Resource Summary Report

Generated by dkNET on Apr 22, 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: 20250420T015824+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 dkNET.

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.