

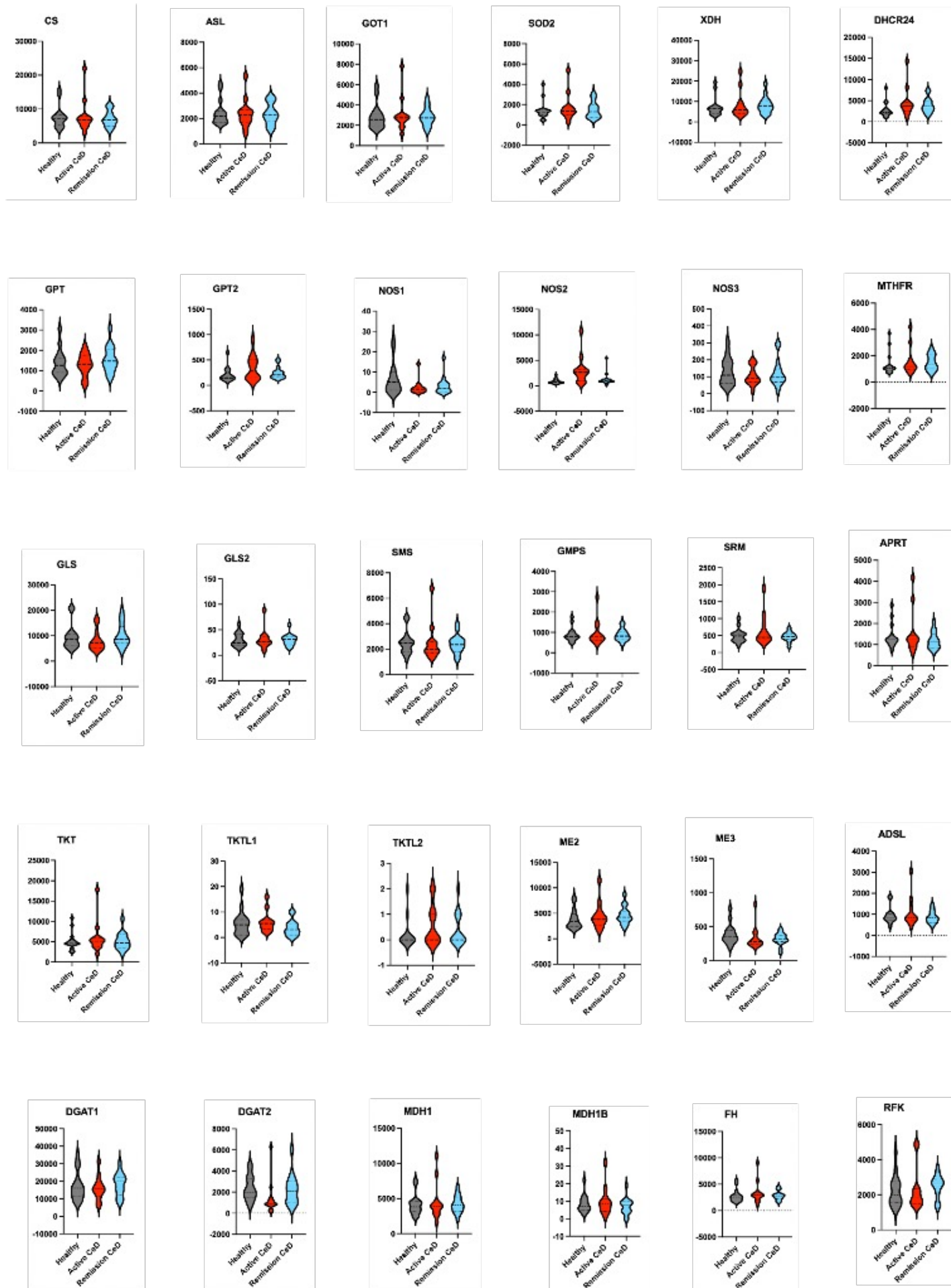
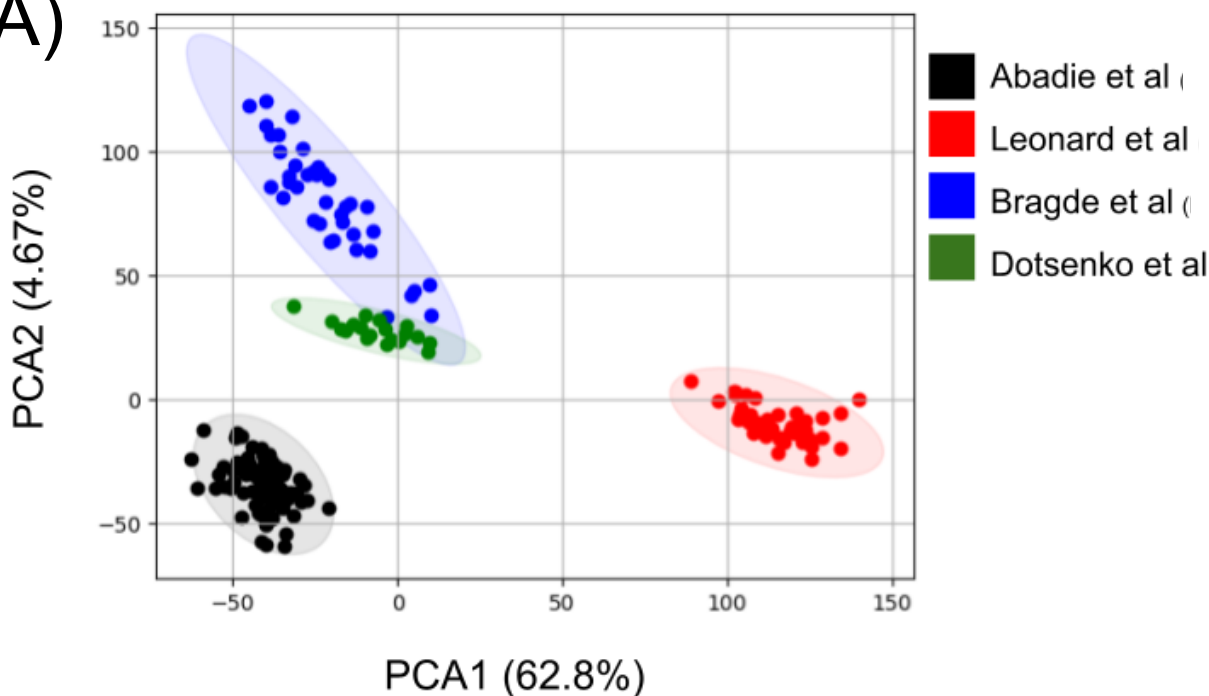


