteriaceae];Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_[Mogibacteriaceae];g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_[Mogibacteriaceae];g_Anaerovorax	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_[Tissierellaceae];g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_[Tissierellaceae];g_Sedimentibacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_[Tissierellaceae];g_Sporanaerobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_[Tissierellaceae];g_Tepidimicrobium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales;f_[Tissierellaceae];g_Tissierella Soehngenia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Halanaerobiales;f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_OPB54;f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p_Firmicutes;c_Clostridia;o_SHA-98;f_D2;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Thermoanaerobacterales;f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Clostridia;o_Thermoanaerobacterales;f_Thermacetogenium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Erysipelotrichi;o_Erysipelotrichales;f_Erysipelotrichaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Firmicutes;c_Erysipelotrichi;o_Erysipelotrichales;f_Erysipelotrichaceae;g_PSB-M-3	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_GAL15;c_o_f_g_	0.4%	0.0%	0.2%	0.0%	1.5%	0.7%	0.0%
k_Bacteria;p_GN02;c_o_f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_GN02;c_3BR-5F;o_f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_GN02;c_BB34;o_f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_GN02;c_BD1-5;o_f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_GN02;c_GKS2-174;o_f_g_	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_GN02;c_GN07;o_f_g_	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%
k_Bacteria;p_GN04;c_o_f_g_	0.0%	0.0%	0.2%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_GN04;c_GN15;o_f_g_	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_GN04;c_MSB-5A5;o_f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Gemmatimonadetes;c_o_f_g_	0.3%	0.0%	0.7%	0.0%	0.0%	0.7%	0.2%
k_Bacteria;p_Gemmatimonadetes;c_Gemm-1;o_f_g_	0.7%	0.7%	0.8%	0.6%	0.2%	1.6%	0.5%
k_Bacteria;p_Gemmatimonadetes;c_Gemm-2;o_f_g_	0.1%	0.2%	0.1%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p_Gemmatimonadetes;c_Gemm-3;o_f_g_	0.1%	0.5%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Gemmatimonadetes;c_Gemm-4;o_f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Gemmatimonadetes;c_Gemm-5;o_f_g_	0.2%	0.4%	0.3%	0.0%	0.0%	0.1%	0.3%
k_Bacteria;p_Gemmatimonadetes;c_Gemmatimonadetes;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Gemmatimonadetes;c_Gemmatimonadetes;o_f_g_	0.3%	0.1%	1.0%	0.1%	0.1%	0.6%	0.2%
k_Bacteria;p_Gemmatimonadetes;c_Gemmatimonadetes;o_C114;f_g_	0.2%	0.0%	0.5%	0.0%	0.0%	0.6%	0.0%
k_Bacteria;p_Gemmatimonadetes;c_Gemmatimonadetes;o_Ellin5290;f_g_	0.0%	0.1%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p_Gemmatimonadetes;c_Gemmatimonadetes;o_Gemmatimonadales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Gemmatimonadetes;c_Gemmatimonadetes;o_Gemmatimonadales;f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Gemmatimonadetes;c_Gemmatimonadetes;o_Gemmatimonadales;f_Ellin5301;g_	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p_Gemmatimonadetes;c_Gemmatimonadetes;o_Gemmatimonadales;f_Gemmatimonadaceae;g_Gemmatimonas	0.1%	0.0%	0.3%	0.1%	0.0%	0.1%	0.0%
k_Bacteria;p_Gemmatimonadetes;c_Gemmatimonadetes;o_KD8-87;f_g_	0.0%	0.1%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Gemmatimonadetes;c_Gemmatimonadetes;o_N1423WL;f_g_	0.3%	0.0%	1.8%	0.0%	0.0%	0.3%	0.0%
k_Bacteria;p_Kazan-3B-28;c_o_f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Lentisphaerae;c_[Lentisphaeria];o_Lentisphaerales;f_Arctic95B-10;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Lentisphaerae;c_[Lentisphaeria];o_Victivallales;f_Victivallaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_MVP-21;c_o_f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_NC10;c_12-24;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_NC10;c_12-24;o_JH-WHS47;f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p_NC10;c_12-24;o_Methylomirabiales;f_Methylomirabiliaceae;g_Candidatus Methylomirabilis	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_NC10;c_wb1-A12;o_f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_NKB19;Other;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_NKB19;c_o_f_g_	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_NKB19;c_TSBW08;o_f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Nitrospirae;c_Nitrospira;o_Nitrospirales;f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p_Nitrospirae;c_Nitrospira;o_Nitrospirales;f_0319-6A21;g_	0.6%	0.1%	0.5%	0.7%	0.3%	1.1%	1.1%

k_Bacteria;p__Nitrospirae;c__Nitrospira;o__Nitrospirales;f__FW;g_4-29	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.2%
k_Bacteria;p__Nitrospirae;c__Nitrospira;o__Nitrospirales;f__Nitrospiraceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Nitrospirae;c__Nitrospira;o__Nitrospirales;f__Nitrospiraceae;g__	1.6%	0.0%	3.1%	0.1%	4.6%	2.0%	0.0%
