Resource Summary Report

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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: 20250519T203939+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 found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET.

Vernia S, et al. (2016) An alternative splicing program promotes adipose tissue thermogenesis. eLife, 5.