

Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

mzStudio

RRID:SCR_017088

Type: Tool

Proper Citation

mzStudio (RRID:SCR_017088)

Resource Information

URL: <https://github.com/BlaisProteomics/mzStudio>

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Description: Software tool for proteomics data analysis, visualization, and notebook application. Dynamic digital canvas for user driven interrogation of mass spectrometry data. Operating system Unix/Linux, Windows.

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

Defining Citation: [PMID:28763045](#)

Keywords: proteomic, data, analysis, visualization, notebooking, mass, spectrometry, modification, gas, phase, fragmentation, behavior

Funding: NCI CA188881;
NCI CA178860;
NCI CA042368;
Dana-Farber Strategic Research Initiative ;
Barr Program in Basic Research ;
Honorable Tina Brozman Foundation for Ovarian Cancer Research ;
Michael J. Fox Foundation

Availability: Free, Available for download, Freely available

Resource Name: mzStudio

Resource ID: SCR_017088

Alternate IDs: OMICS_26946

Alternate URLs: <https://omictools.com/mzstudio-tool>

License: GNU GPL v3

Record Creation Time: 20220129T080333+0000

Record Last Update: 20260731T043016+0000

Ratings and Alerts

No rating or validation information has been found for mzStudio.

No alerts have been found for mzStudio.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Chen H, et al. (2023) A small molecule exerts selective antiviral activity by targeting the human cytomegalovirus nuclear egress complex. PLoS pathogens, 19(11), e1011781.