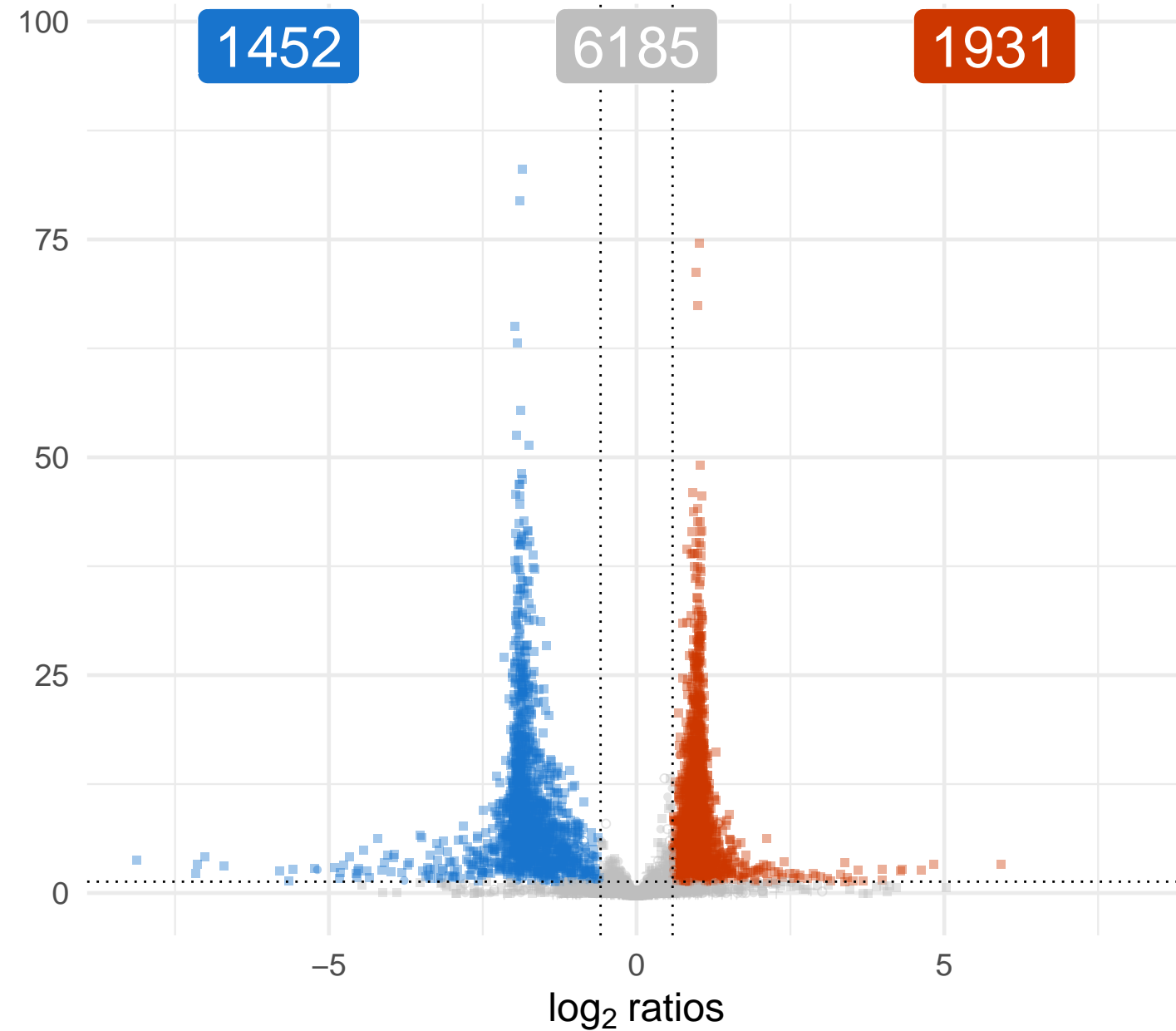


HYE mix A/
HYE mix B

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

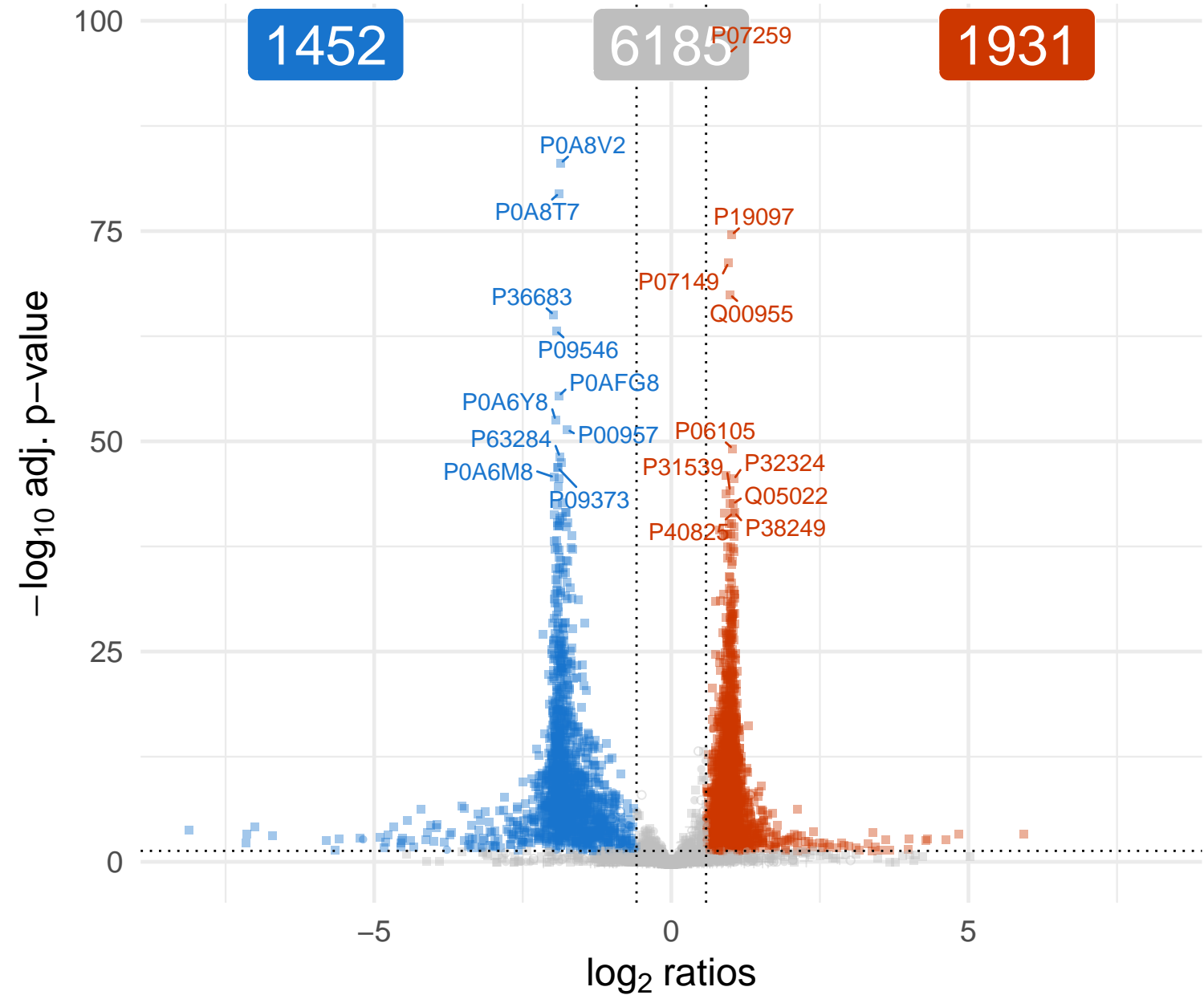


Cohen's eff. size magnitude: + negligible ○ small ● medium ■ large

protein data filtered for equal or more 2 peptides

HYE mix A/
HYE mix B

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