

Resource Summary Report

Generated by [nidm-terms](#) on Sep 23, 2026

ShrinkSeq

RRID:SCR_012022

Type: Tool

Proper Citation

ShrinkSeq (RRID:SCR_012022)

Resource Information

URL: <http://www.few.vu.nl/~mavdwiel/ShrinkBayes.html>

Proper Citation: ShrinkSeq (RRID:SCR_012022)

Description: Software for detecting differential features across the entire spectrum, including the lower counts.

Abbreviations: ShrinkSeq

Resource Type: software resource

Defining Citation: [PMID:22988280](#)

Keywords: rna-seq, r

Resource Name: ShrinkSeq

Resource ID: SCR_012022

Alternate IDs: OMICS_01961

Record Creation Time: 20220129T080308+0000

Record Last Update: 20260919T195222+0000

Ratings and Alerts

No rating or validation information has been found for ShrinkSeq.

No alerts have been found for ShrinkSeq.

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 [nidm-terms](#) .

Neerincx M, et al. (2015) MiR expression profiles of paired primary colorectal cancer and metastases by next-generation sequencing. *Oncogenesis*, 4(10), e170.

van de Wiel MA, et al. (2014) ShrinkBayes: a versatile R-package for analysis of count-based sequencing data in complex study designs. *BMC bioinformatics*, 15, 116.