

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

[qusage](#)

RRID:SCR_024255

Type: Tool

Proper Citation

qusage (RRID:SCR_024255)

Resource Information

URL: <https://bioconductor.org/packages/qusage/>

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Description: Software R package is implementation Quantitative Set Analysis for Gene Expression method. Used to provide faster, more accurate, and easier to understand test for gene expression studies.

Synonyms: Quantitative Set Analysis for Gene Expression

Resource Type: software resource, software toolkit

Defining Citation: [PMID:23921631](#)

Keywords: test for gene expression studies,

Availability: Free, Available for download, Freely available,

Resource Name: qusage

Resource ID: SCR_024255

Alternate URLs: <https://sources.debian.org/src/r-bioc-qusage/>

Record Creation Time: 20230830T050217+0000

Record Last Update: 20260905T133106+0000

Ratings and Alerts

No rating or validation information has been found for qusage.

No alerts have been found for qusage.

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 [Kravitz](#) .

Zhang T, et al. (2025) Intratumoral Fusobacterium nucleatum Recruits Tumor-Associated Neutrophils to Promote Gastric Cancer Progression and Immune Evasion. Cancer research, 85(10), 1819.

Santos de Macedo BG, et al. (2024) Autophagy and inflammasome activation are associated with poor response to FLT3 inhibitors in patients with FLT3-ITD acute myeloid leukemia. Scientific reports, 14(1), 23882.

Menegatti S, et al. (2021) Immune response profiling of patients with spondyloarthritis reveals signalling networks mediating TNF-blocker function in vivo. Annals of the rheumatic diseases, 80(4), 475.