

Resource Summary Report

Generated by [dkNET](#) on May 22, 2025

[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: source code, data analysis software, software resource, software application, data processing software, sequence analysis software

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

Funding:

Availability: Free, Available for download

Resource Name: rpkmforgenes.py

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

Record Creation Time: 20220129T080323+0000

Record Last Update: 20250522T060927+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 [dkNET](#).

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.