

# Resource Summary Report

Generated by [Kravitz](#) on Sep 17, 2026

## RnaSeqGeneEdgeRQL

RRID:SCR\_016699

Type: Tool

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### Proper Citation

RnaSeqGeneEdgeRQL (RRID:SCR\_016699)

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### Resource Information

**URL:**

<https://www.bioconductor.org/packages/release/workflows/html/RnaSeqGeneEdgeRQL.html>

**Proper Citation:** RnaSeqGeneEdgeRQL (RRID:SCR\_016699)

**Description:** Software to study analysis of an RNA-Seq experiment using the Rsubread and edgeR packages. The workflow starts from read alignment and continues on to data exploration, to differential expression and, finally, to pathway analysis. The analysis includes plots, GO and KEGG analyses, and the analysis of a expression signature as generated by a prior experiment.

**Resource Type:** data analysis software, data processing software, software application, software resource, software toolkit

**Keywords:** study, analysis, RNA, sequencing, read, alignment, data

**Availability:** Free, Available for download, Freely available

**Resource Name:** RnaSeqGeneEdgeRQL

**Resource ID:** SCR\_016699

**Record Creation Time:** 20220129T080331+0000

**Record Last Update:** 20260912T195843+0000

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### Ratings and Alerts

No rating or validation information has been found for RnaSeqGeneEdgeRQL.

No alerts have been found for RnaSeqGeneEdgeRQL.

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### Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.