Resource Summary Report

Generated by NIF on Apr 27, 2025

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

Funding:

Resource Name: ShrinkSeq

Resource ID: SCR_012022

Alternate IDs: OMICS_01961

Record Creation Time: 20220129T080308+0000

Record Last Update: 20250420T014604+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: <u>SciCrunch Registry</u>

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>NIF</u>.

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 countbased sequencing data in complex study designs. BMC bioinformatics, 15, 116.