AD 2 months Versus AD 12 months

sample_group												
-0.43	-0.43	-0.43	-0.43	1.85	2.41	-0.43	-0.43	-0.43	-0.43	-0.43	-0.43	s__GGB28792_SGB41445
2.04	-0.54	-0.54	-0.54	0.95	1.83	-0.54	-0.54	-0.54	-0.54	-0.54	-0.54	s__GGB31838_SGB45216
1.22	0.89	-1.50	0.99	0.47	0.81	-0.16	-0.18	0.29	-1.50	0.16	-1.50	s__GGB42586_SGB59781
0.92	1.73	-0.66	-0.66	1.64	1.00	-0.66	-0.66	-0.66	-0.66	-0.66	-0.66	s__GGB28870_SGB41549
1.01	0.65	0.60	0.51	0.68	1.31	-1.29	-1.29	-1.29	0.04	0.34	-1.29	s__GGB31440_SGB44770
1.08	1.59	0.55	0.16	0.42	0.51	-0.36	-2.38	-0.11	-0.44	-0.66	-0.35	s__GGB27892_SGB40326
-0.58	0.13	0.33	-0.06	0.39	-2.85	0.47	1.10	0.21	-0.17	0.18	0.85	s__Clostridium_cocleatum
-0.65	-0.65	-0.65	-0.65	-0.65	-0.65	-0.65	0.60	1.18	-0.65	2.00	1.38	s__GGB25041_SGB36960
-0.53	-0.53	-0.53	-0.53	-0.53	-0.53	-0.53	0.74	-0.53	-0.53	1.95	2.04	s__GGB29683_SGB42492
0.71	-0.70	-1.70	-1.59	-0.02	0.45	-0.15	1.61	-0.22	1.11	0.67	-0.17	s__Clostridiaceae_bacterium
0.02	-1.36	-1.25	-0.46	1.48	-0.08	0.11	2.21	-0.39	-0.24	0.17	-0.22	s__GGB28949_SGB41656
-0.28	-0.98	-0.43	-1.11	-0.12	0.23	-1.07	2.52	0.43	-0.33	0.47	0.67	s__GGB28849_SGB41516
0.07	-1.05	-1.49	-0.87	0.19	0.12	-0.47	2.28	-0.30	0.44	-0.01	1.09	s__GGB28846_SGB41513
0.68	0.30	0.10	-1.68	-1.19	-1.10	0.38	2.06	0.58	0.06	-0.45	0.25	s__Streptococcus_danieliae
0.40	0.03	-1.12	-1.42	-0.59	-1.50	-0.08	1.43	0.56	0.85	0.06	1.38	s__GGB27885_SGB40319
0.16	-0.33	-0.49	-0.81	-0.80	-0.17	-0.35	2.83	-0.04	-0.66	-0.17	0.84	s__GGB28945_SGB41650
0.76	0.12	-0.05	-1.37	-0.75	0.43	-0.68	2.16	-0.32	-0.96	-0.51	1.17	s__Ligilactobacillus_murinus
-0.18	-0.57	-0.38	-1.36	0.00	-0.68	0.41	2.07	-0.29	-0.56	-0.27	1.81	s__Bacteroides_acidifaciens
-0.52	-0.05	-0.58	-1.50	-0.78	-0.52	0.37	1.96	0.87	-0.18	-0.55	1.48	s__GGB31179_SGB44472
-0.50	0.44	-0.50	-0.50	-0.50	-0.50	-0.50	1.49	2.52	-0.50	-0.50	-0.50	s__Jeotgalicoccus_halotolerans
-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	3.18	-0.29	-0.29	-0.29	s__Facklamia_tabacinasalis
-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	3.18	-0.29	-0.29	-0.29	s__GGB28383_SGB40963
-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	3.18	-0.29	-0.29	-0.29	s__Corynebacterium_stationis
-0.54	-0.54	-0.54	-0.54	-0.54	-0.54	-0.54	-0.54	1.91	0.98	1.95	-0.54	s__GGB30461_SGB43527
-0.15	-0.42	-1.39	0.12	-0.59	-0.93	0.23	0.78	1.38	0.16	1.94	-1.14	s__GGB28621_SGB41239
-0.53	1.37	-0.53	-0.53	-0.53	0.97	-0.53	2.39	-0.53	-0.53	-0.53	-0.53	s__GGB29006_SGB41724
-0.29	3.18	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	s__Staphylococcus_nepalensis
-0.29	3.18	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	s__GGB28456_SGB41039
-0.29	3.18	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	s__Mammaliicoccus_lentus
-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	3.18	-0.29	-0.29	s__GGB28382_SGB40962
-0.29	-0.29	-0.29	3.18	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	s__bacterium_1xD42_87
-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	3.18	-0.29	-0.29	-0.29	-0.29	-0.29	s__GGB29468_SGB42247
-0.29	-0.29	-0.29	-0.29	-0.29	-0.29	3.18	-0.29	-0.29	-0.29	-0.29	-0.29	s__GGB28781_SGB41434

sample_group
AD_2M
AD_12M