

	Class	Order	DNA		RNA	
			min	max	min	max
Otu000001	Micrococcaceae	Micrococcus	0	0.02	0	0.73
Otu000002	unclassified	unclassified	0.01	0.06	0	0.36
Otu000003	unclassified	unclassified	0	0.14	0	0.14
Otu000004	Moraxellaceae	Acinetobacter	0	0	0	0.87
Otu000005	Pseudomonadaceae	Pseudomonas	0	0.04	0	0.73
Otu000006	unclassified	unclassified	0	0.15	0	0.05
Otu000007	Burkholderiaceae	Burkholderia	0	0.02	0	0.4
Otu000008	Pseudomonadaceae	Pseudomonas	0	0.05	0	0.27
Otu000009	unclassified	unclassified	0	0.06	0	0.08
Otu000010	unclassified	unclassified	0	0.02	0	0.22
Otu000011	Comamonadaceae	unclassified	0	0.02	0	0.28
Otu000012	unclassified	unclassified	0	0.08	0	0.1
Otu000013	Microbacteriaceae	unclassified	0	0.03	0	0.11
Otu000014	Clostridiales_Incertae_Sedis_XI	Finegoldia	0	0	0	0.48
Otu000016	Comamonadaceae	unclassified	0	0.02	0	0.1
Otu000017	Chitinophagaceae	Sediminibacterium	0	0.04	0	0.09
Otu000018	Saprospiraceae	unclassified	0	0.01	0	0.13
Otu000019	unclassified	unclassified	0	0.04	0	0.26
Otu000020	Xanthomonadaceae	Stenotrophomonas	0	0	0	0.38
Otu000021	unclassified	unclassified	0	0.08	0	0.02
Otu000022	Cyclobacteriaceae	Algoriphagus	0	0.03	0	0.06
Otu000023	unclassified	unclassified	0	0.07	0	0.03
Otu000024	unclassified	unclassified	0	0.02	0	0.08
Otu000025	Sphingomonadaceae	Sphingomonas	0	0.12	0	0.07
Otu000026	unclassified	unclassified	0	0.01	0	0.05
Otu000029	Acidobacteria_Gp4_family_incertae_sedis	Gp4	0	0	0	0.19
Otu000030	Microbacteriaceae	unclassified	0	0.03	0	0.03
Otu000032	Bradyrhizobiaceae	Bradyrhizobium	0	0.01	0	0.13
Otu000033	Pseudomonadaceae	Pseudomonas	0	0	0	0.27
Otu000034	unclassified	unclassified	0	0.02	0	0.1
Otu000035	Comamonadaceae	Comamonas	0	0.01	0	0.06
Otu000036	Cytophagaceae	unclassified	0	0.09	0	0.01
Otu000037	unclassified	unclassified	0	0.02	0	0.11
Otu000038	Rhodobacteraceae	Rhodobacter	0	0	0	0.17
Otu000039	unclassified	unclassified	0	0.02	0	0.07
Otu000042	Flavobacteriaceae	Flavobacterium	0	0.05	0	0.03
Otu000043	Micrococcaceae	Arthrobacter	0	0.05	0	0.01
Otu000045	unclassified	unclassified	0	0	0	0.33
Otu000049	Flavobacteriaceae	Flavobacterium	0	0.02	0	0.02
Otu000051	Spartobacteria_family_incertae_sedis	Spartobacteria_genera_incertae_sedis	0	0.01	0	0
Otu000053	Acidobacteria_Gp6_family_incertae_sedis	Gp6	0	0.01	0	0
Otu000057	Comamonadaceae	unclassified	0	0.02	0	0.01
Otu000065	Bacillaceae_1	Bacillus	0	0.03	0	0
Otu000067	Comamonadaceae	unclassified	0	0.01	0	0.02
Otu000073	Planococcaceae	Lysinibacillus	0	0.06	0	0
Otu000077	Pseudomonadaceae	Pseudomonas	0	0.02	0	0
Otu000083	Sphingobacteriaceae	Mucilaginibacter	0	0.02	0	0
Otu000084	Oxalobacteraceae	Janthinobacterium	0	0.02	0	0
Otu000088	Oxalobacteraceae	Duganella	0	0.02	0	0
Otu000132	Enterobacteriaceae	Yersinia	0	0.01	0	0
Otu000164	unclassified	unclassified	0	0	0	0
Otu000191	Solirubrobacteraceae	Solirubrobacter	0	0	0	0
Otu000222	unclassified	unclassified	0	0	0	0
Otu000223	Sphingomonadaceae	unclassified	0	0	0	0
Otu000309	unclassified	unclassified	0	0	0	0
Otu000432	Acidobacteria_Gp1_family_incertae_sedis	Gp1	0	0	0	0
Otu000532	unclassified	unclassified	0	0	0	0

	Class	Order	DNA		RNA	
			min	max	min	max
Otu000015	Burkholderiaceae	Polynucleobacter	0	0.03	0	0.18
Otu000027	unclassified	unclassified	0	0.02	0	0.18
Otu000031	unclassified	unclassified	0	0.03	0	0.11
Otu000044	Chitinophagaceae	unclassified	0	0.01	0	0.02
Otu000056	unclassified	unclassified	0	0.02	0	0.02
Otu000058	unclassified	unclassified	0	0	0	0.14
Otu000087	unclassified	unclassified	0	0.01	0	0.03
Otu000098	unclassified	unclassified	0	0.02	0	0
Otu000219	Rubrobacteraceae	Rubrobacter	0	0.01	0	0
Otu000225	Micrococcaceae	unclassified	0	0.01	0	0