

Resource Summary Report

Generated by [RRID](#) on Aug 12, 2026

VarFish

RRID:SCR_023710

Type: Tool

Proper Citation

VarFish (RRID:SCR_023710)

Resource Information

URL: <https://www.cubi.bihealth.org/software/varfish/>

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Description: Web application for quality control, filtering, prioritization, analysis, and user based annotation of DNA variant data with focus on rare disease genetics.
Comprehensive DNA variant analysis for diagnostics and research.

Resource Type: data analysis software, data processing software, software resource, software application

Defining Citation: [DOI:10.1093/nar/gkaa241](https://doi.org/10.1093/nar/gkaa241)

Keywords: Genetics, Variant Call Format, Filtering, Quality control,

Funding: Berlin Institute of Health Charité funds ;
Stiftung Charité

Availability: Free, Available for download, Freely available

Resource Name: VarFish

Resource ID: SCR_023710

Alternate URLs: <https://github.com/bihealth/varfish-server>

License: MIT license

Record Creation Time: 20230622T050213+0000

Record Last Update: 20260812T044437+0000

Ratings and Alerts

No rating or validation information has been found for VarFish.

No alerts have been found for VarFish.

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](#) .

Kakar N, et al. (2025) Further evidence of biallelic NAV3 variants associated with recessive neurodevelopmental disorder with dysmorphism, developmental delay, intellectual disability, and behavioral abnormalities. Human genetics, 144(1), 55.

Nieminen M, et al. (2022) SODAR: managing multiomics study data and metadata. GigaScience, 12.