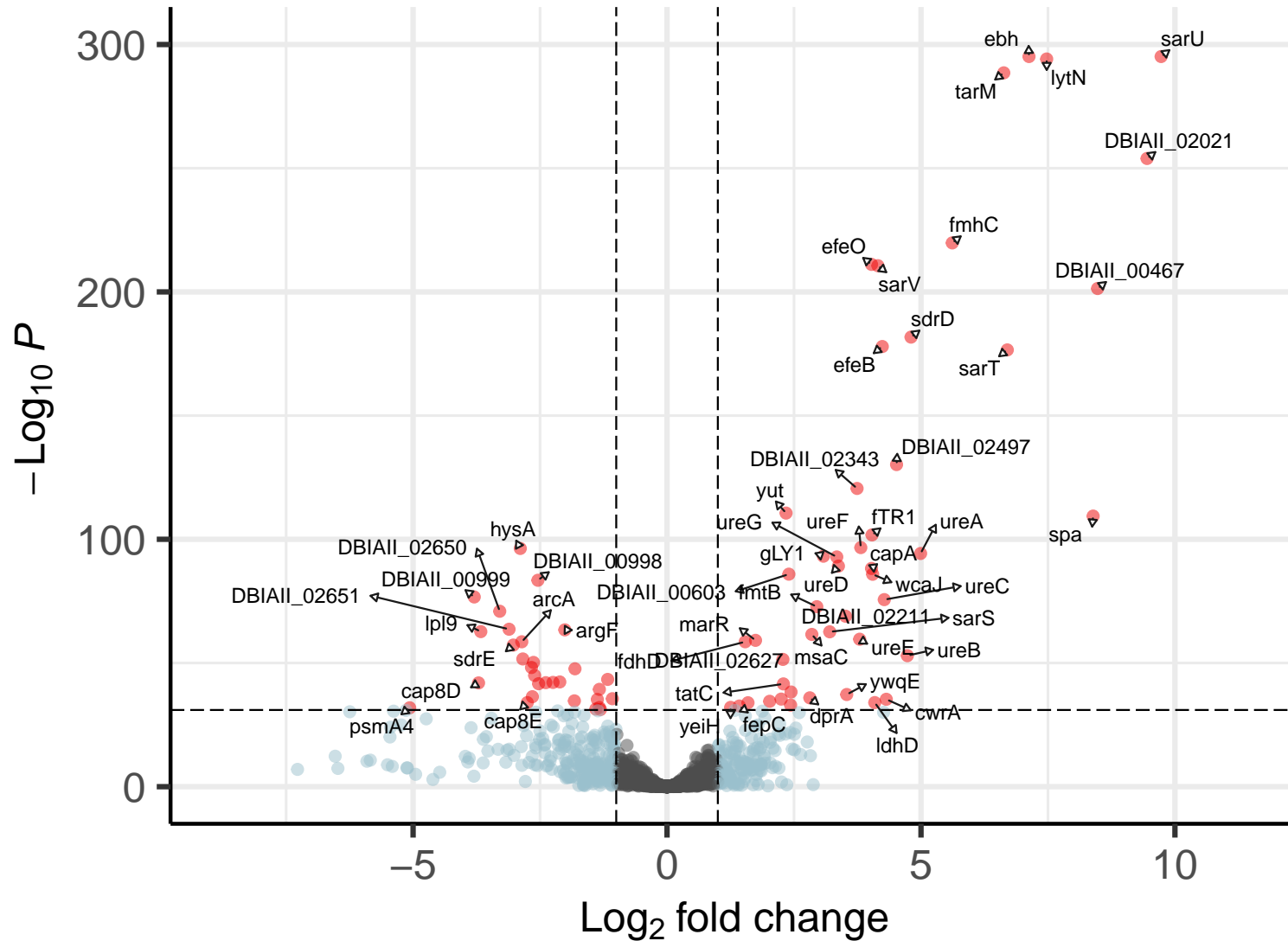


# MW2 RNA-seq data

(control vs treated)

● NS    ● Log<sub>2</sub> FC    ● p-value and log<sub>2</sub> FC



total = 2248 variables