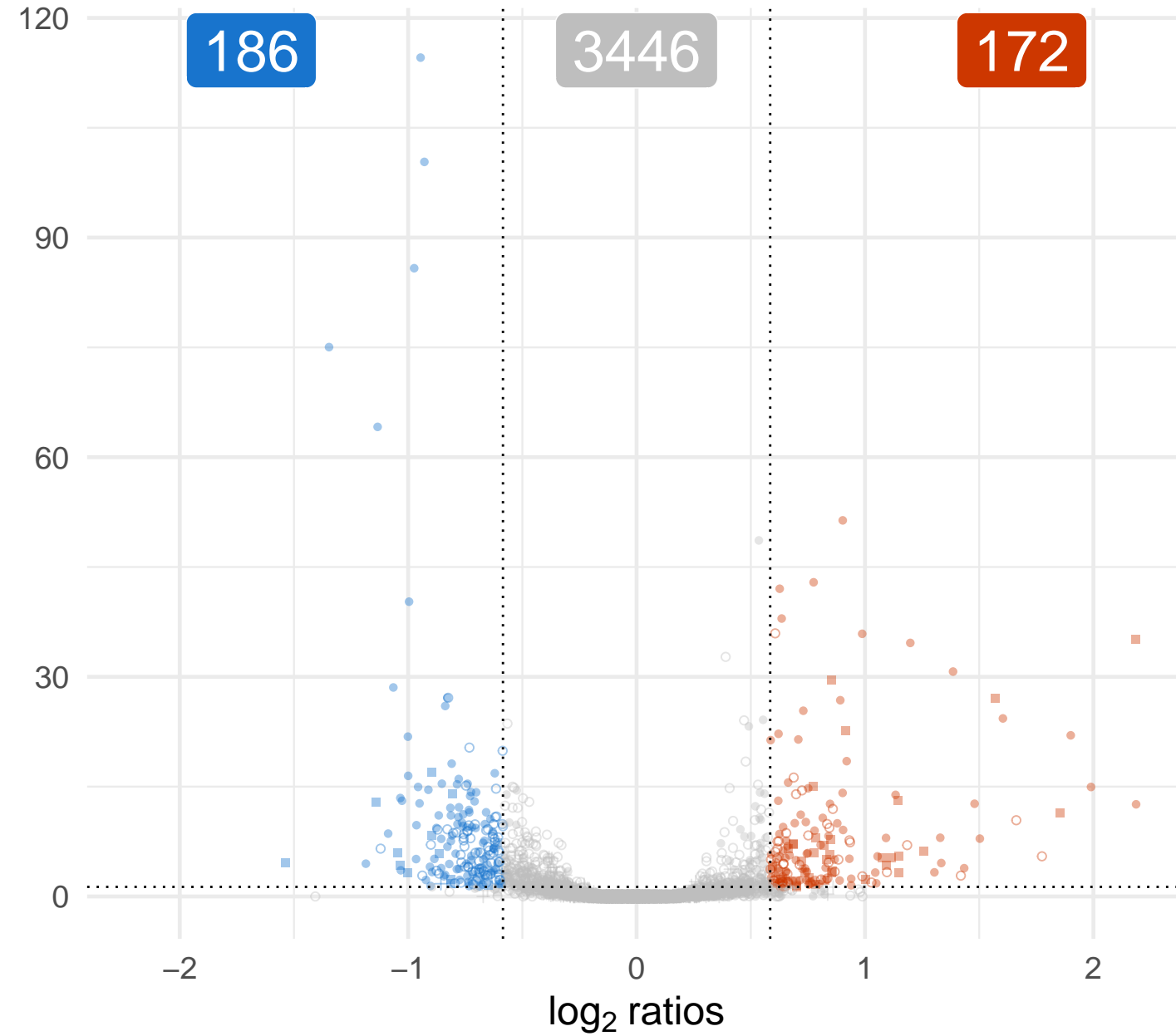


cancer/
control

cutoffs: fold-change=1.5 / adj. p-value=0.05

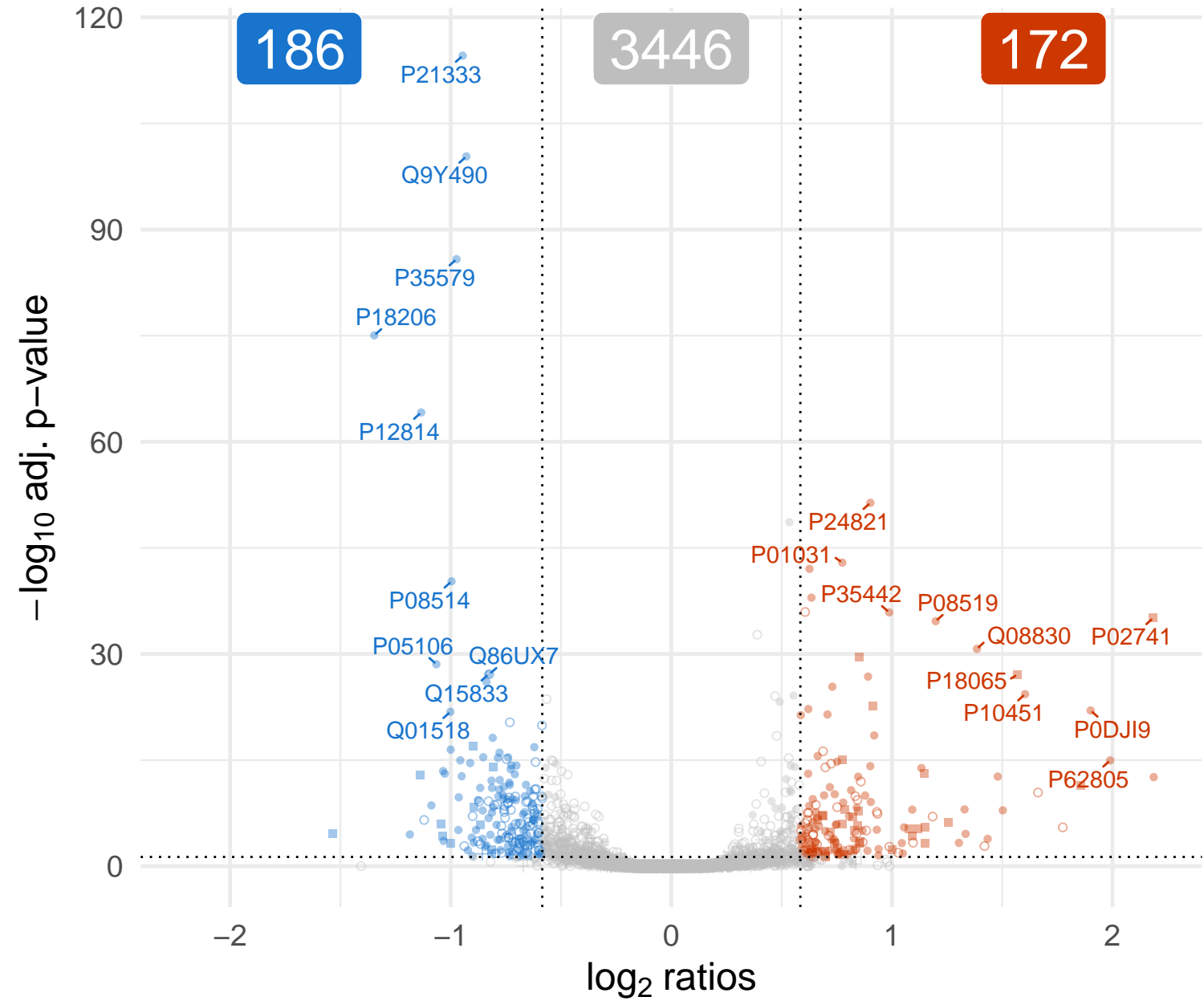


Cohen's eff. size magnitude: + negligible ○ small ● medium ■ large

protein data filtered for equal or more 2 peptides

cancer/
control

cutoffs: fold-change=1.5 / adj. p-value=0.05



Cohen's eff. size magnitude: + negligible ○ small ● medium ■ large

protein data filtered for equal or more 2 peptides