





OTU	Number of reads		Greengenes taxonomic classification
	Wol-	Wol+	
Otu111	7	0	Bacteria(100);Proteobacteria(100);Alphaproteobacteria(100);PB21(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu112	1	2	Bacteria(100);Cyanobacteria(100);Chloroplasts(100);Dinophysis(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu113	2	0	Bacteria(100);Bacteroidetes(100);Bacteroidales(100);vadinBC27(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu114	1	1	Bacteria(100);Firmicutes(100);Clostridia(100);Clostridiales(100);Ruminococcus(100);p-1082-a5(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu115	8	0	Bacteria(100);Proteobacteria(100);Gammaproteobacteria(100);Enterobacteriales_Enterobacteriaceae(100);Yersinia(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu116	5	0	Bacteria(100);Firmicutes(100);Clostridia(100);Clostridiales(100);Johnsonella(100);Eubacterium_cf_saburreum_oral_strain_C27KA(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu117	2	1	Bacteria(100);TM7(100);TM7-1(100);K20-27(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu118	1	0	Bacteria(100);Bacteroidetes(100);Bacteroidales(100);Porphyromonadaceae(100);Porphyromonas_canis(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu119	1	0	Bacteria(100);ABY1_OD1(100);FW129(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu120	1	1	Bacteria(100);Firmicutes(100);Clostridia(100);Peptostreptococaceae(100);Frigovirgula_patagoniensis(100);Clostridium_sticklandii(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu121	4	1	Bacteria(100);Actinobacteria(100);Actinobacteridae(100);Bifidobacteriaceae(100);Gardnerella(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu122	2	2	Bacteria(100);ZB3(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu123	1	0	Bacteria(100);Proteobacteria(100);Gammaproteobacteria(100);Methylococaceae(100);Hyd24-01(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu124	1	0	Bacteria(100);Lentisphaerae(100);Victivallales(100);Victivallaceae(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu125	1	0	Bacteria(100);Proteobacteria(100);Gammaproteobacteria(100);Ferrimonadaceae(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu126	1	1	Bacteria(100);Firmicutes(100);Mollicutes(100);Spiroplasma_citri(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu127	1	1	Bacteria(100);Firmicutes(100);Clostridia(100);Peptostreptococaceae(100);Peptoniphilus(100);Peptoniphilus_indolicus(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu128	1	2	Bacteria(100);Firmicutes(100);Desulfotomaculum(100);Desulfotomaculum_thermocisternum(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu129	1	0	Bacteria(100);Verrucomicrobia(100);Spartobacteria(100);Chthoniobacter(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu130	1	2	Bacteria(100);WS6(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu131	1	0	Bacteria(100);Firmicutes(100);Clostridia(100);Peptostreptococaceae(100);Soehngenia(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu132	81	0	Bacteria(100);Firmicutes(100);Clostridia(100);Clostridiales(100);Coprococcus(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu133	0	1	Bacteria(100);Proteobacteria(100);Gammaproteobacteria(100);Alteromonadales(100);Pseudoalteromonadaceae(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu134	0	1	Bacteria(100);Proteobacteria(100);Gammaproteobacteria(100);Moraxellaceae(100);Moraxella_Psychrobacter(100);Psychrobacter(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu135	5	2	Bacteria(100);Proteobacteria(100);Gammaproteobacteria(100);Betaproteobacteria(100);Comamonadaceae(100);Comamonas(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu136	8	0	Bacteria(100);Firmicutes(100);Clostridia(100);Clostridiales(100);bacterium_MDA2477(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu137	2	4	Bacteria(100);SR1(100);BH1(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu138	1	4	Bacteria(100);Bacteroidetes(100);Sphingobacteriaceae(100);Sphingobacterium(100);Sphingobacterium_multivorum(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu139	5	0	Bacteria(100);Bacteroidetes(100);Bacteroidales(100);Tannerellaceae(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu140	0	3	Bacteria(100);Actinobacteria(100);Actinobacteridae(100);Brachyacterium(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu141	1	0	Bacteria(100);Bacteroidetes(100);Bacteroidales(100);Bacteroidaceae(100);B_coprocota(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu142	0	4	Bacteria(100);Actinobacteria(100);Actinobacteridae(100);Rhodococcus_B(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu143	1	0	Bacteria(100);Actinobacteria(100);Acidimicrobiae(100);EB1017_group(100);Acidimicrobiae_bacterium_Ellin7143(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu144	3	0	Bacteria(100);Proteobacteria(100);Gammaproteobacteria(100);Betaproteobacteria(100);Nitrosomonadales(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu145	1	0	Bacteria(100);Nitrospirae(100);Thermodesulfobiontales(100);Thermodesulfobiontales(100);Thermodesulfobiontales(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu146	0	3	Bacteria(100);Proteobacteria(100);Gammaproteobacteria(100);Betaproteobacteria(100);Alcaligenaceae(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu147	2	0	Bacteria(100);Bacteroidetes(100);Bacteroidales(100);p-184-o5(100);rc5-47(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu148	4	3	Bacteria(100);Proteobacteria(100);Alphaproteobacteria(100);JP57(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu149	1	0	Bacteria(100);Proteobacteria(100);Gammaproteobacteria(100);Nevskiaceae(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu150	1	0	Bacteria(100);Actinobacteria(100);Actinobacteridae(100);Propionibacterineae(100);Nocardioideae(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu151	3	0	Bacteria(100);Proteobacteria(100);Pseudomonadaceae(100);glacial_ice_bacterium_M3C47K-2(100);Pseudomonas_psychrotolerans(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu152	5	0	Bacteria(100);Proteobacteria(100);Epsilonproteobacteria(100);Helicobacteriales(100);Helicobacteraceae(100);Wolinella(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu153	1	0	Bacteria(100);Firmicutes(100);Clostridia(100);Clostridiales(100);Clostridiaceae(100);pDH-A(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu154	0	1	Bacteria(100);Spirochaetes(100);Treponemaceae(100);Treponema(100);Rs-D01(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu155	4	0	Bacteria(100);Lactobacillales(100);Lactobacillaceae(100);Lactobacillus(100);Lactobacillus_delbrueckii(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu156	0	1	Bacteria(100);Proteobacteria(100);Gammaproteobacteria(100);Betaproteobacteria(100);Sutterellaceae(100);HuCA4(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu157	1	0	Bacteria(100);Proteobacteria(100);Alphaproteobacteria(100);Ellin314(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu158	1	0	Bacteria(100);Firmicutes(100);Clostridia(100);Clostridiales(100);C21_k11(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu159	0	1	Bacteria(100);Proteobacteria(100);Gammaproteobacteria(100);Betaproteobacteria(100);Comamonadaceae(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu160	6	3	Bacteria(100);Proteobacteria(100);Alphaproteobacteria(100);Sphingomonadales(100);Sphingobium(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu161	0	2	Bacteria(100);Acidobacteria(100);iii1-15(100);uncultivated_soil_bacterium_clone_C112(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu162	1	1	Bacteria(100);Actinobacteria(100);Actinobacteridae(100);Segniliparaceae(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu163	2	1	Bacteria(100);Firmicutes(100);Bacilli(100);Lactobacillales(100);Aerococcaceae(100);Unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu164	1	0	Bacteria(100);Proteobacteria(100);Alphaproteobacteria(100);Crabtreeella(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);
Otu165	2	0	Bacteria(100);Firmicutes(100);Clostridia(100);Peptostreptococaceae(100);Clostridium_felsineum(100);Clostridium_formicaceticum(100);unclassified(100);unclassified(100);unclassified(100);unclassified(100);

