

Resource Summary Report

Generated by [kravitz2](#) on Sep 18, 2026

RNAseqViewer

RRID:SCR_011900

Type: Tool

Proper Citation

RNAseqViewer (RRID:SCR_011900)

Resource Information

URL: <http://bioinfo.au.tsinghua.edu.cn/software/RNAseqViewer/>

Proper Citation: RNAseqViewer (RRID:SCR_011900)

Description: Software to visualize the various data from the RNA-Seq analyzing process, for single or multiple samples.

Abbreviations: RNAseqViewer

Resource Type: software resource

Defining Citation: [PMID:24215023](#)

Availability: Free to academic users

Resource Name: RNAseqViewer

Resource ID: SCR_011900

Alternate IDs: OMICS_01363

Record Creation Time: 20220129T080307+0000

Record Last Update: 20260912T195741+0000

Ratings and Alerts

No rating or validation information has been found for RNAseqViewer.

No alerts have been found for RNAseqViewer.

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 [kravitz2](#) .

Hersh CP, et al. (2019) High-Throughput Sequencing in Respiratory, Critical Care, and Sleep Medicine Research. An Official American Thoracic Society Workshop Report. Annals of the American Thoracic Society, 16(1), 1.

Kurt-K?z?ldo?an A, et al. (2017) Global transcriptome analysis of Halolamina sp. to decipher the salt tolerance in extremely halophilic archaea. Gene, 601, 56.

Pavlopoulos GA, et al. (2015) Visualizing genome and systems biology: technologies, tools, implementation techniques and trends, past, present and future. GigaScience, 4, 38.

Mazzanti CM, et al. (2015) Human saliva as route of inter-human infection for mouse mammary tumor virus. Oncotarget, 6(21), 18355.