

Phylum	Family	Genus	Species	OTU
Firmicutes	Lachnospiraceae	Lachnoclostridium	[Clostridium]_aldenense	OTU854
Firmicutes	Ruminococcaceae	GCA-900066225	unclassified_g_GCA-900066225	OTU928
Bacteroidetes	unclassified_o_Bacteroidales	unclassified_o_Bacteroidales	unclassified_o_Bacteroidales	OTU695
unclassified_k_norank_d_Bacteria	unclassified_k_norank_d_Bacteria	unclassified_k_norank_d_Bacteria	unclassified_k_norank_d_Bacteria	OTU856
Bacteroidetes	Muribaculaceae	norank_f_Muribaculaceae	uncultured_bacterium_g_norank_f_Muribaculaceae	OTU925
Bacteroidetes	Muribaculaceae	norank_f_Muribaculaceae	uncultured_bacterium_g_norank_f_Muribaculaceae	OTU920
Firmicutes	Lachnospiraceae	Lachnoclostridium	unclassified_g_Lachnoclostridium	OTU810
Firmicutes	Lachnospiraceae	Hungatella	uncultured_bacterium_g_Hungatella	OTU735
Bacteroidetes	Rikenellaceae	Alistipes	unclassified_g_Alistipes	OTU837
Bacteroidetes	Tannerellaceae	Parabacteroides	unclassified_g_Parabacteroides	OTU765
Bacteroidetes	unclassified_o_Bacteroidales	unclassified_o_Bacteroidales	unclassified_o_Bacteroidales	OTU960
Bacteroidetes	Bacteroidaceae	Bacteroides	unclassified_g_Bacteroides	OTU924
Bacteroidetes	unclassified_o_Bacteroidales	unclassified_o_Bacteroidales	unclassified_o_Bacteroidales	OTU923
Proteobacteria	Enterobacteriaceae	Klebsiella	Klebsiella_pneumoniae_subsp_pneumoniae_g_Klebsiella	OTU896
Firmicutes	Lachnospiraceae	[Ruminococcus]_gnavus_group	uncultured_bacterium_g_[Ruminococcus]_gnavus_group	OTU892
Firmicutes	Lachnospiraceae	Robinsoniella	Robinsoniella_peoriensis_g_Robinsoniella	OTU914
Bacteroidetes	unclassified_o_Bacteroidales	unclassified_o_Bacteroidales	unclassified_o_Bacteroidales	OTU487
Bacteroidetes	Bacteroidaceae	Bacteroides	unclassified_g_Bacteroides	OTU855
Bacteroidetes	Bacteroidaceae	Bacteroides	unclassified_g_Bacteroides	OTU857
Firmicutes	Lachnospiraceae	Anaerostipes	Anaerostipes_caccae	OTU748
Bacteroidetes	Muribaculaceae	norank_f_Muribaculaceae	uncultured_bacterium_g_norank_f_Muribaculaceae	OTU852
Proteobacteria	norank_o_Rhodospirillales	norank_f_norank_o_Rhodospirillales	gut_metagenome_g_norank_f_norank_o_Rhodospirillales	OTU895
Bacteroidetes	Bacteroidaceae	Bacteroides	Bacteroides_caecimuris	OTU802
Bacteroidetes	Tannerellaceae	Parabacteroides	unclassified_g_Parabacteroides	OTU758
Firmicutes	Ruminococcaceae	Flavonifractor	unclassified_g_Flavonifractor	OTU766
Bacteroidetes	Bacteroidaceae	Bacteroides	unclassified_g_Bacteroides	OTU768
Firmicutes	Lachnospiraceae	Hungatella	uncultured_organism_g_Hungatella	OTU844
Bacteroidetes	unclassified_o_Bacteroidales	unclassified_o_Bacteroidales	unclassified_o_Bacteroidales	OTU948
Firmicutes	Erysipelotrichaceae	Coprobacillus	Coprobacillus_cateniformis	OTU738
Bacteroidetes	Bacteroidaceae	Bacteroides	Bacteroides_acidifaciens	OTU744
Verrucomicrobia	Akkermansiaceae	Akkermansia	unclassified_g_Akkermansia	OTU691
Bacteroidetes	unclassified_o_Bacteroidales	unclassified_o_Bacteroidales	unclassified_o_Bacteroidales	OTU908
Firmicutes	Peptostreptococcaceae	Clostridioides	Clostridioides_difficile_g_Clostridioides	OTU877
Firmicutes	Lachnospiraceae	Lachnoclostridium	unclassified_g_Lachnoclostridium	OTU876
Bacteroidetes	Muribaculaceae	norank_f_Muribaculaceae	unclassified_g_norank_f_Muribaculaceae	OTU915
Bacteroidetes	Muribaculaceae	norank_f_Muribaculaceae	unclassified_g_norank_f_Muribaculaceae	OTU849
Verrucomicrobia	Akkermansiaceae	Akkermansia	Akkermansia_muciniphila	OTU890