

# Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 17, 2026

## [rpkmforgenes.py](#)

RRID:SCR\_014938

Type: Tool

### Proper Citation

rpkmforgenes.py (RRID:SCR\_014938)

### Resource Information

**URL:** <http://sandberg.cmb.ki.se/media/data/rnaseq/rpkmforgenes.py>

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**Description:** Python script which calculates gene expression for RNA-Sequencing data. It analyzes files in formats such as BED, BAM, and SAM to output data about RNA.

**Resource Type:** data analysis software, data processing software, sequence analysis software, software application, software resource, source code

**Keywords:** rna, sequencing, rna-seq, python, script, data analysis, genetic, gene analysis

**Availability:** Free, Available for download

**Resource Name:** rpkmforgenes.py

**Resource ID:** SCR\_014938

**Alternate URLs:** <http://sandberg.cmb.ki.se/media/data/rnaseq/rpkmforgenes.py>

**Record Creation Time:** 20220129T080323+0000

**Record Last Update:** 20260912T195823+0000

### Ratings and Alerts

No rating or validation information has been found for rpkmforgenes.py.

No alerts have been found for rpkmforgenes.py.

### Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Di Martino E, et al. (2024) Inflammatory, metabolic, and sex-dependent gene-regulatory dynamics of microglia and macrophages in neonatal hippocampus after hypoxia-ischemia. *iScience*, 27(4), 109346.

Chaparro V, et al. (2020) Translational profiling of macrophages infected with *Leishmania donovani* identifies mTOR- and eIF4A-sensitive immune-related transcripts. *PLoS pathogens*, 16(6), e1008291.