

# Resource Summary Report

Generated by [kravitz2](#) on Aug 1, 2026

## SplicePlot

RRID:SCR\_001172

Type: Tool

### Proper Citation

SplicePlot (RRID:SCR\_001172)

### Resource Information

**URL:** <http://montgomerylab.stanford.edu/spliceplot/index.html>

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**Description:** A software tool for visualizing alternative splicing and the effects of splicing quantitative trait loci (sQTLs) from RNA-seq data. It provides a simple command line interface for drawing sashimi plots, hive plots, and structure plots of alternative splicing events from .bam, .gtf, and .vcf files.

**Abbreviations:** SplicePlot

**Synonyms:** SplicePlot: a tool for visualizing alternative splicing

**Resource Type:** software resource

**Defining Citation:** [PMID:24363378](#)

**Keywords:** visualization, alternative splicing, splicing quantitative trait loci, quantitative trait loci, rna-seq, hive plot, structure plot, sashimi plot, python, linux, mac os, bio.tools

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** SplicePlot

**Resource ID:** SCR\_001172

**Alternate IDs:** OMICS\_02185, biotools:spliceplot

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

**Record Creation Time:** 20220129T080206+0000

**Record Last Update:** 20260725T190456+0000

### Ratings and Alerts

No rating or validation information has been found for SplicePlot.

No alerts have been found for SplicePlot.

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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.