Log Fold Changes by Diet Composition

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Order:Lachnospirales -	-0.07	0.22	0.02
Family:uncultured -	0.35	0.53	1.92
Family:UCG-010 -	0.03	0.32	-0.46
Family:Tannerellaceae -	-0.28	1.96	1.88
Family:Sutterellaceae -	0.69	1.71	1.79
Family:Staphylococcaceae -	1.03	-2	0.02
Family:Ruminococcaceae -	0.73	-0.42	-0.76
Family:Rikenellaceae -	0.16	0.44	0.54
Family:RF39 -	-0.36	0.24	-1.66
Family:Peptostreptococcaceae -	0.14	0.45	2.11
Family:Peptococcaceae -	-0.17	0.86	1.07
Family:Oxalobacteraceae -	3.07	-2.77	-3.29
Family:Oscillospiraceae -	0.18	-0.52	-0.54
Family:Muribaculaceae -	0.65	1.04	-2.16
Family:Monoglobaceae -	0.51	0.99	0.08
Family:Marinifilaceae -	-0.05	-0.64	2.37
Family:Lactobacillaceae -	0.57	0.44	-1.57
Family:Lachnospiraceae -	0.77	-0.12	-2.12
Family:Erysipelotrichaceae -	0.05	1.41	2.12
Family:Erysipelatoclostridiaceae -	0.4	0.15	0.31
Family:Enterococcaceae -	0.09	0.43	1.04
Family:Enterobacteriaceae -	0.58	2.5	2.81
Family:Eggerthellaceae -	0.45	-0.81	-1.49
Family:Defluviitaleaceae -	0.22	-0.41	0.45
Family:Deferribacteraceae -	0.57	-0.32	-0.17
Family:Clostridiaceae -	0.83	0.52	-0.85
Family:Clostridia_vadinBB60_group -	-0.34	1.39	-0.04
Family:Clostridia_UCG-014 -	-0.02	1.05	-1.22
Family:Christensenellaceae -	-0.22	-0.78	0.15
Family:Butyricicoccaceae -	-0.17	0	-0.93
Family:Bacteroidaceae -	-0.06	-0.75	-0.55
Family:Anaerovoracaceae -	0.2	-1.24	-1.25
Family:Anaerofustaceae -	0.16	-0.02	-0.32
Family:Akkermansiaceae -	0.28	-0.38	0.28
Family:Acholeplasmataceae -	-0.46	1.04	-1.43
Family:[Eubacterium]_coprostanoligenes_group -	0.38	-0.64	-0.69
	High Fat	High Fiber	Purified Diet