k_Bacteria;p__Nitrospirae;c__Nitrospira;o__Nitrospirales;f__Nitrospiraceae;g_JG37-AG-70	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Nitrospirae;c__Nitrospira;o__Nitrospirales;f__Nitrospiraceae;g_Nitrospira	3.6%	0.8%	6.0%	1.5%	9.5%	2.0%	2.0%
k_Bacteria;p__Nitrospirae;c__Nitrospira;o__Nitrospirales;f__(Leptospirillaceae);g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Nitrospirae;c__Nitrospira;o__Nitrospirales;f__(Thermodesulfobionaceae);g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p__Nitrospirae;c__Nitrospira;o__Nitrospirales;f__(Thermodesulfobionaceae);g_GOUTA19	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p__Nitrospirae;c__Nitrospira;o__Nitrospirales;f__(Thermodesulfobionaceae);g_HB118	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__OD1;c__o__f__g__	0.0%	0.1%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__OD1;c__ABY1;o__f__g__	0.1%	0.0%	0.0%	0.0%	0.2%	0.1%	0.0%
k_Bacteria;p__OD1;c__Mb-NB09;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__OD1;c__SM2F11;o__f__g__	0.2%	0.1%	0.2%	0.1%	0.2%	0.1%	0.2%
k_Bacteria;p__OD1;c__ZB2;o__f__g__	0.1%	0.1%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__OP1;c__[Acetothermia];o__[Acetothermales];f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__OP11;c__o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__OP11;c__OP11-3;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__OP11;c__OP11-4;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%
k_Bacteria;p__OP11;c__WCHB1-64;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__OP11;c__WCHB1-64;o__K2-4-19;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__OP11;c__WCHB1-64;o__d153;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__OP3;c__o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__OP3;c__PBS-25;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__OP3;c__kol11;o__f__g__	0.0%	0.0%	0.1%	0.0%	0.0%	0.1%	0.0%
k_Bacteria;p__OP3;c__kol11;o__GIF10;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__OP3;c__kol11;o__GIF10;f__kpj58rc;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__028H05-P-BN-P5;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__BD7-11;o__f__g__	0.0%	0.0%	0.1%	0.0%	0.0%	0.1%	0.0%
k_Bacteria;p__Planctomycetes;c__C6;o__MVS-107;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__C6;o__Ucm1571;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__C6;o__d113;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__ODP123;o__T8-B82;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__OM190;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__OM190;o__CL500-15;f__g__	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__OM190;o__agg27;f__g__	0.1%	0.1%	0.1%	0.1%	0.0%	0.1%	0.1%
k_Bacteria;p__Planctomycetes;c__Phycisphaerae;o__AKAU3564;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__Phycisphaerae;o__CCM11a;f__g__	0.0%	0.0%	0.2%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__Phycisphaerae;o__Phycisphaerales;f__g__	0.3%	0.4%	0.3%	0.4%	0.0%	0.3%	0.2%
k_Bacteria;p__Planctomycetes;c__Phycisphaerae;o__Phycisphaerales;f__Phycisphaeraeae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__Phycisphaerae;o__Pla1;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__Phycisphaerae;o__S-70;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__Phycisphaerae;o__WD2101;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__Phycisphaerae;o__mle1-8;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__Pla3;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%
k_Bacteria;p__Planctomycetes;c__Pla4;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__Planctomycetia;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__Planctomycetia;o__B97;f__g__	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__Planctomycetia;o__Gemmatales;f__Gemmataceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__Planctomycetia;o__Gemmatales;f__Gemmataceae;g__	1.0%	0.2%	0.6%	0.9%	2.8%	0.7%	0.8%
k_Bacteria;p__Planctomycetes;c__Planctomycetia;o__Gemmatales;f__Gemmataceae;g_Gemmata	0.1%	0.1%	0.1%	0.2%	0.0%	0.0%	0.1%
k_Bacteria;p__Planctomycetes;c__Planctomycetia;o__Gemmatales;f__Isosphaeraeae;g__	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__Planctomycetia;o__Gemmatales;f__Isosphaeraeae;g_Isosphaera	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__Planctomycetia;o__Pirellulales;f__Pirellulaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__Planctomycetia;o__Pirellulales;f__Pirellulaceae;g__	0.1%	0.2%	0.1%	0.2%	0.0%	0.1%	0.2%
k_Bacteria;p__Planctomycetes;c__Planctomycetia;o__Pirellulales;f__Pirellulaceae;g_A17	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__Planctomycetia;o__Pirellulales;f__Pirellulaceae;g_Pirellula	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.1%
k_Bacteria;p__Planctomycetes;c__Planctomycetia;o__Planctomycetiales;f__Planctomycetaceae;g_Planctomyces	0.0%	0.1%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__[Brocadia];o__Brocadiales;f__Brocadiaaceae;g_Candidatus Brocadia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__vadinHA49;o__DH61;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__vadinHA49;o__PHOS-HE93;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Planctomycetes;c__vadinHA49;o__p04_C01;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;Other;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Alphaproteobacteria;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__f__g__	0.4%	1.1%	0.5%	0.4%	0.0%	0.1%	0.0%
