

Resource Summary Report

Generated by [RRID](#) on Aug 4, 2026

Human Variation DB

RRID:SCR_009014

Type: Tool

Proper Citation

Human Variation DB (RRID:SCR_009014)

Resource Information

URL: http://gwas.biosciencedbc.jp/cgi-bin/hvdb/hv_top.cgi

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Description: A repository database to achieve continuous and intensive management of GWAS data and variation data identified by next generation sequencing (NGS) and data-sharing among researchers. In this database, variations including short/long insertions / deletions and structural variations related to disease susceptibility, virus resistance, and drug response are registered along with statistical genetic results and simple clinical characteristics to clarify the locus specific characteristics. Currently this database contains information extracted from scientific papers and next generation sequencing results and other small scale experimental results of several research laboratories. Mutation data submission is greatly appreciated.

Abbreviations: Human Variation DB

Synonyms: Human Genome Variation Database

Resource Type: storage service resource, data repository, service resource, database, data or information resource

Defining Citation: [PMID:19629137](#), [PMID:21385384](#)

Keywords: gene, genome, disease, snp, next generation sequencing, gwas, genetic variation, mutation, copy number variation, insertion, deletion, structural variation

Funding: Japan Science and Technology Agency

Availability: The community can contribute to this resource

Resource Name: Human Variation DB

Resource ID: SCR_009014

Alternate IDs: nlx_153886

Record Creation Time: 20220129T080250+0000

Record Last Update: 20260805T044217+0000

Ratings and Alerts

No rating or validation information has been found for Human Variation DB.

No alerts have been found for Human Variation DB.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.