

Resource Summary Report

Generated by T1D on Sep 5, 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 analysis software, data processing software, data visualization software, software application, software resource

Defining Citation: [PMID:28763045](https://pubmed.ncbi.nlm.nih.gov/28763045/)

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

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

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: 20260903T235414+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 [T1D](#).

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.