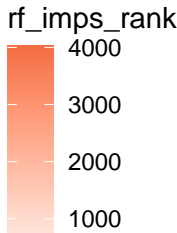


feature_Taxon

ASV_16200__d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Burkholderiales; f__Neisseriaceae; g__Eikenella; s__uncultured_bacterium	3835	188	108	42	41	34	22	174	99
ASV_4633__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Bacteroidales; f__Prevotellaceae; g__Prevotella_7	97	54	8	3	150	144	205	6	141
ASV_5782__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Flavobacteriales; f__Flavobacteriaceae; g__Capnocytophaga; s__Capnocytophaga_gingivalis	26	37	160	197	121	156	3888	73	129
ASV_4976__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Bacteroidales; f__Porphyromonadaceae; g__Porphyromonas; s__Porphyromonas_catoniae	170	233	3146	260	368	94	30	5	120
ASV_8226__d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacterales; f__Pasteurellaceae; g__Actinobacillus	201	87	61	21	73	31	82	10	15
ASV_1665__d__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales	140	3	51	95	25	19	2	9	4
ASV_9113__d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacterales; f__Pasteurellaceae; g__Haemophilus	52	192	12	433	234	158	64	215	49
ASV_6949__d__Bacteria; p__Actinobacteriota; c__Actinobacteria; o__Micrococcales; f__Micrococcaceae; g__Rothia	3838	20	16	105	4	191	95	26	162
ASV_16756__d__Bacteria; p__Actinobacteriota; c__Actinobacteria; o__Actinomycetales; f__Actinomycetaceae; g__Actinomyces	11	4	4	19	36	5	34	53	6
ASV_7797__d__Bacteria; p__Actinobacteriota; c__Actinobacteria; o__Corynebacteriales; f__Corynebacteriaceae; g__Corynebacterium; s__Corynebacterium_matruchotii	6	7	102	177	98	52	56	124	7
ASV_6951__d__Bacteria; p__Actinobacteriota; c__Actinobacteria; o__Micrococcales; f__Micrococcaceae; g__Rothia	132	9	3	63	3	11	7	366	3
ASV_4823__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Bacteroidales; f__Tannerellaceae; g__Tannerella; s__Tannerella_sp.	274	76	508	4	39	40	12	29	482
ASV_5306__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Bacteroidales; f__Prevotellaceae; g__Prevotella_7	74	183	31	37	369	3	6	11	22
ASV_3643__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Flavobacteriales; f__Flavobacteriaceae; g__Capnocytophaga	161	123	59	20	9	137	226	55	157
ASV_12302__d__Bacteria; p__Firmicutes; c__Negativicutes; o__Veillonellales–Selenomonadales; f__Veillonellaceae; g__Veillonella	3812	46	14	11	277	108	256	96	112
ASV_3559__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Flavobacteriales; f__Flavobacteriaceae; g__Capnocytophaga; s__Capnocytophaga_sputigena	130	93	357	46	54	341	149	3464	2
ASV_4847__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Bacteroidales; f__Porphyromonadaceae; g__Porphyromonas	125	45	154	47	30	182	65	3429	118
ASV_1938__d__Bacteria; p__Firmicutes; c__Bacilli; o__Staphylococcales; f__Gemellaceae; g__Gemella	25	198	214	58	37	2300	18	101	20
ASV_7886__d__Bacteria; p__Actinobacteriota; c__Actinobacteria; o__Propionibacteriales; f__Propionibacteriaceae; g__Propionibacterium	17	2	30	5	12	4	11	46	5
ASV_1186__d__Bacteria; p__Campylobacterota; c__Campylobacteria; o__Campylobacterales; f__Campylobacteraceae; g__Campylobacter	36	69	26	35	17	2459	25	63	507
ASV_4983__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Bacteroidales; f__Porphyromonadaceae; g__Porphyromonas	13	17	82	299	232	165	40	18	14
ASV_10606__d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Cardiobacteriales; f__Cardiobacteriaceae; g__Cardiobacterium	213	90	17	99	14	12	17	19	4017
ASV_12878__d__Bacteria; p__Actinobacteriota; c__Coriobacteriia; o__Coriobacteriales; f__Atopobiaceae; g__Atopobium	236	136	197	23	238	51	99	43	106
ASV_4904__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Bacteroidales; f__Porphyromonadaceae; g__Porphyromonas	2	24	78	49	53	15	5	2	25
ASV_2679__d__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales; f__Streptococcaceae; g__Streptococcus	204	8	140	6	116	255	416	273	4020
ASV_7755__d__Bacteria; p__Actinobacteriota; c__Actinobacteria; o__Corynebacteriales; f__Corynebacteriaceae; g__Corynebacterium; s__Corynebacterium_matruchotii	56	6	37	27	28	117	126	157	3988
