

Resource Summary Report

Generated by [NIF](#) on Aug 27, 2026

RNA-eXpress

RRID:SCR_005167

Type: Tool

Proper Citation

RNA-eXpress (RRID:SCR_005167)

Resource Information

URL: <http://rnaexpress.org/>

Proper Citation: RNA-eXpress (RRID:SCR_005167)

Description: Software designed as a user friendly solution to extract and annotate biologically important transcripts from next generation RNA sequencing data.

Abbreviations: RNA-eXpress

Resource Type: software resource

Defining Citation: [PMID:23396121](#)

Keywords: bio.tools

Availability: Acknowledgement requested, Non-Profit

Resource Name: RNA-eXpress

Resource ID: SCR_005167

Alternate IDs: biotools:rna-express, OMICS_01285

Alternate URLs: <https://bio.tools/rna-express>

Record Creation Time: 20220129T080228+0000

Record Last Update: 20260815T182312+0000

Ratings and Alerts

No rating or validation information has been found for RNA-eXpress.

No alerts have been found for RNA-eXpress.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Afzal M, et al. (2020) Legume genomics and transcriptomics: From classic breeding to modern technologies. Saudi journal of biological sciences, 27(1), 543.

Hur EH, et al. (2017) Induction of immunoglobulin transcription factor 2 and resistance to MEK inhibitor in melanoma cells. Oncotarget, 8(25), 41387.

Pavli??ev M, et al. (2017) Single-cell transcriptomics of the human placenta: inferring the cell communication network of the maternal-fetal interface. Genome research, 27(3), 349.

Huang WH, et al. (2013) RNA editing and drug discovery for cancer therapy. TheScientificWorldJournal, 2013, 804505.

Forster SC, et al. (2013) RNA-eXpress annotates novel transcript features in RNA-seq data. Bioinformatics (Oxford, England), 29(6), 810.