

Resource Summary Report

Generated by [NIF](#) on Aug 9, 2026

SashimipLOT

RRID:SCR_016861

Type: Tool

Proper Citation

SashimipLOT (RRID:SCR_016861)

Resource Information

URL: <https://github.com/yarden/MISO/blob/fastmiso/docs/source/sashimi.rst>

Proper Citation: SashimipLOT (RRID:SCR_016861)

Description: Software tool for quantitative visualization of aligned RNA-Seq reads that enables quantitative comparison of exon usage across samples or experimental conditions.

Abbreviations: sashimipLOT

Synonyms: sashimi_plot

Resource Type: software resource, software application, data visualization software, data processing software

Defining Citation: [PMID:25617416](#), [DOI:10.1093/bioinformatics/btv034](#)

Keywords: quantitative, visualization, aligned, RNA-Seq, read, data, compare, exon, usage, sample, experiment, condition, MISO

Funding: NCI R01 CA157304;
Starr Cancer Consortium ;
NIGMS R01 GM096193;
NSF IIS 1149662;
Alfred P. Sloan research fellowship ;
NIGMS R01 GM085319;
NCI U01 CA184897;
NHGRI R01 HG002439

Availability: Free, Available for download, Freely available

Resource Name: SashimipLOT

Resource ID: SCR_016861

Alternate URLs: <http://miso.readthedocs.org/en/fastmiso/sashimi.html>

Record Creation Time: 20220129T080332+0000

Record Last Update: 20260808T190036+0000

Ratings and Alerts

No rating or validation information has been found for Sashimiplot.

No alerts have been found for Sashimiplot.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.