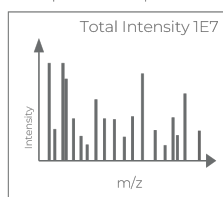
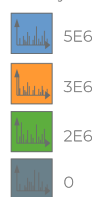
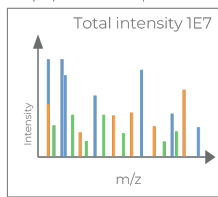
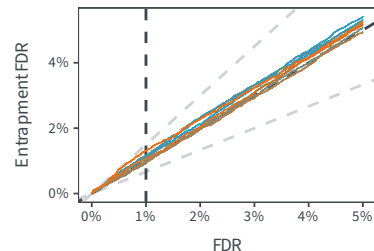
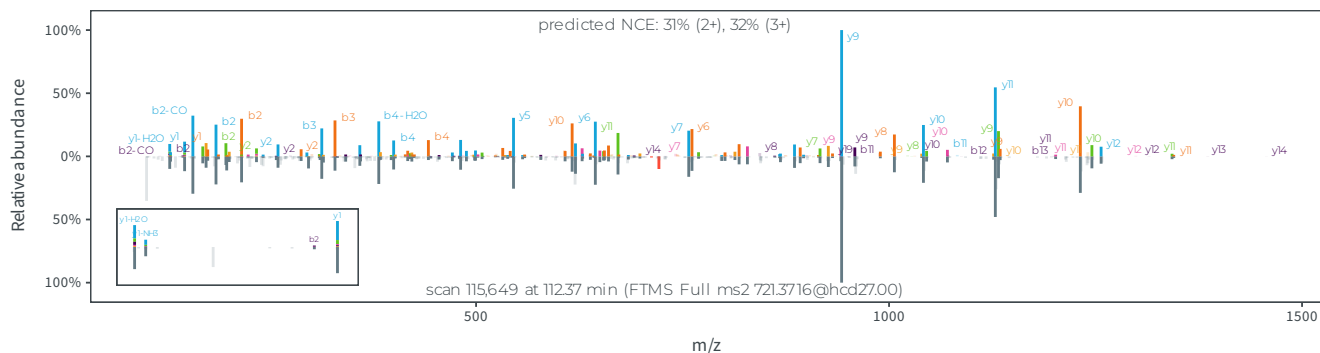


A

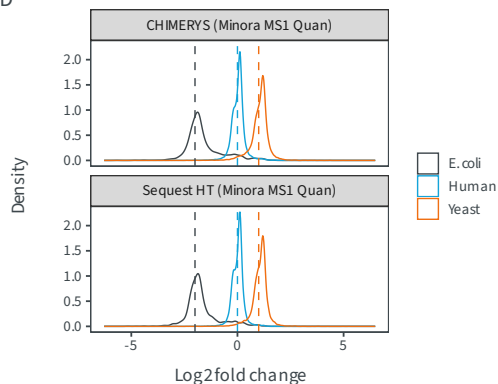
(Chimeric) DDA, DIA or PRM  
Peptide MS2 SpectrumPrediction of candidates  
from full isolation windowSpectral  
deconvolution  
Non-negative  
regularized  
linear regressionDistributed  
intensityResulting deconvoluted  
peptide MS2 spectrumB Isolation  
width  
[Th]1  
3  
6  
8  
10  
12  
15  
20

C

— measured — ALQSVGQNGEVLK (2+) — VPLSEDFSWSK (2+) — GSPMEISLPALSK (2+)  
 — precursor — TNLMNYLPQNMTQDELK (3+) — SINILDIYSTSK (2+) — GSNTIASAAADKIPGLGLGVFQK (3+)



D



E

■ PSM FDR ■ PCM FDR ■ Peptide group FDR

