

# Resource Summary Report

Generated by [SPARC SAWG](#) on Jul 28, 2026

## [jmzML](#)

RRID:SCR\_001119

Type: Tool

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### Proper Citation

jmzML (RRID:SCR\_001119)

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### Resource Information

**URL:** <https://code.google.com/p/jmzml/>

**Proper Citation:** jmzML (RRID:SCR\_001119)

**Description:** A Java application programming interface (API) for the Proteomics Standards Initiative mzML data standard.

**Resource Type:** software resource

**Defining Citation:** [PMID:20127693](#)

**Keywords:** standalone software, mac os x, unix/linux, windows, java, bio.tools

**Availability:** Free, Available for download, Freely available

**Resource Name:** jmzML

**Resource ID:** SCR\_001119

**Alternate IDs:** biotools:jmzml, OMICS\_03340

**Alternate URLs:** <https://bio.tools/jmzml>

**Record Creation Time:** 20220129T080205+0000

**Record Last Update:** 20260725T190455+0000

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### Ratings and Alerts

No rating or validation information has been found for jmzML.

No alerts have been found for jmzML.

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### Data and Source Information

**Source:** [SciCrunch Registry](#)

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## 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](#) .

Rocca-Serra P, et al. (2016) Data standards can boost metabolomics research, and if there is a will, there is a way. Metabolomics : Official journal of the Metabolomic Society, 12, 14.