k_Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__BD7-3;f__g__	0.1%	0.2%	0.1%	0.1%	0.0%	0.0%	0.0%

k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales;f_g_	0.3%	1.3%	0.0%	0.1%	0.0%	0.2%	0.2%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales;f_Acetobacteraceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales;f_Acetobacteraceae;g_Roseococcus	0.1%	0.0%	0.0%	0.3%	0.4%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales;f_Acetobacteraceae;g_Roseococcus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales;f_Rhodospirillaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales;f_Rhodospirillaceae;g_	2.2%	3.1%	0.7%	2.5%	4.0%	0.8%	2.3%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales;f_Rhodospirillaceae;g_Azospirillum	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales;f_Rhodospirillaceae;g_Inquilinus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales;f_Rhodospirillaceae;g_Magnetospirillum	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales;f_Rhodospirillaceae;g_Phaeospirillum	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales;f_Rhodospirillaceae;g_Rhodovibrio	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales;f_Rhodospirillaceae;g_Skermanella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales;f_Rhodospirillaceae;g_Telmatospirillum	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rickettsiales;f_g_	0.1%	0.0%	0.1%	0.1%	0.1%	0.0%	0.1%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rickettsiales;f_Holosporaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rickettsiales;f_Rickettsiaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rickettsiales;f_Rickettsiaceae;g_Orientia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rickettsiales;f_Rickettsiaceae;g_Rickettsia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rickettsiales;f_mitochondria;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rickettsiales;f_mitochondria;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales;f_g_	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales;f_Erythrobacteraceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales;f_Erythrobacteraceae;g_	0.1%	0.2%	0.2%	0.2%	0.0%	0.0%	0.1%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales;f_Erythrobacteraceae;g_Lutibacterium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales;f_Sphingomonadaceae;Other	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales;f_Sphingomonadaceae;g_	0.4%	0.7%	0.5%	0.5%	0.0%	0.3%	0.1%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales;f_Sphingomonadaceae;g_Blastomonas	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales;f_Sphingomonadaceae;g_Kaistobacter	1.4%	0.9%	5.8%	0.8%	0.0%	0.6%	0.5%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales;f_Sphingomonadaceae;g_Novosphingobium	0.1%	0.0%	0.2%	0.2%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales;f_Sphingomonadaceae;g_Sphingobium	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales;f_Sphingomonadaceae;g_Sphingomonas	0.1%	0.1%	0.4%	0.2%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales;f_Sphingomonadaceae;g_Sphingopyxis	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.2%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_f_g_	1.4%	1.2%	1.3%	1.8%	0.0%	1.8%	2.3%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_A21b;f_EB1003;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_A21b;f_UD5;g_	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Alcaligenaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Alcaligenaceae;g_	0.2%	0.1%	0.4%	0.1%	0.0%	0.1%	0.6%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Alcaligenaceae;g_Achromobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Alcaligenaceae;g_Tetrathibacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Burkholderiaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Burkholderiaceae;g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Burkholderiaceae;g_Burkholderia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Burkholderiaceae;g_Lautropia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Burkholderiaceae;g_Pandoraea	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;Other	0.1%	0.1%	0.1%	0.2%	0.0%	0.1%	0.1%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_	0.3%	0.2%	0.5%	0.5%	0.1%	0.1%	0.1%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_Acidovorax	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_Azohydromonas	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_Comamonas	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_Delftia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_Hydrogenophaga	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_Hylemonella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_Leptothrix	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_Limnobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_Limnohabitans	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_Methylibium	0.0%	0.0%	0.1%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_Polaromonas	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_Ramlibacter	0.1%	0.1%	0.2%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_Rhodoferax	0.1%	0.0%	0.1%	0.0%	0.1%	0.1%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_Roseateae	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_Rubrivivax	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales;f_Comamonadaceae;g_Simplicispira	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%