ASV_8872__d__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Enterobacterales; f__Pasteurellaceae; g__Aggregatibacter	5	30	18	24	11	6	63	3	122
ASV_10960__d__Bacteria; p__Fusobacteriota; c__Fusobacteriia; o__Fusobacteriales; f__Leptotrichiaceae; g__Leptotrichia; s__Leptotrichia_sp.	16	2626	134	30	123	73	518	56	16
ASV_1822__d__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales; f__Carnobacteriaceae; g__Granulicatella	92	25	22	22	31	29	16	28	3943
ASV_3980__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Bacteroidales; f__Prevotellaceae; g__Prevotella; s__Prevotella_saccharolytica	103	49	85	82	2732	151	15	75	309
ASV_7424__d__Bacteria; p__Actinobacteriota; c__Actinobacteria; o__Actinomycetales; f__Actinomycetaceae; g__Actinomyces	3815	44	5	186	144	77	322	81	121
ASV_1989__d__Bacteria; p__Firmicutes; c__Bacilli; o__Staphylococcales; f__Gemellaceae; g__Gemella	60	79	44	29	34	88	88	291	38
ASV_5726__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Flavobacteriales; f__Flavobacteriaceae; g__Capnocytophaga; s__Capnocytophaga_granulosa	54	26	13	44	79	2449	13	32	12
ASV_2160__d__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales; f__Streptococcaceae; g__Streptococcus; s__Streptococcus_mutans	1	1	1	1	2	1	1	1	1
ASV_1443__d__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales; f__Lactobacillaceae; g__Ligilactobacillus	373	75	28	7	67	212	97	16	154
ASV_4653__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Bacteroidales; f__Prevotellaceae; g__Prevotella_7; s__Prevotella_histicola	30	33	10	12	13	9	84	179	396
ASV_8145__d__Bacteria; p__Firmicutes; c__Clostridia; o__Lachnospirales; f__Lachnospiraceae; g__Oribacterium	149	85	164	14	50	17	21	235	102
ASV_7121__d__Bacteria; p__Actinobacteriota; c__Actinobacteria; o__Actinomycetales; f__Actinomycetaceae; g__Actinomyces	86	19	11	15	286	10	59	14	27
ASV_5346__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Bacteroidales; f__Prevotellaceae	148	427	29	154	2755	249	193	230	273
ASV_7510__d__Bacteria; p__Actinobacteriota; c__Actinobacteria; o__Bifidobacteriales; f__Bifidobacteriaceae; g__Scardovia; s__Scardovia_wiggisiae	259	14	36	2	23	2	28	64	8
ASV_5000__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Bacteroidales; f__Porphyromonadaceae; g__Porphyromonas	50	331	141	48	160	64	9	27	3977
ASV_5586__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Bacteroidales; f__Prevotellaceae; g__Prevotella	84	330	433	122	2825	72	80	30	142
ASV_5893__d__Bacteria; p__Patescibacteria; c__Saccharimonadia; o__Saccharimonadales; f__Saccharimonadaceae; g__TM7x	4	114	122	128	321	2425	66	136	44
ASV_6531__d__Bacteria; p__Actinobacteriota; c__Actinobacteria; o__Actinomycetales; f__Actinomycetaceae; g__Actinomyces; s__Schaalia_odontolytica	72	12	34	8	1	30	8	248	138
ASV_12257__d__Bacteria; p__Firmicutes; c__Negativicutes; o__Veillonellales–Selenomonadales; f__Veillonellaceae; g__Veillonella	9	41	143	33	107	14	54	145	90
ASV_7597__d__Bacteria; p__Actinobacteriota; c__Actinobacteria; o__Corynebacteriales; f__Corynebacteriaceae; g__Corynebacterium; s__Corynebacterium_durum	147	59	70	18	18	28	27	8	26
ASV_7774__d__Bacteria; p__Actinobacteriota; c__Actinobacteria; o__Corynebacteriales; f__Corynebacteriaceae; g__Corynebacterium; s__Corynebacterium_matruchotii	311	147	218	2306	168	210	528	598	10
ASV_3575__d__Bacteria; p__Bacteroidota; c__Bacteroidia; o__Flavobacteriales; f__Flavobacteriaceae; g__Capnocytophaga; s__Capnocytophaga_sputigena	33	250	132	137	244	71	112	3552	9
ASV_9466__d__Bacteria; p__Fusobacteriota; c__Fusobacteriia; o__Fusobacteriales; f__Fusobacteriaceae; g__Fusobacterium	18	71	9	9	20	24	10	21	71
ASV_7149__d__Bacteria; p__Actinobacteriota; c__Actinobacteria; o__Actinomycetales; f__Actinomycetaceae; g__Actinomyces	77	131	56	28	7	38	3	401	50
ASV_1504__d__Bacteria; p__Firmicutes; c__Bacilli; o__Lactobacillales; f__Lactobacillaceae; g__Limosilactobacillus	194	40	424	13	398	2112	92	51	199



dataset