

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

## [jmzTab](#)

RRID:SCR\_003481

Type: Tool

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### Proper Citation

jmzTab (RRID:SCR\_003481)

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### Resource Information

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

**Proper Citation:** jmzTab (RRID:SCR\_003481)

**Description:** A Java interface to the mzTab data exchange format for reporting a summary of proteomics results.

**Resource Type:** software resource

**Defining Citation:** [PMID:24659499](#)

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

**Availability:** Apache License, v2

**Resource Name:** jmzTab

**Resource ID:** SCR\_003481

**Alternate IDs:** OMICS\_04542, biotools:jmztab

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

**Record Creation Time:** 20220129T080219+0000

**Record Last Update:** 20260905T132504+0000

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### Ratings and Alerts

No rating or validation information has been found for jmzTab.

No alerts have been found for jmzTab.

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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 [Kravitz](#) .

Qi D, et al. (2015) The mzqLibrary--An open source Java library supporting the HUPO-PSI quantitative proteomics standard. Proteomics, 15(18), 3152.