k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Burkholderiales;f__Comamonadaceae;g__Tepidimonas	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Burkholderiales;f__Comamonadaceae;g__Variovorax	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Burkholderiales;f__Oxalobacteraceae;Other	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Burkholderiales;f__Oxalobacteraceae;g__	0.4%	0.1%	0.5%	0.6%	0.6%	0.3%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Burkholderiales;f__Oxalobacteraceae;g__Cupriavidus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Burkholderiales;f__Oxalobacteraceae;g__Janthinobacterium	0.2%	0.0%	0.4%	0.4%	0.0%	0.1%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Burkholderiales;f__Oxalobacteraceae;g__Oxalobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Burkholderiales;f__Oxalobacteraceae;g__Ralstonia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Ellin6067;f__g__	0.3%	0.2%	0.4%	0.5%	0.0%	0.2%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Gallionellales;f__Gallionellaceae;g__Gallionella	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Hydrogenophilales;f__Hydrogenophilaceae;g__Thiobacillus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__IS-44;f__g__	0.1%	0.2%	0.1%	0.4%	0.0%	0.1%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__MND1;f__g__	0.6%	0.4%	0.7%	0.3%	0.0%	0.5%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Methylophilales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Methylophilales;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Methylophilales;f__Methylophilaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Methylophilales;f__Methylophilaceae;g__	0.1%	0.0%	0.1%	0.0%	0.2%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Methylophilales;f__Methylophilaceae;g__Methylothera	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Neisseriales;f__Neisseriaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Neisseriales;f__Neisseriaceae;g__	0.0%	0.0%	0.0%	0.0%	0.2%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Neisseriales;f__Neisseriaceae;g__Deefgea	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Neisseriales;f__Neisseriaceae;g__Vogesella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Nitrosomonadales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Nitrosomonadales;f__g__	0.3%	0.0%	0.1%	0.2%	1.9%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Nitrosomonadales;f__Nitrosomonadaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Nitrosomonadales;f__Nitrosomonadaceae;g__	0.5%	0.0%	0.5%	0.1%	2.0%	0.4%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Nitrosomonadales;f__Nitrosomonadaceae;g__Nitrosovibrio	0.2%	0.0%	0.1%	0.0%	0.9%	0.1%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Procabacteriales;f__Procabacteriaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Procabacteriales;f__Procabacteriaceae;g__Procabacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Rhodocyclales;f__Rhodocyclaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Rhodocyclales;f__Rhodocyclaceae;g__	0.0%	0.0%	0.0%	0.2%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Rhodocyclales;f__Rhodocyclaceae;g__Azoarcus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Rhodocyclales;f__Rhodocyclaceae;g__Azospira	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Rhodocyclales;f__Rhodocyclaceae;g__Dechloromonas	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Rhodocyclales;f__Rhodocyclaceae;g__Dok59	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Rhodocyclales;f__Rhodocyclaceae;g__Propionivibrio	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Rhodocyclales;f__Rhodocyclaceae;g__Rhodocycilus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Rhodocyclales;f__Rhodocyclaceae;g__Sulfuritalea	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Rhodocyclales;f__Rhodocyclaceae;g__Thauera	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Rhodocyclales;f__Rhodocyclaceae;g__Uliginosibacterium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__SBla14;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__SC-I-84;f__g__	0.3%	0.1%	0.5%	0.4%	0.1%	0.2%