Figure S1. Differential expression analysis of genes encoding differentially active essential metabolic tasks.

A)



B)

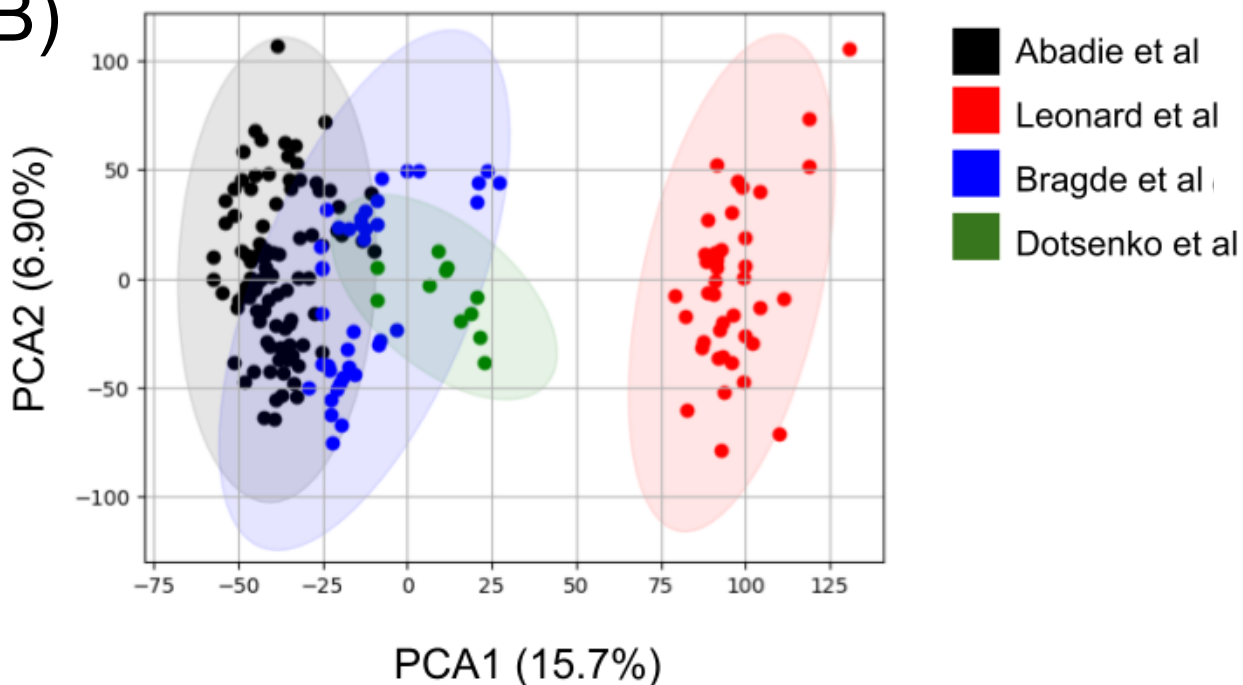


Figure S2. Principal Component Analysis (PCA) on RNA sequencing data from the four datasets analyzed in this study (A) before, and (B) after batch correction using ComBat.