Log Fold Changes by Diet Composition

Log Fold Changes by Diet Compos			
Order:Lachnospirales -	0	0.22	0
Family:uncultured -	0.35	0.53	1.92
Family:UCG-010 -	0	0.32	-0.46
Family:Tannerellaceae -	0	1.96	1.88
Family:Sutterellaceae -	0.69	1.71	1.79
Family:Staphylococcaceae -	1.03	-2	0
Family:Ruminococcaceae -	0.73	-0.42	-0.76
Family:Rikenellaceae -	0	0.44	0.54
Family:RF39 -	-0.36	0.24	-1.66
Family:Peptostreptococcaceae -	0	0	2.11
Family:Peptococcaceae -	0	0.86	1.07
Family:Oxalobacteraceae -	3.07	-2.77	-3.29
Family:Oscillospiraceae -	0	-0.52	-0.54
Family:Muribaculaceae -	0.65	1.04	-2.16
Family:Monoglobaceae -	0.51	0.99	0
Family:Marinifilaceae -	0	-0.64	2.37
Family:Lactobacillaceae -	0.57	0	-1.57
Family:Lachnospiraceae -	0.77	0	-2.12
Family:Erysipelotrichaceae -	0	1.41	2.12
Family:Erysipelatoclostridiaceae -	0.4	0	0
Family:Enterococcaceae -	0	0	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
Family:Clostridiaceae -	0.83	0.52	-0.85
Family:Clostridia_vadinBB60_group -	-0.34	1.39	0
Family:Clostridia_UCG-014 -	0	1.05	-1.22
Family:Christensenellaceae -	-0.22	-0.78	0
Family:Butyricicoccaceae -	-0.17	0	-0.93
Family:Bacteroidaceae -	0	-0.75	-0.55
Family:Anaerovoracaceae -	0	-1.24	-1.25
Family:Anaerofustaceae -	0	0	-0.32
Family:Acholeplasmataceae -	-0.46	1.04	-1.43
Family:[Eubacterium]_coprostanoligenes_group -	0.38	-0.64	-0.69
_	High Fat	High Fiber	Purified Diet

3 2 1 0 -1 -2 -3