

	Class	Order	DNA		RNA	
			min	max	min	max
Otu000001	Actinobacteria	Actinomycetales	0	0.02	0	0.73
Otu000002	unclassified	unclassified	0.01	0.06	0	0.36
Otu000003	Actinobacteria	Actinomycetales	0	0.14	0	0.14
Otu000004	Gammaproteobacteria	Pseudomonadales	0	0	0	0.87
Otu000005	Gammaproteobacteria	Pseudomonadales	0	0.04	0	0.73
Otu000006	unclassified	unclassified	0	0.15	0	0.05
Otu000007	Betaproteobacteria	Burkholderiales	0	0.02	0	0.4
Otu000008	Gammaproteobacteria	Pseudomonadales	0	0.05	0	0.27
Otu000009	Actinobacteria	Actinomycetales	0	0.06	0	0.08
Otu000010	Sphingobacteria	Sphingobacteriales	0	0.02	0	0.22
Otu000011	Betaproteobacteria	Burkholderiales	0	0.02	0	0.28
Otu000012	Betaproteobacteria	unclassified	0	0.08	0	0.1
Otu000013	Actinobacteria	Actinomycetales	0	0.03	0	0.11
Otu000014	Clostridia	Clostridiales	0	0	0	0.48
Otu000016	Betaproteobacteria	Burkholderiales	0	0.02	0	0.1
Otu000017	Sphingobacteria	Sphingobacteriales	0	0.04	0	0.09
Otu000018	Sphingobacteria	Sphingobacteriales	0	0.01	0	0.13
Otu000019	Alphaproteobacteria	Rhizobiales	0	0.04	0	0.26
Otu000020	Gammaproteobacteria	Xanthomonadales	0	0	0	0.38
Otu000021	Actinobacteria	unclassified	0	0.08	0	0.02
Otu000022	Sphingobacteria	Sphingobacteriales	0	0.03	0	0.06
Otu000023	Actinobacteria	Actinomycetales	0	0.07	0	0.03
Otu000024	Actinobacteria	Actinomycetales	0	0.02	0	0.08
Otu000025	Alphaproteobacteria	Sphingomonadales	0	0.12	0	0.07
Otu000026	unclassified	unclassified	0	0.01	0	0.05
Otu000029	Acidobacteria_Gp4	Acidobacteria_Gp4_order_incertae_sedis	0	0	0	0.19
Otu000030	Actinobacteria	Actinomycetales	0	0.03	0	0.03
Otu000032	Alphaproteobacteria	Rhizobiales	0	0.01	0	0.13
Otu000033	Gammaproteobacteria	Pseudomonadales	0	0	0	0.27
Otu000034	unclassified	unclassified	0	0.02	0	0.1
Otu000035	Betaproteobacteria	Burkholderiales	0	0.01	0	0.06
Otu000036	Sphingobacteria	Sphingobacteriales	0	0.09	0	0.01
Otu000037	unclassified	unclassified	0	0.02	0	0.11
Otu000038	Alphaproteobacteria	Rhodobacterales	0	0	0	0.17
Otu000039	Gammaproteobacteria	unclassified	0	0.02	0	0.07
Otu000042	Flavobacteria	Flavobacteriales	0	0.05	0	0.03
Otu000043	Actinobacteria	Actinomycetales	0	0.05	0	0.01
Otu000045	Betaproteobacteria	unclassified	0	0	0	0.33
Otu000049	Flavobacteria	Flavobacteriales	0	0.02	0	0.02
Otu000051	Spartobacteria	Spartobacteria_order_incertae_sedis	0	0.01	0	0
Otu000053	Acidobacteria_Gp6	Acidobacteria_Gp6_order_incertae_sedis	0	0.01	0	0
Otu000057	Betaproteobacteria	Burkholderiales	0	0.02	0	0.01
Otu000065	Bacilli	Bacillales	0	0.03	0	0
Otu000067	Betaproteobacteria	Burkholderiales	0	0.01	0	0.02
Otu000070	unclassified	unclassified	0	0.01	0	0.02
Otu000073	Bacilli	Bacillales	0	0.06	0	0
Otu000074	Betaproteobacteria	Burkholderiales	0	0.02	0	0.01
Otu000077	Gammaproteobacteria	Pseudomonadales	0	0.02	0	0
Otu000083	Sphingobacteria	Sphingobacteriales	0	0.02	0	0
Otu000084	Betaproteobacteria	Burkholderiales	0	0.02	0	0
Otu000088	Betaproteobacteria	Burkholderiales	0	0.02	0	0
Otu000099	Opitutae	Opitutales	0	0.01	0	0.01
Otu000132	Gammaproteobacteria	Enterobacteriales	0	0.01	0	0
Otu000164	Bacilli	Bacillales	0	0	0	0
Otu000191	Actinobacteria	Solirubrobacterales	0	0	0	0
Otu000222	Gammaproteobacteria	unclassified	0	0	0	0
Otu000223	Alphaproteobacteria	Sphingomonadales	0	0	0	0
Otu000309	Actinobacteria	Solirubrobacterales	0	0	0	0
Otu000432	Acidobacteria_Gp1	Acidobacteria_Gp1_order_incertae_sedis	0	0	0	0
Otu000532	Deltaproteobacteria	Myxococcales	0	0	0	0

	Class	Order	DNA		RNA	
			min	max	min	max
Otu000015	Betaproteobacteria	Burkholderiales	0	0.03	0	0.18
Otu000027	unclassified	unclassified	0	0.02	0	0.18
Otu000031	Actinobacteria	Actinomycetales	0	0.03	0	0.11
Otu000044	Sphingobacteria	Sphingobacteriales	0	0.01	0	0.02
Otu000055	Planctomycetacia	Planctomycetales	0	0.02	0	0.02
Otu000056	unclassified	unclassified	0	0.02	0	0.02
Otu000058	unclassified	unclassified	0	0	0	0.14
Otu000061	Opitutae	unclassified	0	0.02	0	0.01
Otu000062	unclassified	unclassified	0	0.01	0	0.05
Otu000087	Actinobacteria	unclassified	0	0.01	0	0.03
Otu000098	Betaproteobacteria	Burkholderiales	0	0.02	0	0
Otu000162	Alphaproteobacteria	Rhodospirillales	0	0	0	0.02
Otu000219	Actinobacteria	Rubrobacterales	0	0.01	0	0
Otu000225	Actinobacteria	Actinomycetales	0	0.01	0	0