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♠ Top-down proteomics data analysis on the NRTDP TDPortal

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This method describes how to analyze data on TD Portal, a top-down proteomics search tool from the National Resource for Translational and Developmental Proteomics.

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43300

In steps of

Overall protocol for top-down LC-MS/MS of human lung tissue Overall protocol for top-down LC-MS/MS of human heart tissue

1 Top-down proteomics data was searched on the TD Portal (portal.nrtdp.northwestern.edu)



- 2 Raw files for analysis were selected for search using Top Down Search v. 4.0.
- 3 Search parameters:

Organism: Homo Sapiens (Human) with Curated Histones: (Taxon 9606 - June 2020)

Use Empirical P-Score: FALSE

Filter by FDR: TRUE

Create SAS Input Sheet for Quant: FALSE Precursor Resolution: High Resolution

Fragmentation Type: Auto Code Set: Standard 4.0.0

Include ProSight Error Tolerant Search: FALSE