HS   CTRL

A

Counts

1000   2000   3000
Mass

Counts

1000   2000   3000
Mass

B

HS   CTRL

Counts

2   3   4   5   ≥6
Z

Counts

2   3   4   ≥6
Z

C

HS   CTRL

Counts

-5   0   5
Mass Error [ppm]

Counts

-5   0   5
Mass Error [ppm]

Figure S1: A) Precursor molecular mass relative to peptides identified in the analysis of HS and CTRL. B) Precursor charges related to peptides identified in the analysis of HS and CTRL. C) Mass error (ppm) related to peptides identified in the analysis of HS and CTRL. Panels were generated using the Perseus software.