Resource Summary Report

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: 20250214T183208+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 RRID.

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.