#OTU ID	B1.1	B1.2	B1.3	W1.1	W1.2	W1.3	E1.1	E1.2	E1.3	Z1.1	Z1.2	Z1.3	Taxonomy
ASV0	25	18	33	12	19	8	5	14	9	11	22	16	k__Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__Acetobacterales;f__Acetobacteraceae;g__Komagataeibacter
ASV1	103	95	158	62183	55446	21035	59871	63450	124	7455	1987	8056	k__Bacteria;p__Actinobacteriota;c__Actinomycetia;o__Mycobacteriales;f__Corynebacteriaceae;g__Corynebacterium
ASV2	45	38	91	46350	40890	15770	42100	49500	98	215	122	89	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Vibrionales;f__Vibrionaceae;g__Vibrio
ASV3	88	105	123	75	92	45	68	81	33	24	31	19	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Enterobacterales;f__Morganellaceae;g__Providencia
ASV4	65	77	99	61	85	39	55	73	28	15	25	12	k__Bacteria;p__Firmicutes;c__Bacilli;o__Lactobacillales;f__Enterococcaceae;g__Enterococcus
ASV5	38560	41234	59870	35	41	29	22	30	18	2560	3120	2890	k__Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__Sphingomonadales;f__Sphingomonadaceae;g__Sphingomonas
ASV6	32	45	61	18	22	11	15	20	9	1890	6123	5987	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Enterobacterales;f__Enterobacteriaceae;g__Pantoea
ASV7	21	19	34	10	14	7	8	11	5	4120	3890	4015	k__Bacteria;p__Firmicutes;c__Bacilli;o__Staphylo

ASV8	2841	52	73	22	28	14	18	25	12	6032	5788	6104	coccales;f__Staphylococcaceae;g__Staphylococcus k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Pseudomonadales;f__Pseudomonadaceae;g__Pseudomonas
ASV9	18	23	35	9	12	6	7	10	4	4876	5011	4950	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Moraxellales;f__Moraxellaceae;g__Acinetobacter
ASV10	321	15	21	5	8	3	4	6	39	31	28	35	k__Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__Hyphomicrobiales;f__Methylobacteriaceae;g__Methylobacterium
ASV11	22	28	40	11	15	8	10	14	7	21	25	23	k__Bacteria;p__Firmicutes;c__Bacilli;o__Lactobacillales;f__Lactobacillaceae;g__Lactobacillus
ASV12	15	19	28	8	11	5	6	9	4	18	22	20	k__Bacteria;p__Firmicutes;c__Bacilli;o__Bacillales;f__Bacillaceae;g__Bacillus
ASV13	10	13	19	4	6	2	3	5	2	12	16	14	k__Bacteria;p__Actinobacteriota;c__Actinomycetia;o__Micrococcales;f__Microbacteriaceae;g__Microbacterium
ASV14	8	11	16	3	5	1	2	4	1	9	12	10	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Enterobacterales;f__Erwiniaceae;g__Erwinia
ASV15	7	9	14	2	4	0	1	3	0	7	10	8	k__Bacteria;p__Firmicutes;c__Bacilli;o__Lactobacillales;f__Streptococcaceae;g__Streptococcus

ASV16	6	8	12	1	3	0	0	2	0	6	8	7	k__Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__Rhodospirillales;f__Acetobacteraceae;g__Roseomonas
ASV17	5	7	10	0	2	0	0	1	0	5	7	6	k__Bacteria;p__Bacteroidota;c__Bacteroidia;o__Flavobacteriales;f__Flavobacteriaceae;g__Chryseobacterium
ASV18	4	6	9	0	1	0	0	0	0	4	6	5	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Burkholderiales;f__Comamonadaceae;g__Massilia
ASV19	3	5	8	0	0	0	0	0	0	3	5	4	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Enterobacterales;f__Enterobacteriaceae;g__Serratia
ASV20	2	4	7	0	0	0	0	0	0	2	4	3	k__Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__Rhizobiales;f__Rhizobiaceae;g__Rhizobium
ASV21	2	3	6	0	0	0	0	0	0	1	3	2	k__Bacteria;p__Actinobacteriota;c__Actinomycetia;o__Streptomycetales;f__Streptomycetaceae;g__Streptomyces
ASV22	1	2	5	0	0	0	0	0	0	0	2	1	k__Bacteria;p__Firmicutes;c__Bacilli;o__Lactobacillales;f__Carnobacteriaceae;g__Granulicatella
ASV23	0	1	7	0	0	0	0	0	0	0	1	0	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Xanthomonadales;f__Xanthomonadaceae;g__Stenotrophomonas

