Resource Summary Report

Generated by NIF on May 19, 2025

Kojak

RRID:SCR_021028
Type: Tool

Proper Citation

Kojak (RRID:SCR_021028)

Resource Information

URL: http://www.kojak-ms.org/

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Description: Software tool for identification of cross-linked peptides from mass spectra. Used for analysis of chemically cross-linked protein complexes. Used to analyze both novel and existing data sets.

Resource Type: software application, software resource, data analysis software, data processing software

Defining Citation: PMID:25812159

Keywords: Mass spectra, cross-linked peptides identification, protein complexes analysis, novel data analysis, existing data analysis

Funding: National Science Foundation MRI grant 0923536;
NIGMS P50 GM076547;
NIGMS P50 GM08722150;
NCRR S10 RR027584;
NIGMS P41 GM103533

Availability: Free, Available for download, Freely available

Resource Name: Kojak

Resource ID: SCR_021028

Record Creation Time: 20220129T080353+0000

Record Last Update: 20250519T204230+0000

Ratings and Alerts

No rating or validation information has been found for Kojak.

No alerts have been found for Kojak.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF.

Dahiya V, et al. (2022) The switch from client holding to folding in the Hsp70/Hsp90 chaperone machineries is regulated by a direct interplay between co-chaperones. Molecular cell, 82(8), 1543.

Brilot AF, et al. (2021) CM1-driven assembly and activation of yeast ?-tubulin small complex underlies microtubule nucleation. eLife, 10.

Popov KI, et al. (2019) Insight into the Structure of the "Unstructured" Tau Protein. Structure (London, England : 1993), 27(11), 1710.