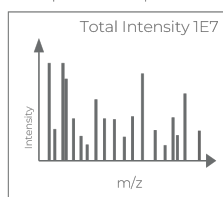
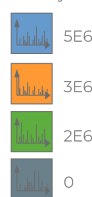
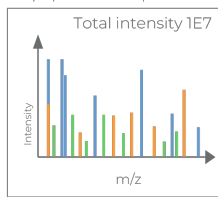
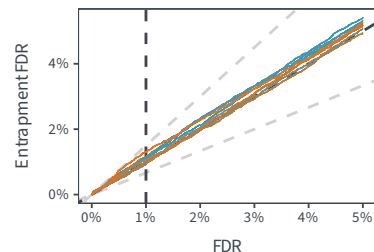


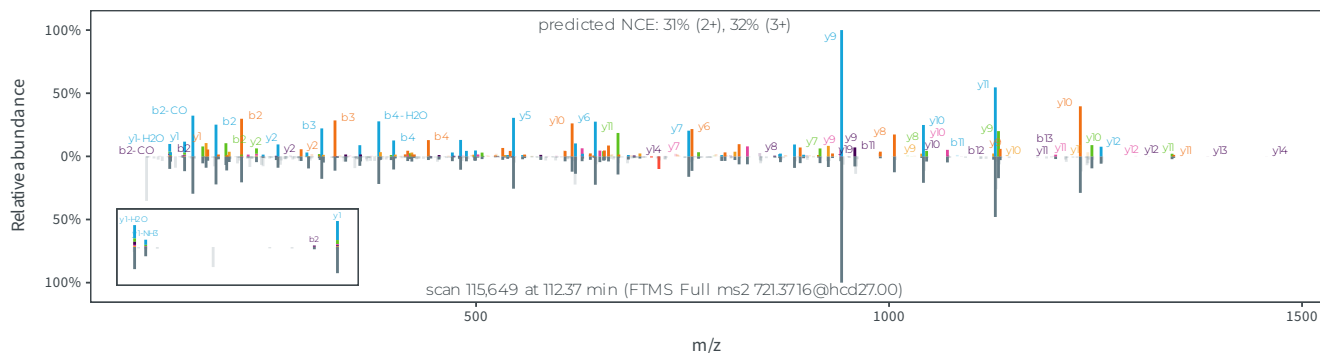
A

(Chimeric) DDA, DIA or PRM
Peptide MS2 SpectrumPrediction of candidates
from full isolation windowSpectral
deconvolutionNon-negative
regularized
linear regressionDistributed
intensityResulting deconvoluted
peptide MS2 spectrumB Isolation
width
[Th]1
3
6
8
10
12
15
20

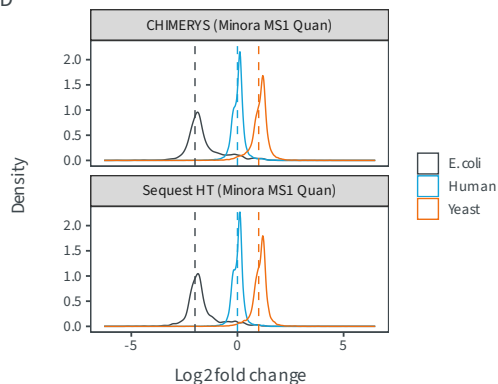
C

— measured — ALQSVGQNGEVLK (2+) — VPLSEDFSWSK (2+) — GSPMEISLPALSK (2+)

— precursor — TNLMNYLPQNMTQDELK (3+) — SINILDIYSTSK (2+) — GSNTIASAAADKIPGLGVFQK (3+)



D



E

■ PSM FDR ■ Precursor FDR ■ Peptide group FDR

