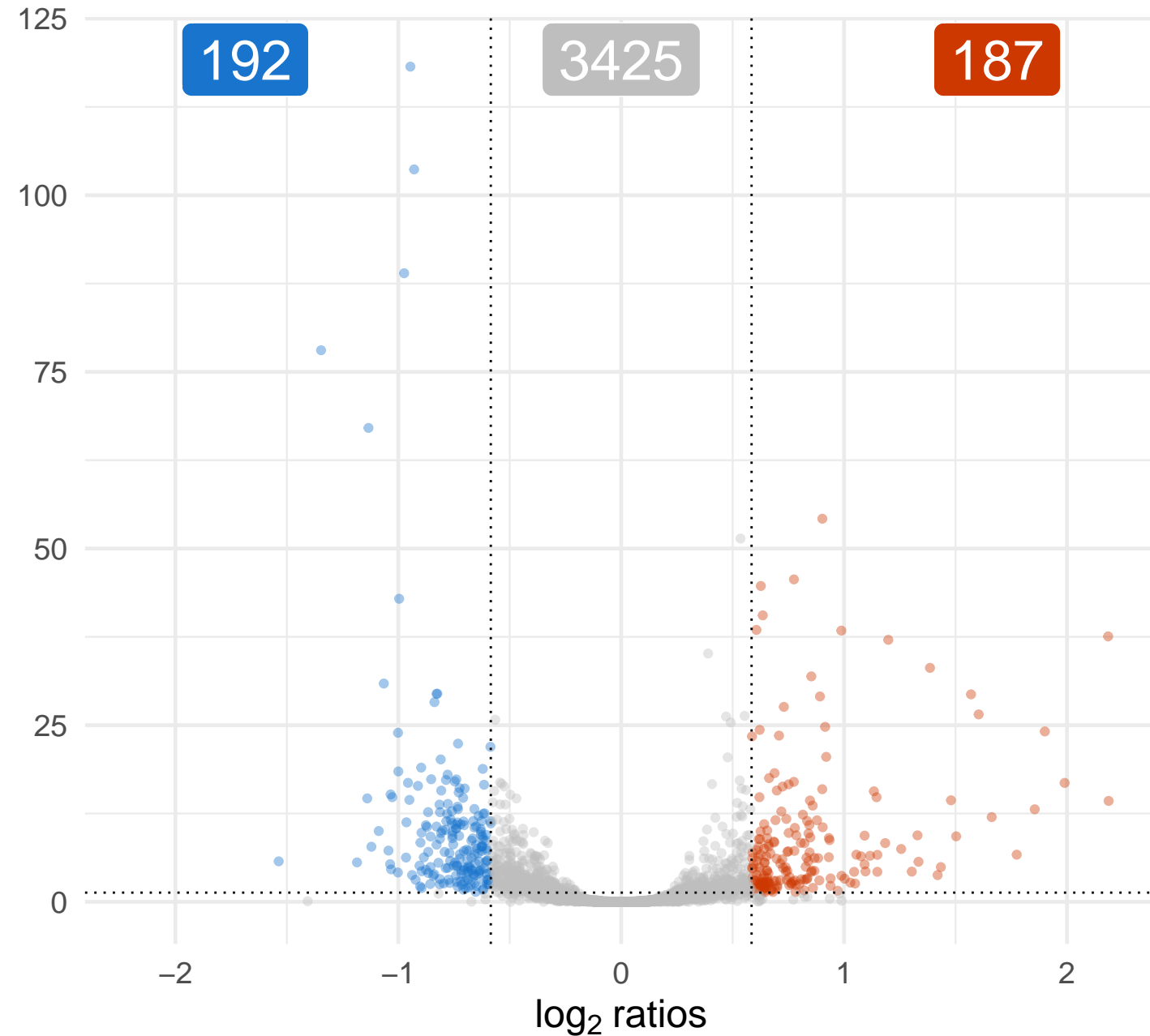


cancer/  
control

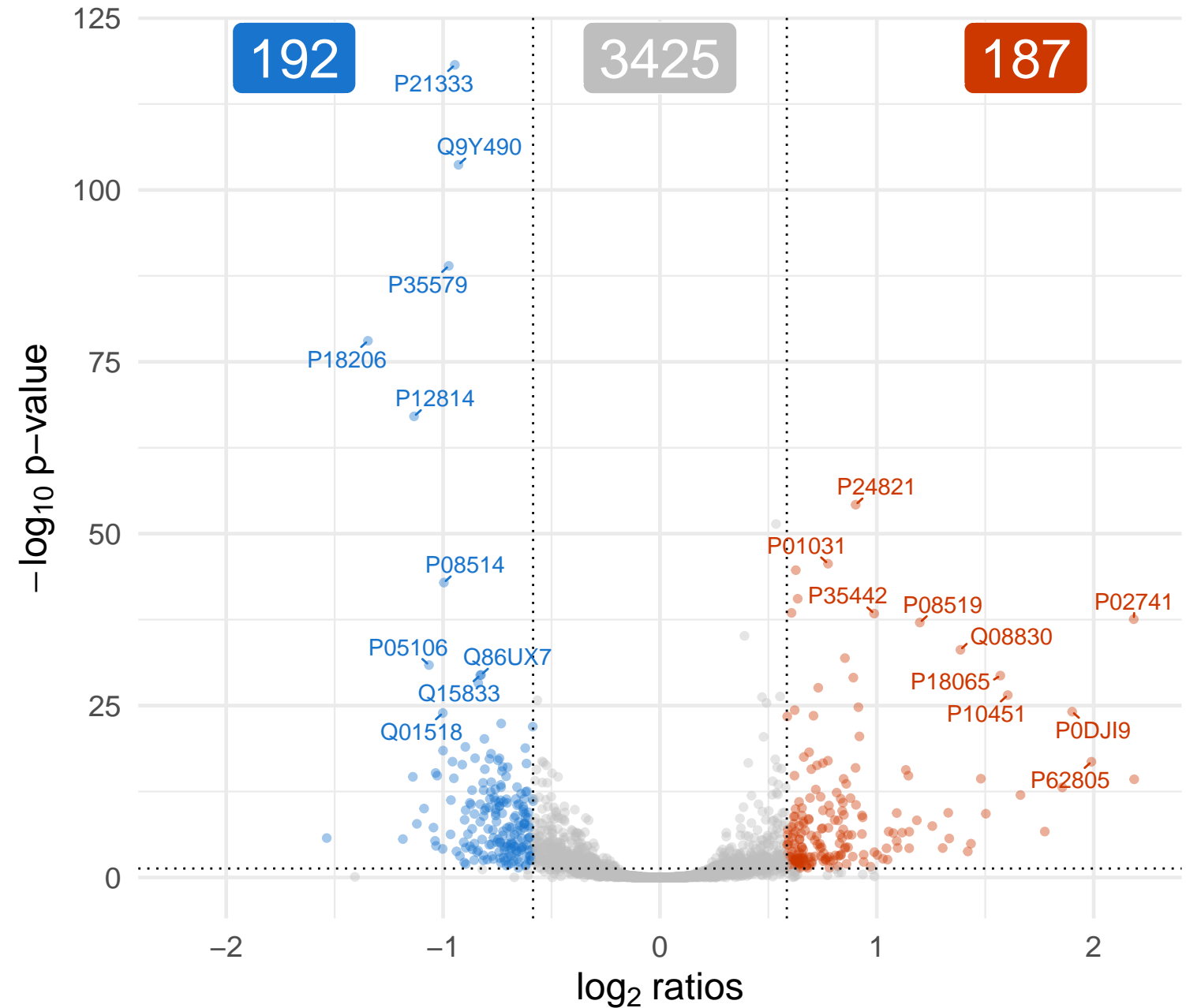
cutoffs: fold-change=1.5 / p-value=0.05



protein data filtered for equal or more 2 peptides

cancer/  
control

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



protein data filtered for equal or more 2 peptides