

Resource Summary Report

Generated by [RRID](#) on Aug 10, 2026

DivBrowse

RRID:SCR_022780

Type: Tool

Proper Citation

DivBrowse (RRID:SCR_022780)

Resource Information

URL: <https://divbrowse.ipk-gatersleben.de/>

Proper Citation: DivBrowse (RRID:SCR_022780)

Description: Web tool for interactive visualization and analysis of diversity of genomic variants. Used for interactive exploration and analysis of very large SNP matrices based on VCF files.

Resource Type: web application, software resource

Keywords: interactive visualization and analysis, diversity of genomic variants, large SNP matrices, VCF files

Availability: Free, Freely available

Resource Name: DivBrowse

Resource ID: SCR_022780

Alternate URLs: <https://github.com/IPK-BIT/divbrowse>, <https://pypi.org/project/divbrowse/>

License: MIT License

Record Creation Time: 20220923T050148+0000

Record Last Update: 20260808T190245+0000

Ratings and Alerts

No rating or validation information has been found for DivBrowse.

No alerts have been found for DivBrowse.

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 [RRID](#) .

K??nig P, et al. (2022) DivBrowse-interactive visualization and exploratory data analysis of variant call matrices. GigaScience, 12.

, et al. (2022) Correction to: DivBrowse-interactive visualization and exploratory data analysis of variant call matrices. GigaScience, 12.