

# Resource Summary Report

Generated by [dkNET](#) on Aug 5, 2026

## SNPSVM

RRID:SCR\_000028

Type: Tool

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### Proper Citation

SNPSVM (RRID:SCR\_000028)

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### Resource Information

**URL:** <https://github.com/brendanofallon/SNPSVM/>

**Proper Citation:** SNPSVM (RRID:SCR\_000028)

**Description:** A support vector machine for calling variants from next-gen sequencing data. It takes as input a BAM-formatted alignment of sequencing reads, and emits a VCF formatted file describing where all the SNPs (single nucleotide polymorphisms) are.

**Resource Type:** software resource

**Defining Citation:** [PMID:23620357](#)

**Keywords:** standalone software

**Availability:** Free, Available for download, Freely available

**Resource Name:** SNPSVM

**Resource ID:** SCR\_000028

**Alternate IDs:** OMICS\_03838

**Record Creation Time:** 20220129T080159+0000

**Record Last Update:** 20260801T190107+0000

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### Ratings and Alerts

No rating or validation information has been found for SNPSVM.

No alerts have been found for SNPSVM.

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### Data and Source Information

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

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.