

# Resource Summary Report

Generated by [RRID](#) on Sep 10, 2026

## [chevreul](#)

RRID:SCR\_026966

Type: Tool

### Proper Citation

chevreul (RRID:SCR\_026966)

### Resource Information

**URL:** <https://github.com/cobriniklab/chevreul>

**Proper Citation:** chevreul (RRID:SCR\_026966)

**Description:** Software R package and shiny app for processing, plotting, and interactive exploratory data analysis of full-length scRNA-seq as SingleCellExperiments.

**Resource Type:** software resource, software toolkit, source code

**Keywords:** SingleCellExperiments, processing, plotting, interactive exploratory, data analysis, full-length scRNA-seq,

**Availability:** Free, Available for download, Freely available

**Resource Name:** chevreul

**Resource ID:** SCR\_026966

**License:** MIT license

**Record Creation Time:** 20250521T061058+0000

**Record Last Update:** 20260905T133536+0000

### Ratings and Alerts

No rating or validation information has been found for chevreul.

No alerts have been found for chevreul.

### 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](#) .

Stachelek K, et al. (2025) Chevreul: An R Bioconductor Package for Exploratory Analysis of Full-Length Single Cell Sequencing. bioRxiv : the preprint server for biology.

Stachelek K, et al. (2025) Chevreul: an R bioconductor package for exploratory analysis of full-length single cell sequencing. GigaByte (Hong Kong, China), 2025, gigabyte158.