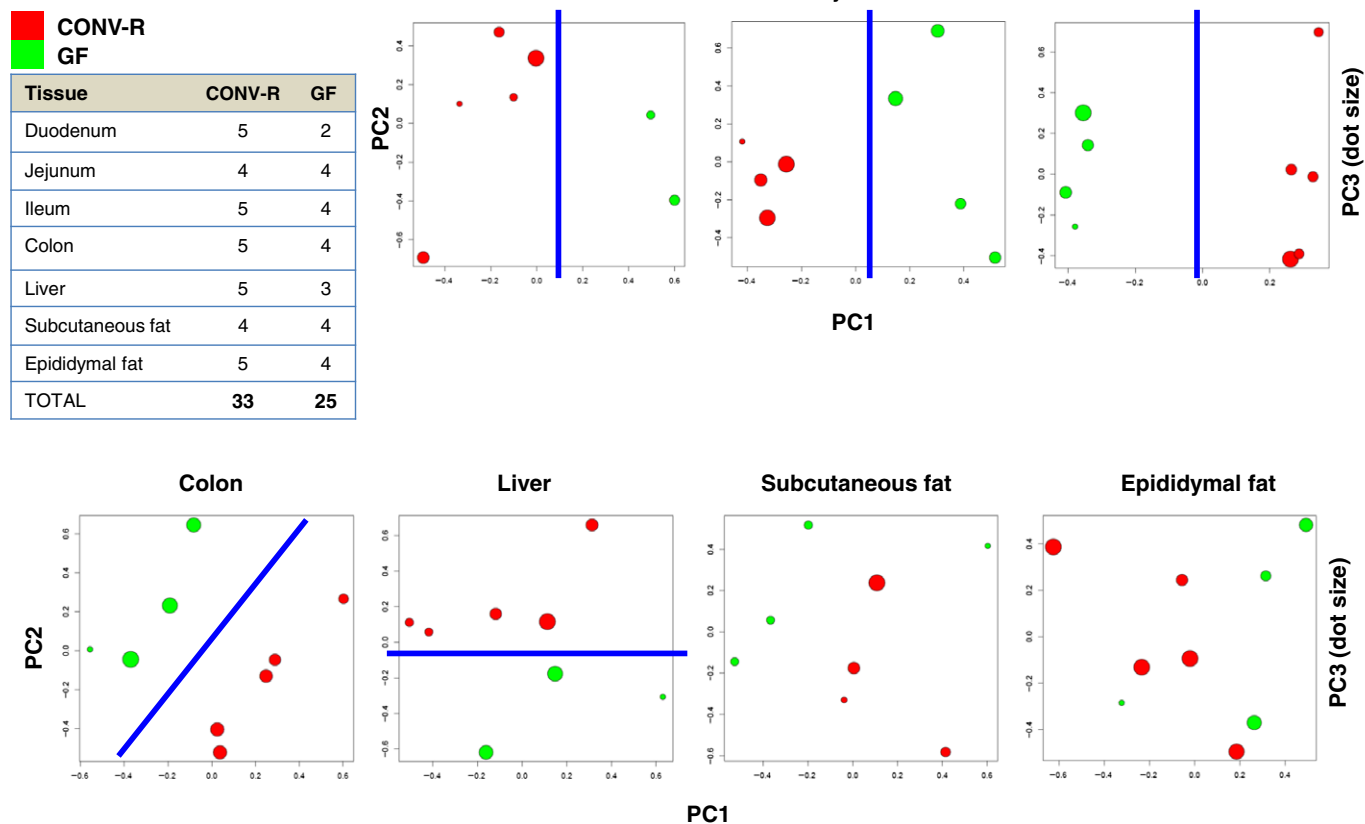# Expanded View Figures

Figure EV1. Gene expression profiling of CONV-R and GF mice tissues.
Global gene expression profile of seven different tissues including liver, epididymal and subcutaneous fat, duodenum, jejunum, ileum, and colon tissues has been generated, and each tissue sample is normalized independently. Principal component analysis (PCA) of transcription profiles on each tissue is presented.