ASV24	0	0	3	0	0	0	0	0	0	0	0	0	k__Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__Rhodobacterales;f__Rhodobacteraceae;g__Paracoccus
ASV25	0	0	6	0	0	0	0	0	0	0	0	0	k__Bacteria;p__Firmicutes;c__Bacilli;o__Paenibacillales;f__Paenibacillaceae;g__Paenibacillus
ASV26	0	0	1	0	0	0	0	0	0	0	0	0	k__Bacteria;p__Actinobacteriota;c__Actinomycetia;o__Kitasatosporales;f__Kitasatosporaceae;g__Kitasatospora
ASV27	3545	0	0	0	0	0	0	0	0	2750	0	0	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Burkholderiales;f__Burkholderiaceae;g__Burkholderia-Caballeronia-Paraburkholderia
ASV28	0	0	0	0	0	0	0	7	0	0	0	0	k__Bacteria;p__Firmicutes;c__Clostridia;o__Eubacteriales;f__Lachnospiraceae;g__Blautia
ASV29	0	0	0	0	0	0	0	0	0	0	2	0	k__Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__Novosphingomonadales;f__Novosphingomonadaceae;g__Novosphingobium
ASV30	0	0	0	0	0	0	0	0	2	0	0	0	k__Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__Reyranellales;f__Reyranellaceae;g__Reyranella
ASV31	0	0	0	0	0	0	0	0	0	0	0	1	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Enterobacterales;f__Enterobacteriaceae;

												g__Enterobacter
ASV32	0	4	0	0	0	0	0	0	0	0	0	k__Bacteria;p__Actinobacteriota;c__Actinomycetia;o__Propionibacteriales;f__Nocardioidaceae;g__Nocardioides
ASV34	0	0	0	0	0	0	0	0	0	4	0	0	k__Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__Rhizobiales;f__Beijerinckiaceae;g__Beijerinckia
ASV35	0	0	8	0	0	0	0	0	0	0	0	0	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Enterobacterales;f__Enterobacteriaceae;g__Klebsiella
ASV36	0	0	0	6	0	0	0	0	0	0	0	0	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Burkholderiales;f__Comamonadaceae;g__Comamonas
ASV37	0	0	0	0	6	0	0	0	0	0	0	0	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Enterobacterales;f__Yersiniaceae;g__Rahnella
ASV38	0	0	0	0	0	0	1	0	0	0	0	0	k__Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__Rhizobiales;f__Phyllobacteriaceae;g__Mesorhizobium
ASV39	13	0	0	0	0	0	0	0	0	0	0	0	k__Bacteria;p__Firmicutes;c__Bacilli;o__Lactobacillales;f__Lactobacillaceae;g__Pediococcus
ASV40	0	0	0	0	0	0	0	0	0	0	0	3	k__Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__Sphingomonadales;f__Sphingomonadacea

ASV43	0	0	0	0	0	0	0	0	9	0	0	0	e;g__Sphingopyxis k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Lysobacterales;f__Lysobacteraceae;g__Lysobacter
ASV44	0	0	0	0	0	5	0	0	0	0	0	0	k__Bacteria;p__Firmicutes;c__Bacilli;o__Caryophanales;f__Paenibacillaceae;g__Brevibacillus
ASV45	0	6	0	0	0	0	0	0	0	0	0	0	k__Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__Rhodobacterales;f__Rhodobacteraceae;g__Gemmobacter
ASV46	0	0	0	6	0	0	0	0	0	0	0	0	k__Bacteria;p__Actinobacteriota;c__Actinomycetia;o__Micrococcales;f__Micrococcaceae;g__Rothia
ASV47	0	0	0	0	0	0	3	0	0	0	0	0	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Cellvibrionales;f__Halieaceae;g__Halioglobus
ASV48	0	0	7	0	0	0	0	0	0	0	0	0	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Burkholderiales;f__Sutterellaceae;g__Parasutterella
ASV49	0	0	0	0	5	0	0	0	0	0	0	0	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Aeromonadales;f__Aeromonadaceae;g__Aeromonas
ASV50	0	0	0	0	0	0	0	0	0	5	0	0	k__Bacteria;p__Firmicutes;c__Bacilli;o__Lactobacillales;f__Leuconostocaceae;g__Weissella

ASV100	8	11	14	24	31	18	852	974	21	2855	15	19	k__Bacteria;p__Actinobacteriota;c__Actinomycetia;o__Propionibacteriales;f__Propionibacteriaceae;g__Cutibacterium
ASV107	5	0	0	0	0	0	0	0	0	0	0	0	k__Bacteria;p__Firmicutes;c__Clostridia;o__Lachnospirales;f__Lachnospiraceae;g__Fusicatenibacter
ASV108	0	0	0	0	0	0	0	8	0	0	0	0	k__Bacteria;p__Bacteroidota;c__Bacteroidia;o__Bacteroidales;f__Tannerellaceae;g__Parabacteroides
ASV109	0	0	0	0	0	0	0	0	0	0	5	0	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Pseudomonadales;f__Pseudomonadaceae;g__Pseudomonas
ASV110	0	0	0	0	0	0	0	0	6	0	0	0	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Enterobacterales;f__Enterobacteriaceae;g__Gibbsiella
ASV111	0	0	0	0	0	0	0	0	0	0	0	2	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Enterobacterales;f__Pectobacteriaceae;g__Pectobacterium
ASV112	0	11	0	0	0	0	0	0	0	0	0	0	k__Bacteria;p__Proteobacteria;c__Alphaproteobacteria;o__Rhizobiales;f__Methylobacteriaceae;g__Methylobacterium-methylorubrum
ASV113	0	0	0	0	0	4	0	0	0	0	0	0	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Burkholderiales;f__Rhodocyclaceae;g__Denitratisoma

ASV114	7	0	0	0	0	0	0	0	0	0	0	k__Bacteria;p__Proteobacteria;c__Gammaproteobacteria;o__Enterobacterales;f__Enterobacteriaceae;g__Izhakiella