k_Bacteria:p__Proteobacteria;c__Betaproteobacteria;o__Thiobacteriales;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__f__g__	0.1%	0.0%	0.1%	0.1%	0.0%	0.1%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__BPC076;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Bdellovibrionales;f__Bacteriovoracaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Bdellovibrionales;f__Bacteriovoracaceae;g__	0.1%	0.1%	0.1%	0.2%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Bdellovibrionales;f__Bacteriovoracaceae;g__Bacteriovorax	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Bdellovibrionales;f__Bacteriovoracaceae;g__Peredibacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Bdellovibrionales;f__Bdellovibrionaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Bdellovibrionales;f__Bdellovibrionaceae;g__Bdellovibrio	0.2%	0.3%	0.2%	0.4%	0.0%	0.1%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Desulfarculales;f__Desulfarculaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Desulfobacteriales;f__Desulfobacteraceae;g__Desulfococcus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Desulfobacteriales;f__Desulfobulbaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Desulfobacteriales;f__Desulfobulbaceae;g__Desulfobulbus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Desulfovibrionales;f__Desulfovibrionaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Desulfovibrionales;f__Desulfovibrionaceae;g__Desulfovibrio	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Desulfuromonadales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Desulfuromonadales;f__Desulfuromonadaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Desulfuromonadales;f__Geobacteraceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Desulfuromonadales;f__Geobacteraceae;g__Geobacter	0.1%	0.0%	0.0%	0.1%	0.0%	0.6%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Desulfuromonadales;f__Pelobacteraceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Desulfuromonadales;f__Pelobacteraceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__Desulfuromonadales;f__Pelobacteraceae;g__Pelobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__FAC87;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p__Proteobacteria;c__Deltaproteobacteria;o__GMD14H09;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%

k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__MBNT15;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__MIZ46;f__g__	0.1%	0.0%	0.1%	0.2%	0.0%	0.1%	0.1%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;Other:Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;f__g__	1.0%	2.1%	0.7%	1.3%	0.3%	0.5%	1.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;f__0319-6G20;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;f__Cystobacterineae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;f__Haliangiaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;f__Haliangiaceae;g__	0.2%	0.3%	0.2%	0.1%	0.0%	0.1%	0.2%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;f__Haliangiaceae;g__Haliangium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;f__Myxococcaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;f__Myxococcaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;f__Myxococcaceae;g__Anaeromyxobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;f__Myxococcaceae;g__Myxococcus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;f__Nannocystaceae;g__Nannocystis	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;f__Nannocystaceae;g__Plesiocystis	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;f__OM27;g__	0.1%	0.2%	0.1%	0.1%	0.0%	0.1%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;f__Polyangiaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Myxococcales;f__Polyangiaceae;g__Chondromyces	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__NB1-j;f__g__	0.1%	0.0%	0.1%	0.1%	0.0%	0.1%	0.1%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__NB1-j;f__JTB38;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__NB1-j;f__MND4;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__NB1-j;f__NB1-i;g__	0.1%	0.0%	0.0%	0.1%	0.0%	0.1%	0.1%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__NB1-j;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__PB19;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Spirobacillales;f__g__	0.1%	0.1%	0.1%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Sva0853;f__JTB36;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Syntrophobacterales;f__Syntrophaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Syntrophobacterales;f__Syntrophaceae;g__Desulfobacca	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Syntrophobacterales;f__Syntrophaceae;g__Smithella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Syntrophobacterales;f__Syntrophobacteraceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Syntrophobacterales;f__Syntrophobacteraceae;g__	1.5%	1.1%	1.0%	1.4%	1.4%	1.2%	2.9%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Syntrophobacterales;f__Syntrophobacteraceae;g__Syntrophobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__Syntrophobacterales;f__Syntrophorhabdaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__[Entotheonellales];f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__[Entotheonellales];f__[Entotheonellaceae];g__	0.3%	0.1%	0.1%	0.3%	0.0%	0.1%	1.0%
