

Resource Summary Report

Generated by [RRID](#) on Aug 14, 2026

GTestimate

RRID:SCR_026562

Type: Tool

Proper Citation

GTestimate (RRID:SCR_026562)

Resource Information

URL: <https://github.com/Martin-Fahrenberger/GTestimate>

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Description: Software R package for scRNA-seq normalization. Uses Simple Good-Turing estimator for the per cell relative gene expression estimation.

Resource Type: data processing software, software resource, software application

Defining Citation: [DOI:10.1101/2024.07.02.601501](https://doi.org/10.1101/2024.07.02.601501)

Keywords: scRNA-seq, normalization, Good-Turing estimation, scRNA-seq normalization, gene expression estimation,

Funding: European Commission ;
Austrian Science Fund

Availability: Free, Available for download, Freely available

Resource Name: GTestimate

Resource ID: SCR_026562

License: GNU GPL-3.0 license

Record Creation Time: 20250313T060330+0000

Record Last Update: 20260813T055349+0000

Ratings and Alerts

No rating or validation information has been found for GTestimate.

No alerts have been found for GTestimate.

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 [RRID](#) .

Schuster VP, et al. (2026) Self-antigen disrupts cDC1 mediated antitumor responses. bioRxiv : the preprint server for biology.

Fahrenberger M, et al. (2025) GTestimate: improving relative gene expression estimation in scRNA-seq using the Good-Turing estimator. GigaScience, 14.