

Resource Summary Report

Generated by [NIF](#) on Jul 30, 2026

[vcfanno](#)

RRID:SCR_024372

Type: Tool

Proper Citation

vcfanno (RRID:SCR_024372)

Resource Information

URL: <https://github.com/brentp/vcfanno>

Proper Citation: vcfanno (RRID:SCR_024372)

Description: Software tool for flexible annotation of genetic variants.Extracts and summarizes attributes from multiple annotation files and integrates annotations within INFO column of the original VCF file.

Resource Type: software application, software resource

Defining Citation: [PMID:27250555](#)

Keywords: annotation of genetic variants, extracts and summarizes attributes, multiple annotation files, integrates annotations, VCF file,

Availability: Free, Available for download, Freely available,

Resource Name: vcfanno

Resource ID: SCR_024372

Alternate IDs: OMICS_11863

Alternate URLs: <https://sources.debian.org/src/vcfanno/>

License: MIT License

Record Creation Time: 20230830T050217+0000

Record Last Update: 20260727T040824+0000

Ratings and Alerts

No rating or validation information has been found for vcfanno.

No alerts have been found for vcfanno.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.