

Resource Summary Report

Generated by [RRID](#) on Aug 14, 2026

[scShapes](#)

RRID:SCR_022838

Type: Tool

Proper Citation

scShapes (RRID:SCR_022838)

Resource Information

URL: <https://github.com/Malindrie/scShapes>

Proper Citation: scShapes (RRID:SCR_022838)

Description: Software tool as statistical framework for identifying distribution shapes in single-cell RNA-sequencing data.

Resource Type: data analysis software, data processing software, software resource, software application

Defining Citation: [DOI:10.1101/2022.02.13.480299](https://doi.org/10.1101/2022.02.13.480299)

Keywords: Differential distributions, single-cell RNA-sequencing, zero-inflation, scRNA-seq data, modeling gene expression read counts

Availability: Free, Available for download, Freely available

Resource Name: scShapes

Resource ID: SCR_022838

Record Creation Time: 20221008T050158+0000

Record Last Update: 20260814T043039+0000

Ratings and Alerts

No rating or validation information has been found for scShapes.

No alerts have been found for scShapes.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Dharmaratne M, et al. (2022) scShapes: a statistical framework for identifying distribution shapes in single-cell RNA-sequencing data. GigaScience, 12.