HYE mix A/
HYE mix B

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

1452 6185 1931

100 75 50 25 0

$-\log_{10}$ adj. p−value

−5 0 5

$\log_2$ ratios

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

1452 6185 1931

P07259
P0A8V2
P0A8T7
P19097
P07149
P36683
Q00955
P09546
P0AFG8
P0A6Y8
P63284
P00957
P06105
P0A6M8
P31539
P32324
P09373
Q05022
P40825
P38249

100 75 50 25 0

$-\log_{10}$ adj. p−value

−5 0 5

$\log_2$ ratios

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

protein data filtered for equal or more 2 peptides