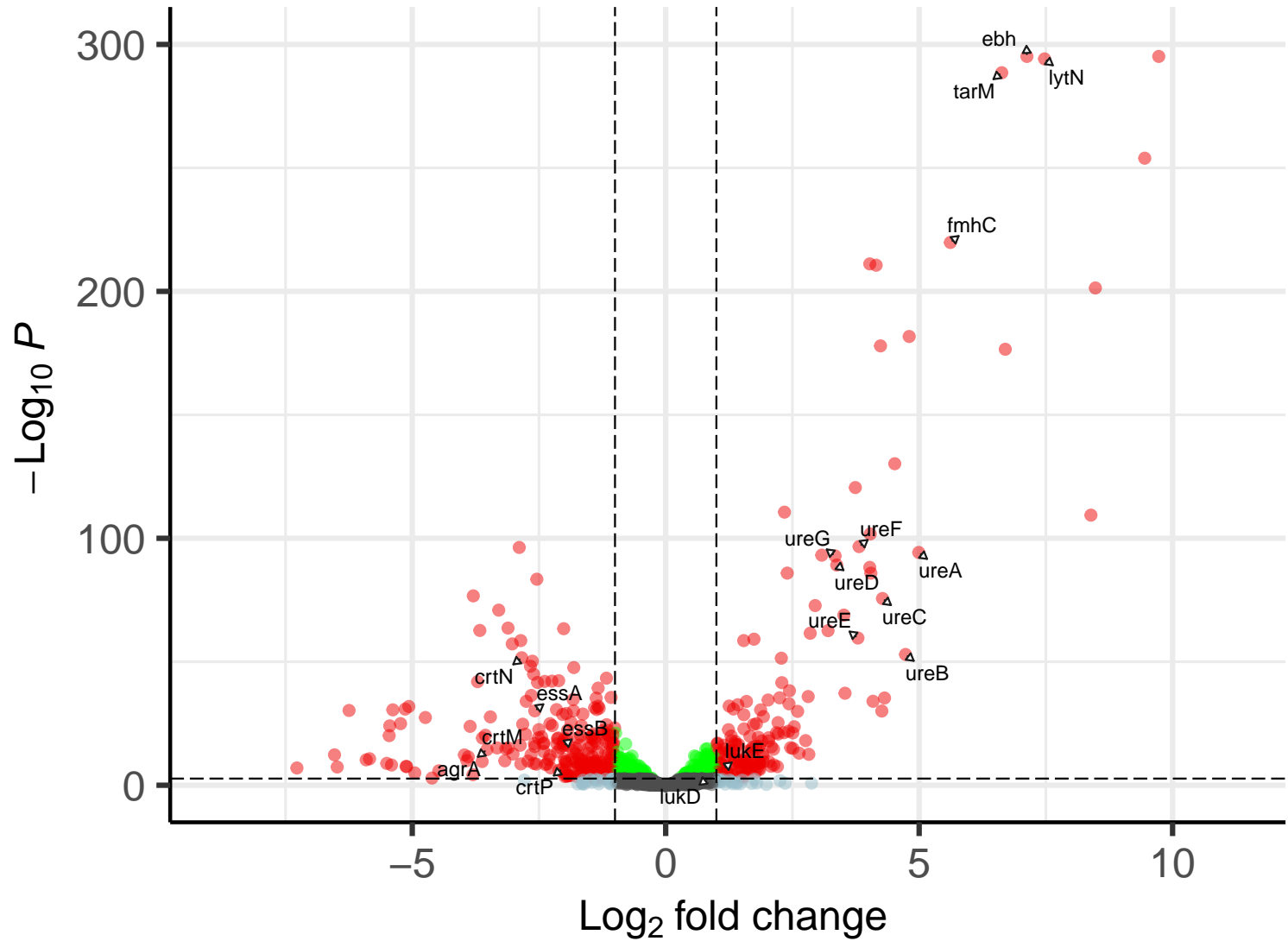


MW2 RNA-seq data

(control vs treated)

● NS ● Log_2 FC ● p-value ● p-value and Log_2 FC



total = 2248 variables