

Resource Summary Report

Generated by [NIF](#) on Jul 30, 2026

RnaSeqGeneEdgeRQL

RRID:SCR_016699

Type: Tool

Proper Citation

RnaSeqGeneEdgeRQL (RRID:SCR_016699)

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 processing software, software toolkit, data analysis software, software resource, software application

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: 20260731T043005+0000

Ratings and Alerts

No rating or validation information has been found for RnaSeqGeneEdgeRQL.

No alerts have been found for RnaSeqGeneEdgeRQL.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.