

# Resource Summary Report

Generated by SPARC SAWG on Aug 10, 2026

## RNASeqBias

RRID:SCR\_001739

Type: Tool

### Proper Citation

RNASeqBias (RRID:SCR\_001739)

### Resource Information

**URL:**

[http://bioinformatics.dreamhosters.com/?page\\_id=113#Genomic\\_Protein\\_Sequence\\_Analysis](http://bioinformatics.dreamhosters.com/?page_id=113#Genomic_Protein_Sequence_Analysis)

**Proper Citation:** RNASeqBias (RRID:SCR\_001739)

**Description:** THIS RESOURCE IS NO LONGER IN SERVICE. Documented on April 10th,2023. An R software package for detecting and correcting biases in RNA-Sequencing data.

**Abbreviations:** RNASeqBias

**Resource Type:** software resource

**Keywords:** rna-seq

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE.

**Resource Name:** RNASeqBias

**Resource ID:** SCR\_001739

**Alternate IDs:** OMICS\_01957

**Record Creation Time:** 20220129T080209+0000

**Record Last Update:** 20260808T185743+0000

### Ratings and Alerts

No rating or validation information has been found for RNASeqBias.

No alerts have been found for RNASeqBias.

### 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.