

# Resource Summary Report

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## Mzkit

RRID:SCR\_023936

Type: Tool

### Proper Citation

Mzkit (RRID:SCR\_023936)

### Resource Information

**URL:** <https://mzkit.org/>

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**Description:** Open source raw data file toolkit for mass spectrometry data analysis, provided by the BioNovoGene corporation. The features of mzkit includes: raw data file content viewer(XIC/TIC/Mass spectrum plot/MS-Imaging), build molecule network, formula de-novo search, de-novo annotation of the unknown metabolite features, MALDI single cell metabolomics data analysis, pathological slide viewer and targeted data quantification.

**Synonyms:** BioNovoGene Mzkit

**Resource Type:** software toolkit, software resource

**Keywords:** BioNovoGene, processing NMR, MALDI MSI, MALDI single cell, Raman Spectroscopy, LC-MS and GC-MS raw data, chemoinformatics data analysis and data visualization,

**Funding:**

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

**Resource Name:** Mzkit

**Resource ID:** SCR\_023936

**Alternate URLs:** 10.5281/zenodo.8076924

**License:** MIT License

**Record Creation Time:** 20230816T050225+0000

**Record Last Update:** 20250329T061647+0000

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## Ratings and Alerts

No rating or validation information has been found for Mzkit.

No alerts have been found for Mzkit.

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## Data and Source Information

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

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.