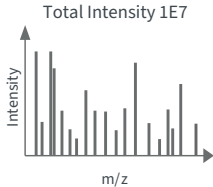


(Chimeric) DDA, DIA or PRM
Peptide MS2 Spectrum



Prediction of candidates
from full isolation window



Spectral
deconvolution

Non-negative
regularized
linear regression

Distributed
intensity



5E6

3E6

2E6

0

Resulting deconvoluted
peptide MS2 spectrum

