

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

riboSeqR

RRID:SCR_016947

Type: Tool

Proper Citation

riboSeqR (RRID:SCR_016947)

Resource Information

URL: <https://bioconductor.org/packages/release/bioc/html/riboSeqR.html>

Proper Citation: riboSeqR (RRID:SCR_016947)

Description: Software tool for analysis of sequencing data from ribosome profiling experiments. Used for plotting functions, frameshift detection and parsing of sequencing data from ribosome profiling experiments.

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

Keywords: analysis, sequencing, data, ribosome, profiling, experiment, plotting, function, frameshift, detect, parsing

Availability: Free, Available for download, Freely available

Resource Name: riboSeqR

Resource ID: SCR_016947

License: GPL v3

Record Creation Time: 20220129T080332+0000

Record Last Update: 20260728T043528+0000

Ratings and Alerts

No rating or validation information has been found for riboSeqR.

No alerts have been found for riboSeqR.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Lim Y, et al. (2021) Multiplexed functional genomic analysis of 5' untranslated region mutations across the spectrum of prostate cancer. Nature communications, 12(1), 4217.

Cai EY, et al. (2020) Selective Translation of Cell Fate Regulators Mediates Tolerance to Broad Oncogenic Stress. Cell stem cell, 27(2), 270.

Gonatopoulos-Pournatzis T, et al. (2020) Autism-Misregulated eIF4G Microexons Control Synaptic Translation and Higher Order Cognitive Functions. Molecular cell, 77(6), 1176.

Sapkota D, et al. (2019) Cell-Type-Specific Profiling of Alternative Translation Identifies Regulated Protein Isoform Variation in the Mouse Brain. Cell reports, 26(3), 594.