k_Bacteria;p__Proteobacteria;c__Deltaproteobacteria;o__[Entotheonellales];f__[Entotheonellaceae];g__Candidatus Entotheonella	0.1%	0.4%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Epsilonproteobacteria;o__Campylobacteriales;f__Helicobacteraceae;g__Sulfuricurvum	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;Other:Other:Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__f__g__	0.1%	0.0%	0.0%	0.1%	0.3%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__34P16;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Acidithiobacillales;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Aeromonadales;f__Aeromonadaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Alteromonadales;Other:Other	0.0%	0.1%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Alteromonadales;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Alteromonadales;f__125ds10;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Alteromonadales;f__211ds20;g__	0.0%	0.2%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Alteromonadales;f__Alteromonadaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Alteromonadales;f__Alteromonadaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Alteromonadales;f__Alteromonadaceae;g__Cellvibrio	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Alteromonadales;f__Alteromonadaceae;g__Marinobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Alteromonadales;f__HTCC2188;g__HTCC	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Alteromonadales;f__OM60;g__	0.0%	0.1%	0.0%	0.2%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Alteromonadales;f__[Chromatiaceae];g__Rheinheimera	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Chromatiales;Other:Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Chromatiales;f__g__	1.0%	0.1%	0.2%	0.0%	0.0%	5.7%	0.2%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Chromatiales;f__Chromatiaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Chromatiales;f__Chromatiaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Chromatiales;f__Chromatiaceae;g__Nitrosococcus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Chromatiales;f__Ectothiorhodospiraceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Chromatiales;f__Ectothiorhodospiraceae;g__Halorhodospira	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Chromatiales;f__Ectothiorhodospiraceae;g__Natronocella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Enterobacteriales;f__Enterobacteriaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Enterobacteriales;f__Enterobacteriaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Enterobacteriales;f__Enterobacteriaceae;g__Providencia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__HOC36;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__HTCC2188;f__HTCC2089;g__	0.1%	0.1%	0.0%	0.3%	0.0%	0.0%	0.1%

k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Legionellales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Legionellales;f__g__	0.1%	0.0%	0.1%	0.1%	0.1%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Legionellales;f__Coxiellaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Legionellales;f__Coxiellaceae;g__	0.2%	0.2%	0.1%	0.3%	0.4%	0.2%	0.1%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Legionellales;f__Coxiellaceae;g__Aquicella	0.1%	0.0%	0.0%	0.1%	0.5%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Legionellales;f__Coxiellaceae;g__Coxiella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Legionellales;f__Coxiellaceae;g__Rickettsiella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Legionellales;f__Legionellaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Legionellales;f__Legionellaceae;g__	0.0%	0.1%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Legionellales;f__Legionellaceae;g__Legionella	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Legionellales;f__Legionellaceae;g__Tatlockia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Methylococcales;f__Crenotrichaceae;g__Crenothrix	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Methylococcales;f__Methylococcaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Methylococcales;f__Methylococcaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Methylococcales;f__Methylococcaceae;g__Methylocaldum	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Methylococcales;f__Methylococcaceae;g__Methylomonas	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Methylococcales;f__Methylococcaceae;g__Methylosarcina	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Oceanospirillales;f__Halomonadaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Oceanospirillales;f__Halomonadaceae;g__Halomonas	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Oceanospirillales;f__Oceanospirillaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Oceanospirillales;f__Oleiphilaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__PYR10d3;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Pseudomonadales;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Pseudomonadales;f__Moraxellaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Pseudomonadales;f__Moraxellaceae;g__	0.1%	0.0%	0.0%	0.0%	0.6%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Pseudomonadales;f__Moraxellaceae;g__Acinetobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Pseudomonadales;f__Moraxellaceae;g__Perluclidibaca	