# Resource Summary Report

Generated by RRID on Aug 14, 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, service resource, portal

**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:** 20260814T043009+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 RRID .

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).