

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 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: source code, software resource, software toolkit

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: 20260728T043746+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 [nidm-terms](#) .

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.