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Pseudomonadales;f__Pseudomonadaceae;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Pseudomonadales;f__Pseudomonadaceae;g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Pseudomonadales;f__Pseudomonadaceae;g__Pseudomonas	0.1%	0.2%	0.1%	0.2%	0.3%	0.1%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Thiotrichales;f__Piscirickettsiaceae;g__	0.6%	2.3%	0.2%	0.6%	0.0%	0.1%	0.4%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Xanthomonadales;f__Sinobacteraceae;g__	1.2%	1.6%	0.6%	1.5%	1.4%	1.1%	0.8%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Xanthomonadales;f__Sinobacteraceae;g__Nevskia	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Xanthomonadales;f__Sinobacteraceae;g__Steroidobacter	0.3%	0.8%	0.4%	0.3%	0.0%	0.1%	0.4%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Xanthomonadales;f__Xanthomonadaceae;Other	0.1%	0.1%	0.6%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Xanthomonadales;f__Xanthomonadaceae;g__	0.3%	0.8%	0.5%	0.5%	0.2%	0.1%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Xanthomonadales;f__Xanthomonadaceae;g__Arenimonas	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Xanthomonadales;f__Xanthomonadaceae;g__Dokdonella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Xanthomonadales;f__Xanthomonadaceae;g__Luteibacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Xanthomonadales;f__Xanthomonadaceae;g__Luteimonas	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Xanthomonadales;f__Xanthomonadaceae;g__Lysobacter	0.1%	0.1%	0.3%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Xanthomonadales;f__Xanthomonadaceae;g__Pseudoxanthomonas	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Xanthomonadales;f__Xanthomonadaceae;g__Rhodanobacter	0.1%	0.0%	0.0%	0.1%	0.5%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Xanthomonadales;f__Xanthomonadaceae;g__Stenotrophomonas	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Xanthomonadales;f__Xanthomonadaceae;g__Thermomonas	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__[Marinicellales];f__[Marinicellaceae];g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__TA18;o__CV90;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Proteobacteria;c__TA18;o__PHOS-HD29;f__g__	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria;p__SBR1093;c__o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.2%	0.0%
k_Bacteria;p__SBR1093;c__VHS-B5-50;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__SC4;c__o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__SR1;c__o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Spirochaetes;c__GN05;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Spirochaetes;c__[Leptospirae];o__[Leptospirales];f__Leptospiraceae;g__Leptonema	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Spirochaetes;c__[Leptospirae];o__[Leptospirales];f__Leptospiraceae;g__Leptospira	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__Spirochaetes;c__[Leptospirae];o__[Leptospirales];f__Leptospiraceae;g__Turneriella	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__TM6;c__SBRH58;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__TM6;c__SJA-4;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__TM6;c__SJA-4;o__f__g__	0.1%	0.2%	0.1%	0.1%	0.1%	0.1%	0.2%
k_Bacteria;p__TM6;c__SJA-4;o__S1198;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__TM6;c__SJA-4;o__YIF2-48;f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__TM7;Other;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__TM7;c__o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__TM7;c__MJK10;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__TM7;c__SC3;o__f__g__	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria;p__TM7;c__TM7-1;o__f__g__	0.4%	0.5%	0.4%	0.5%	0.1%	0.3%	0.4%
k_Bacteria;p__TM7;c__TM7-3;Other;Other;Other	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%

k_Bacteria:p_TM7;c_TM7-3;o_:f_:g_	0.1%	0.1%	0.2%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_TM7;c_TM7-3;o_Blg118:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_TM7;c_TM7-3;o_CW040:f_F16:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_TM7;c_TM7-3;o_EW055:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_TM7;c_TM7-3;o_I025:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Tenericutes;c_Mollicutes;o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Tenericutes;c_Mollicutes;o_Anaeroplasmatales:f_Anaeroplasmataceae:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Tenericutes;c_Mollicutes;o_Anaeroplasmatales:f_Anaeroplasmataceae:g_Asteroleplasma	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_:o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_Opitutae;o_:f_:g_	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_Opitutae;o_Opitutales:f_Opitutaceae:g_	0.0%	0.1%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_Opitutae;o_Opitutales:f_Opitutaceae:g_Opitutus	0.1%	0.2%	0.1%	0.1%	0.0%	0.0%	0.1%
k_Bacteria:p_Verrucomicrobia;c_Opitutae;o_[Cericococcales]:f_[Cericococcaceae]:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_Verruco-5;o_LD1-PB3:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_Verruco-5;o_WCHB1-41:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_Verrucomicrobiae;o_Verrucomicrobiales:f_Verrucomicrobiaceae:g_	0.1%	0.1%	0.0%	0.2%	0.0%	0.0%	0.1%
