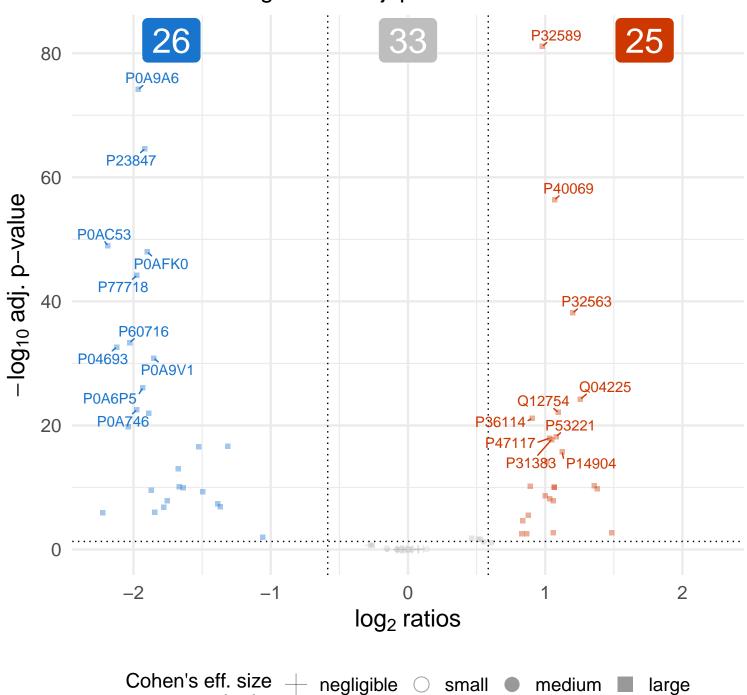


-log₁₀ adj. p-value

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



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



magnitude:

protein data filtered for equal or more 2 peptides

protein data filtered for equal or more 2 peptides