

# Resource Summary Report

Generated by SPARC SAWG on Sep 17, 2026

## BSVF

RRID:SCR\_015727

Type: Tool

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### Proper Citation

BSVF (RRID:SCR\_015727)

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### Resource Information

**URL:** <https://github.com/BGI-SZ/BSVF>

**Proper Citation:** BSVF (RRID:SCR\_015727)

**Description:** Software code for bisulfite sequencing virus integration. This finder is for directional libraries only and does not support PBAT and indirectional libraries.

**Abbreviations:** BSVF

**Synonyms:** BSVF: Bisulfite Sequencing Virus integration Finder, Bisulfite Sequencing Virus integration Finder

**Resource Type:** data analysis software, data processing software, sequence analysis software, software application, software resource, source code

**Keywords:** virus integration, sequencing analysis, virus assembly, integration, bisulfite, bio.tools

**Availability:** Open source

**Resource Name:** BSVF

**Resource ID:** SCR\_015727

**Alternate IDs:** biotools:bs-virus-finder

**Alternate URLs:** <https://bio.tools/bs-virus-finder>

**License:** LGPLv3

**License URLs:** <https://github.com/BGI-SZ/BSVF/blob/master/LICENSE>

**Record Creation Time:** 20220129T080327+0000

**Record Last Update:** 20260912T195829+0000

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## Ratings and Alerts

No rating or validation information has been found for BSVF.

No alerts have been found for BSVF.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [SPARC SAWG](#) .

Gao S, et al. (2018) BS-virus-finder: virus integration calling using bisulfite sequencing data. GigaScience, 7(1), 1.