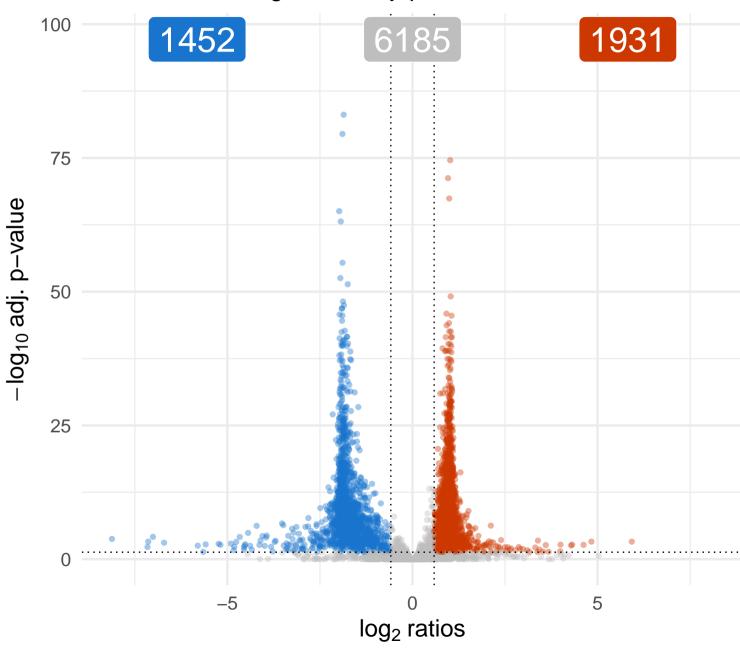
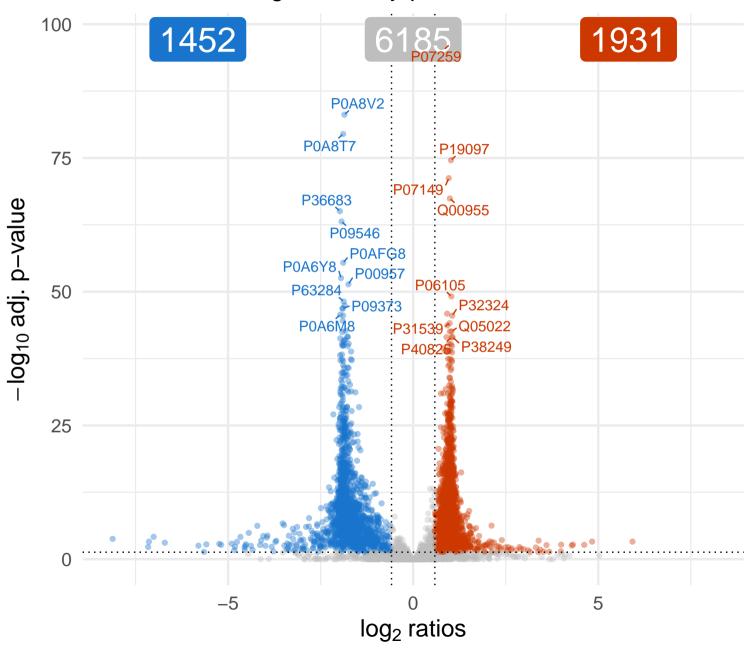


cutoffs: fold-change=1.5 / adj. p-value=0.05



HYE mix A/ HYE mix B

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



protein data filtered for equal or more 2 peptides

protein data filtered for equal or more 2 peptides