# **Resource Summary Report**

Generated by RRID on May 5, 2025

## **GBRS**

RRID:SCR\_020963

Type: Tool

### **Proper Citation**

GBRS (RRID:SCR\_020963)

#### Resource Information

URL: https://gbrs.readthedocs.io/en/latest/

**Proper Citation:** GBRS (RRID:SCR\_020963)

**Description:** Software suite of tools for reconstructing genomes using RNA-Seq data from

multiparent population and for quantifying allele specific expression.

Resource Type: software toolkit, software resource

Keywords: Reconstructing genomes, RNA-Seq data, multiparent population, quantifying

allele specific expression

**Funding:** 

Availability: Free, Available for download, Freely available

Resource Name: GBRS

Resource ID: SCR\_020963

License: GPLv3

**Record Creation Time:** 20220129T080353+0000

**Record Last Update:** 20250503T060846+0000

### Ratings and Alerts

No rating or validation information has been found for GBRS.

No alerts have been found for GBRS.

### **Data and Source Information**

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 3 mentions in open access literature.

**Listed below are recent publications.** The full list is available at RRID.

Gerdes Gyuricza I, et al. (2022) Genome-wide transcript and protein analysis highlights the role of protein homeostasis in the aging mouse heart. Genome research, 32(5), 838.

Lee JY, et al. (2021) Misexpression of genes lacking CpG islands drives degenerative changes during aging. Science advances, 7(51), eabj9111.

Takemon Y, et al. (2021) Proteomic and transcriptomic profiling reveal different aspects of aging in the kidney. eLife, 10.