

Resource Summary Report

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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: 20260815T182203+0000

Ratings and Alerts

No rating or validation information has been found for SplicePlot.

No alerts have been found for SplicePlot.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.