

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

BSVF

RRID:SCR_015727

Type: Tool

Proper Citation

BSVF (RRID:SCR_015727)

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: 20260905T132758+0000

Ratings and Alerts

No rating or validation information has been found for BSVF.

No alerts have been found for BSVF.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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