Log Fold Changes of Inflammation by Tissue

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Order:Lachnospirales -	0.17	-0.19	
Family:uncultured -	-0.11	0.1	
Family:UCG-010 -	-0.1	0.19	
Family:Tannerellaceae -	0.16	0.01	
Family:Sutterellaceae -	-0.07	-0.06	
Family:Staphylococcaceae -	0.03	-0.05	
Family:Ruminococcaceae -	-0.06	0.12	
Family:Rikenellaceae -	0.22	0.04	
Family:RF39 -	0.06	-0.1	
Family:Peptostreptococcaceae -	0.24	0.11	
Family:Peptococcaceae -	0.07	-0.32	
Family:Oxalobacteraceae -	0.25	-0.29	
Family:Oscillospiraceae -	-0.06	0.18	
Family:Muribaculaceae -	-0.18	0.27	
Family:Monoglobaceae -	0.04	0.14	
Family:Marinifilaceae -	0.28	-0.22	
Family:Lactobacillaceae -	-0.15	-0.02	
Family:Lachnospiraceae -	-0.18	0.22	
Family:Erysipelotrichaceae -	0.02	-0.16	
Family:Erysipelatoclostridiaceae -	0.17	-0.33	
Family:Enterococcaceae -	-0.13	-0.47	
Family:Enterobacteriaceae -	0.2	-0.41	
Family:Eggerthellaceae -	-0.03	-0.02	
Family:Dysgonomonadaceae -	-0.44	-0.28	
Family:Defluviitaleaceae -	0.01	0.1	
Family:Deferribacteraceae -	-0.05	0.08	
Family:Clostridiaceae -	-0.07	0.04	
Family:Clostridia_vadinBB60_group -	0.07	0.06	
Family:Clostridia_UCG-014 -	-0.15	0	
Family:Christensenellaceae -	0.03	-0.01	
Family:Butyricicoccaceae -	-0.05	0.09	
Family:Bacteroidaceae -	-1.3	-0.06	
Family:Anaerovoracaceae -	-0.03	-0.06	
Family:Anaerofustaceae -	0.09	-0.2	
Family:Akkermansiaceae -	0.36	-0.72	
Family:Acholeplasmataceae -	0.02	-0.13	
Family:[Eubacterium]_coprostanoligenes_group -	-0.01	0.09	
Family:[Clostridium]_methylpentosum_group -	0.16	-0.1	
	Cecum	Colon	

0.0