# Resource Summary Report

Generated by NIF on Sep 1, 2026

## SV-plaudit

RRID:SCR_016285
Type: Tool

### Proper Citation

SV-plaudit (RRID:SCR_016285)

### Resource Information

**URL:** https://github.com/jbelyeu/SV-plaudit

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**Description:** Software for rapidly curating structural variant (SVs) predictions. SV-plaudit provides a pipeline for creating image views of genomic intervals, automatically storing them in the cloud, deploying a website to view/score them, and retrieving scores for analysis.

**Resource Type:** data analysis software, data processing software, image analysis software, software application, software resource

**Keywords:** genomics, structural, variants, visualization, manual, curation, prediction, image, alignment

**Funding:** NCI U24 CA209999;
NHGRI K99 HG009532;
NHGRI R01 HG006693;
NIGMS R01 GM124355

**Availability:** Free, Available for download

**Resource Name:** SV-plaudit

**Resource ID:** SCR_016285

**License:** MIT License

**Record Creation Time:** 20220129T080329+0000

**Record Last Update:** 20260829T182510+0000

### Ratings and Alerts

No rating or validation information has been found for SV-plaudit.

No alerts have been found for SV-plaudit.

## 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 NIF .

Bertolotti AC, et al. (2020) The structural variation landscape in 492 Atlantic salmon genomes. Nature communications, 11(1), 5176.

Belyeu JR, et al. (2018) SV-plaudit: A cloud-based framework for manually curating thousands of structural variants. GigaScience, 7(7).