

Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 5, 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: 20260801T190416+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 [SPARC SAWG](#) .

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.