

# Resource Summary Report

Generated by [nidm-terms](#) on Jul 29, 2026

## PriVar

RRID:SCR\_010784

Type: Tool

### Proper Citation

PriVar (RRID:SCR\_010784)

### Resource Information

**URL:** <http://paed.hku.hk/uploadarea/yangwl/html/software.html>

**Proper Citation:** PriVar (RRID:SCR\_010784)

**Description:** A toolkit for prioritizing SNVs and indels from next-generation sequencing data.

**Abbreviations:** PriVar

**Resource Type:** software resource

**Resource Name:** PriVar

**Resource ID:** SCR\_010784

**Alternate IDs:** OMICS\_00160

**Record Creation Time:** 20220129T080300+0000

**Record Last Update:** 20260725T190706+0000

### Ratings and Alerts

No rating or validation information has been found for PriVar.

No alerts have been found for PriVar.

### 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 [nidm-terms](#) .

Shi X, et al. (2025) Novel clinical and genetic insights into Gitelman syndrome from 95 Chinese patients. Human genomics, 19(1), 114.

He M, et al. (2021) The Added Value of Whole-Exome Sequencing for Anomalous Fetuses With Detailed Prenatal Ultrasound and Postnatal Phenotype. Frontiers in genetics, 12, 627204.