# NIST Peptide Mass Spectral Libraries: *Freely Available*



## LC-ESI-MS/MS Libraries of Annotated Spectra

- 4.4 million spectra
- 912,182 tryptic peptides

# **Tryptic Peptide Libraries**

- Human (from CPTAC and ProteoeXchange, and ProteomeTools)
  - -Label-free, iTRAQ and TMT
  - -Hair label-free (NIST, forensics use)
  - -Phosphopeptides
  - -Synthetic peptides
- Chinese hamster (from Johns Hopkins University and NIST)

#### **NIST Human Peptide Mass Spectral Library Statistics**

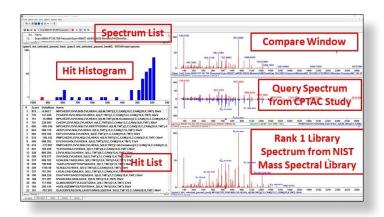
 Spectra
 4 260 741
 Proteins
 33 019

Size 33 GB Variants 17 GVPs (hair)

Peptides 837 673 Modifications 8



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#### How to use libraries

- Formats: (1) ASCII text format (MSP) and
   (2) MS Search binary format spectra
- Library searching with major proteomics workflows, including Mascot (Matrix Science) and Proteome Discoverer (Thermo Fisher Scientific)
- Compatible with Skyline
- Individual spectrum search with the full-featured user-interface program: NIST MS Search
- Batch peptide identification with the *NIST MS PepSearch* software

## **Library Downloads**

Peptide.nist.gov

Web <u>chemdata.nist.gov</u>

Email <u>massspec@nist.gov</u>



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