

Resource Summary Report

Generated by [NIF](#) on Sep 3, 2026

GWAS Quality Control Pipeline

RRID:SCR_019241

Type: Tool

Proper Citation

GWAS Quality Control Pipeline (RRID:SCR_019241)

Resource Information

URL: <https://github.com/ikmb/gwas-qc/>

Proper Citation: GWAS Quality Control Pipeline (RRID:SCR_019241)

Description: Software tool as scalable GWAS Quality Control pipeline for biobank scale datasets.

Synonyms: BIGwas QC

Resource Type: software resource, software toolkit

Keywords: GWAS, data quality control, biobank scale datasets, genome wide association study

Availability: Free, Available for download, Freely available

Resource Name: GWAS Quality Control Pipeline

Resource ID: SCR_019241

License: MIT

Record Creation Time: 20220129T080344+0000

Record Last Update: 20260904T000456+0000

Ratings and Alerts

No rating or validation information has been found for GWAS Quality Control Pipeline.

No alerts have been found for GWAS Quality Control Pipeline.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at [NIF](#) .

Çubuk H, et al. (2026) Mechanistic modeling of recessive disease through allelic integration of variant effects. Cell systems, 17(6), 101595.

Rausch P, et al. (2025) First insights into microbial changes within an Inflammatory Bowel Disease Family Cohort study. Gut microbes, 17(1), 2559119.

Moitinho-Silva L, et al. (2022) Host genetic factors related to innate immunity, environmental sensing and cellular functions are associated with human skin microbiota. Nature communications, 13(1), 6204.

Kässens JC, et al. (2021) BIGwas: Single-command quality control and association testing for multi-cohort and biobank-scale GWAS/PheWAS data. GigaScience, 10(6).