

Figure S1. Scatterplots of baseline cross-sectional associations between DNAm in genes within metabolically-relevant KEGG pathways and gut microbial taxa abundance

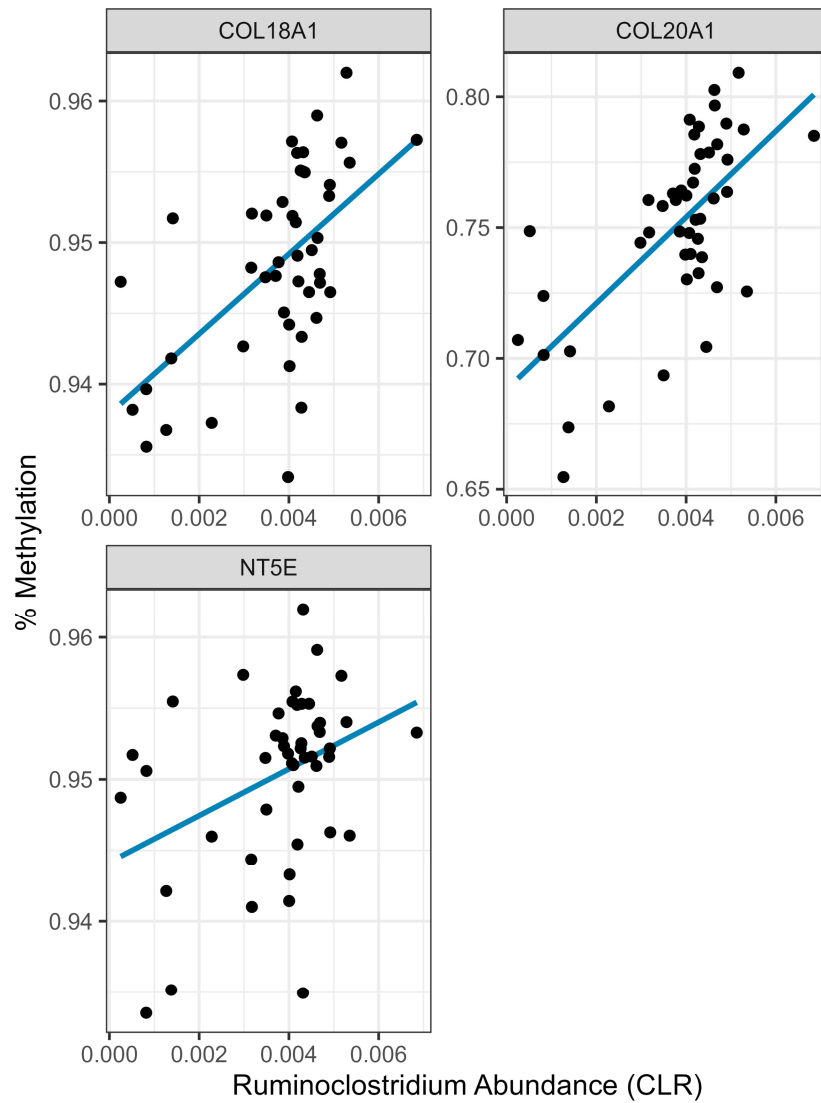


Figure S2. Scatterplots of 3-month cross-sectional associations between DNAm in genes within metabolically-relevant KEGG pathways and gut microbial taxa abundance