k_Bacteria:p_Verrucomicrobia;c_Verrucomicrobiae;o_Verrucomicrobiales:f_Verrucomicrobiaceae:g_Luteolibacter	0.0%	0.1%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_Verrucomicrobiae;o_Verrucomicrobiales:f_Verrucomicrobiaceae:g_Prosthecobacter	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_Verrucomicrobiae;o_Verrucomicrobiales:f_Verrucomicrobiaceae:g_Verrucomicrobium	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_[Methylacidiphilae];o_Methylacidiphilales:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_[Methylacidiphilae];o_Methylacidiphilales:f_LD19:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_[Methylacidiphilae];o_S-BQ2-57:f_:g_	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_[Pedosphaerae];o_[Pedosphaerales];Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_[Pedosphaerae];o_[Pedosphaerales]:f_:g_	0.3%	0.2%	0.1%	0.5%	0.0%	0.1%	0.6%
k_Bacteria:p_Verrucomicrobia;c_[Pedosphaerae];o_[Pedosphaerales]:f_Ellin515:g_	0.1%	0.0%	0.0%	0.2%	0.0%	0.0%	0.4%
k_Bacteria:p_Verrucomicrobia;c_[Pedosphaerae];o_[Pedosphaerales]:f_Ellin517:g_	0.1%	0.2%	0.4%	0.1%	0.0%	0.0%	0.1%
k_Bacteria:p_Verrucomicrobia;c_[Pedosphaerae];o_[Pedosphaerales]:f_OPB35:g_	0.1%	0.0%	0.2%	0.0%	0.0%	0.1%	0.1%
k_Bacteria:p_Verrucomicrobia;c_[Pedosphaerae];o_[Pedosphaerales]:f_R4-41B:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_[Pedosphaerae];o_[Pedosphaerales]:f_[Pedosphaeraceae];Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_[Pedosphaerae];o_[Pedosphaerales]:f_[Pedosphaeraceae]:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_[Pedosphaerae];o_[Pedosphaerales]:f_[Pedosphaeraceae]:g_Pedosphaera	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_[Pedosphaerae];o_[Pedosphaerales]:f_auto67_4W:g_	0.1%	0.1%	0.0%	0.3%	0.0%	0.0%	0.1%
k_Bacteria:p_Verrucomicrobia;c_[Spartobacteria];o_[Chthoniobacteriales]:f_01D2Z36:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_[Spartobacteria];o_[Chthoniobacteriales]:f_[Chthoniobacteraceae];Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_[Spartobacteria];o_[Chthoniobacteriales]:f_[Chthoniobacteraceae]:g_	0.1%	0.1%	0.1%	0.2%	0.0%	0.0%	0.2%
k_Bacteria:p_Verrucomicrobia;c_[Spartobacteria];o_[Chthoniobacteriales]:f_[Chthoniobacteraceae]:g_Candidatus Xiphinema tobacter	0.4%	0.1%	0.0%	0.3%	0.5%	0.2%	1.1%
k_Bacteria:p_Verrucomicrobia;c_[Spartobacteria];o_[Chthoniobacteriales]:f_[Chthoniobacteraceae]:g_Chthoniobacter	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_[Spartobacteria];o_[Chthoniobacteriales]:f_[Chthoniobacteraceae]:g_DA101	0.4%	0.0%	0.0%	1.0%	0.1%	0.0%	1.0%
k_Bacteria:p_Verrucomicrobia;c_[Spartobacteria];o_[Chthoniobacteriales]:f_[Chthoniobacteraceae]:g_Ellin506	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_[Spartobacteria];o_[Chthoniobacteriales]:f_[Chthoniobacteraceae]:g_OR-59	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_Verrucomicrobia;c_[Spartobacteria];o_[Chthoniobacteriales]:f_[Chthoniobacteraceae]:g_heteroC45_4W	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_WPS-2;c_:o_:f_:g_	0.3%	0.0%	0.0%	0.0%	1.6%	0.0%	0.0%
k_Bacteria:p_WS2;c_Kazan-3B-09;o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_WS2;c_SHA-109;o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_WS3;c_PRR-12;o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_WS3;c_PRR-12;o_LD1-PA13:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%
k_Bacteria:p_WS3;c_PRR-12;o_PBS-III-9:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_WS3;c_PRR-12;o_Sediment-1;Other;Other	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_WS3;c_PRR-12;o_Sediment-1:f_:g_	0.1%	0.0%	0.2%	0.1%	0.0%	0.3%	0.2%
k_Bacteria:p_WS3;c_PRR-12;o_Sediment-1:f_CV106:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_WS3;c_PRR-12;o_Sediment-1:f_PRR-10:g_	0.1%	0.0%	0.2%	0.2%	0.0%	0.4%	0.2%
k_Bacteria:p_WS3;c_PRR-12;o_wb1_H11:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_WS4;c_:o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_WS5;c_:o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_WS6;c_:o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_WS6;c_B142;o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_WS6;c_SC72;o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_ZB3;c_:o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_ZB3;c_BS119;o_:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_[Caldithrix];c_KSB1;o_Ucn15732:f_:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_[Thermi];c_Deinococci;o_Deinococcales:f_Deinococcaceae:g_Deinococcus	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_[Thermi];c_Deinococci;o_Deinococcales:f_Deinococcaceae:g_R18-435	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_[Thermi];c_Deinococci;o_Deinococcales:f_Trueperaceae:g_	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_[Thermi];c_Deinococci;o_Deinococcales:f_Trueperaceae:g_B-42	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
k_Bacteria:p_[Thermi];c_Deinococci;o_Deinococcales:f_Trueperaceae:g_Truepera	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%