# Resource Summary Report

Generated by SPARC SAWG on Aug 9, 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:** software resource, software application, data analysis software, data visualization software, data processing software

**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:** 20260808T190043+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 SPARC SAWG .

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.