

Resource Summary Report

Generated by [nidm-terms](#) on Sep 21, 2026

Supersplat

RRID:SCR_009826

Type: Tool

Proper Citation

Supersplat (RRID:SCR_009826)

Resource Information

URL: <http://mocklerlab.org/tools/1>

Proper Citation: Supersplat (RRID:SCR_009826)

Description: An application for discovering potential splice junctions in high throughput sequencing (HTS) data.

Abbreviations: Supersplat

Resource Type: software resource

Keywords: bio.tools

Resource Name: Supersplat

Resource ID: SCR_009826

Alternate IDs: OMICS_01256, biotools:supersplat

Alternate URLs: <https://bio.tools/supersplat>

Record Creation Time: 20220129T080255+0000

Record Last Update: 20260919T195151+0000

Ratings and Alerts

No rating or validation information has been found for Supersplat.

No alerts have been found for Supersplat.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Thangam M, et al. (2015) CRCDA--Comprehensive resources for cancer NGS data analysis. Database : the journal of biological databases and curation, 2015.

Uyar B, et al. (2012) RNA-seq analysis of the C. briggsae transcriptome. Genome research, 22(8), 1567.