

ASSOCIATION BETWEEN METADATA AND RELATIVE ABUNDANCE OF GROUPS OF MICROBES

SMOKING HABIT - UC and CD	UC		CD	
	SMOKERS	NON-SMOKERS	SMOKERS-2	NON-SMOKERS-2
p__Firmicutes;c__Clostridia;o__Clostridiales;f__Peptostreptococcaceae;g__	0.000000	0.000081	0.000275	0.000009
p__Bacteroidetes;c__Bacteroidia;o__Bacteroidales;f__Prevotellaceae;g__Prevotella	0.133849	0.046244	0.000403	0.020871
p__Bacteroidetes;c__Bacteroidia;o__Bacteroidales;f__[Odoribacteraceae];g__Butyricimonas	0.000126	0.000242	0.000037	0.000079
p__Actinoc__Coriobacteridae;o__Coriobacterineae;f__Coriobacteriaceae;g__Eggerthella;s__lenta	0.000202	0.000073	0.000018	0.000070
p__Actinoc__Actinobacteridae;o__Bifidobacteriales;f__Bifidobacteriaceae;g__Bifidobacterium;s__adole	0.002345	0.036023	0.001375	0.005323
p__Firmicutes;c__Clostridia;o__Clostridiales;f__Clostridiaceae	0.001361	0.007342	0.032509	0.002131
p__Firmicutes;c__Clostridia;o__Clostridiales;f__Veillonellaceae	0.009328	0.004139	0.000843	0.002920

SITE OF DISEASE - CD	L1	L3
p__Firmicutes;c__Erysipelotrichia;o__Erysipelotrichales;f__Erysipelotrichaceae;g__	8.633E-03	2.244E-02
p__Firmicutes;c__Bacilli;o__Lactobacillales;f__Enterococcaceae;g__Enterococcus;s__faecalis	2.017E-04	4.034E-05

SITE OF DISEASE - UC	E1	E2	E3
p__Firmicutes;c__Clostridia;o__Clostridiales;f__[Mogibacteriaceae]	0.00235	0.00114	0.00071
p__Firmicutes;c__Clostridia;o__Clostridiales;f__	0.06815	0.05187	0.02447
p__Firmicutes;c__Clostridia;o__Clostridiales;f__Clostridiaceae;g__Clostridium	0.00903	0.00051	0.00062
p__Firmicutes;c__Clostridia;o__Clostridiales;f__Peptostreptococcaceae;g__	0.00020	0.00000	0.00002