

Resource Summary Report

Generated by [kravitz2](#) on Aug 11, 2026

VariantDB

RRID:SCR_018010

Type: Tool

Proper Citation

VariantDB (RRID:SCR_018010)

Resource Information

URL: <http://143.169.238.105/variantdb/index.php?page=variants>

Proper Citation: VariantDB (RRID:SCR_018010)

Description: Web based interactive annotation and filtering platform that automatically annotates variants with allele frequencies, functional impact, pathogenicity predictions and pathway information. Allows filtering by all annotations, under dominant, recessive or de novo inheritance models. Flexible annotation and filtering portal for next generation sequencing data.

Resource Type: data or information resource, analysis service resource, production service resource, portal, service resource

Defining Citation: [PMID:25352915](#)

Keywords: Annotation, filtering, portal, next generation, sequencing data, variant, allele frequency, functional impact, pathogenicity prediction, pathway information

Funding: Belgian National Fund for Scientific Research-Flanders ;
Fondation Leducq

Availability: Restricted

Resource Name: VariantDB

Resource ID: SCR_018010

Old URLs: <http://www.biomina.be/app/variantdb/>

License: GPLv3

Record Creation Time: 20220129T080338+0000

Record Last Update: 20260811T043622+0000

Ratings and Alerts

No rating or validation information has been found for VariantDB.

No alerts have been found for VariantDB.

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 [kravitz2](#) .

Peeters S, et al. (2020) DNA Methylation Profiling and Genomic Analysis in 20 Children with Short Stature Who Were Born Small for Gestational Age. The Journal of clinical endocrinology and metabolism, 105(12).