Wild Type versus AD 2 months sample_group sample group 2 WT_2M -2.07 0.02 -0.29-0.84-1.140.21 0.89 0.14 -0.050.87 1.59 s__GGB30454_SGB43514 AD_2M s__GGB28765_SGB41415 0.29 -0.14-0.61-0.83-0.010.41 0.80 0.37 0.87 0.89 0.60 1 -0.04-0.12-0.24-0.64-0.470.83 0.68 0.39 0.87 0.66 s__Clostridiales_bacterium 0 -1.30s__GGB27849_SGB40283 0.02 0.20 -1.37-0.65-1.040.93 1.91 0.91 0.09 0.66 -0.35-1.71-0.470.41 -1.310.21 -1.320.94 1.51 0.80 0.47 0.38 0.10 s__Muribaculum_gordoncarteri -1 -0.540.82 -0.97-1.36 -0.74-1.521.45 0.76 0.09 s__Atopobiaceae_bacterium_FL090493 0.18 0.96 0.86 -2 s__GGB30290_SGB43252 -1.100.28 -1.10-1.10-1.10-1.100.35 1.08 0.90 1.06 1.01 0.81 -1.100.08 -1.10-1.10-1.10-1.100.91 0.91 0.91 0.85 0.95 0.89 s__GGB31835_SGB45211 s__GGB28988_SGB41701 -0.47-0.12-0.23-1.82-1.820.42 0.80 1.25 0.28 -0.061.05 0.70 s__GGB28845_SGB41512 -0.88-0.88-0.88-0.88-0.88-0.880.15 1.03 0.22 1.14 1.63 1.13 s__GGB31841_SGB45219 -0.92-0.92-0.92-0.92-0.92-0.921.33 1.01 0.47 1.01 1.29 0.43 0.98 s__GGB31179_SGB44472 -0.70-0.65-0.87-0.68-1.24-1.400.98 1.18 0.96 0.58 0.87 0.07 0.99 0.80 0.93 0.13 1.01 -0.00-0.07-0.03-1.930.03 -1.93s__Candidatus_Arthromitus_sp_SFB_mouse 0.89 0.77 0.62 0.78 0.82 0.59 -1.590.11 -1.590.15 -1.59 s__GGB29003_SGB41719 0.03 1.37 1.49 -0.65-0.650.54 1.80 -0.65-0.65-0.65-0.65-0.65-0.65s__GGB28860_SGB41529 1.31 1.25 0.80 -0.370.05 1.25 -0.84-1.16-0.67-1.58-0.450.41 s__GGB28404_SGB40986 0.26 0.58 0.80 0.74 0.88 1.11 -0.140.14 -1.16-0.39-0.47s__Neglecta_sp_X4 0.87 0.49 1.17 0.89 1.19 1.04 -0.94-0.94-0.94-0.94-0.94-0.94s__GGB28851_SGB41518 s__GGB29683_SGB42492 0.92 1.08 0.94 1.12 0.81 0.84 -0.95-0.95-0.95-0.95-0.95-0.95s__GGB28879_SGB41559 0.29 0.44 1.28 0.99 1.00 0.85 -0.04-1.26-1.26-1.26-1.260.24 -0.650.22 0.57 1.26 1.40 1.00 0.45 -0.87-1.67-0.05-1.34-0.33s__GGB30303_SGB43268 -1.55s__GGB28870_SGB41545 -0.720.60 1.87 0.48 0.89 0.58 0.34 -0.62-1.52-0.28-0.08-0.510.46 0.61 1.09 0.36 -0.70-1.23-1.28-0.27-0.65-0.07s__GGB28927_SGB41625 -0.40s__GGB28904_SGB41597 0.41 -1.16-1.31-0.85-0.65-0.380.63 0.28 1.03 0.23 0.20 0.60 0.35 0.30 1.95 0.85 0.35 -1.24-1.79-0.59-0.21-0.77s__GGB30457_SGB43521 s__GGB30454_SGB43516 0.33 0.66 1.62 0.87 0.75 0.22 -0.26-0.15-1.03-0.30-0.460.28 0.73 1.48 0.71 0.68 0.67 -0.560.06 -1.77-1.77-0.01-0.50s__GGB28870_SGB41549 0.55 0.85 1.67 1.18 1.22 -0.78-0.78-0.78-0.78-0.78-0.78-0.78s__GGB30461_SGB43527 s__GGB28856_SGB41525 -0.681.39 1.52 1.29 1.21 -0.68-0.68-0.68-0.68-0.68-0.68-0.68-0.55s__GGB28383_SGB40963 -0.551.43 1.84 -0.55-0.551.69 -0.55-0.55-0.55-0.55-0.55s__GGB25052_SGB36974 -0.050.78 1.08 -0.65-0.35-0.07-0.20-1.49-0.91-0.540.12 -1.52 s GGB28873 SGB41553 -1.520.92 1.05 0.93 0.79 0.74 0.03 -0.100.15 -1.520.06 0.31 0.67 1.28 0.32 0.93 0.66 -0.17-1.90-0.24-1.900.24 -0.22s__GGB31853_SGB45233 0.89 1.43 0.93 0.60 0.55 -0.94-1.71-1.62-0.41-0.04s Clostridiaceae bacterium 0.16 0.16 -0.230.46 1.49 -1.06-2.16 0.09 s__GGB28888_SGB41569 0.65 0.93 0.53 0.27 -1.040.07