

# Resource Summary Report

Generated by NIF on May 22, 2025

## SeuratWrappers

RRID:SCR\_022555

Type: Tool

### Proper Citation

SeuratWrappers (RRID:SCR\_022555)

### Resource Information

**URL:** <https://github.com/satijalab/seurat-wrappers>

**Proper Citation:** SeuratWrappers (RRID:SCR\_022555)

**Description:** Software R package that helps format Seurat objects into objects that can be used for trajectory analysis in other packages such as Monocle

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

**Keywords:** helps format Seurat objects, trajectory analysis, Monocle

**Funding:**

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

**Resource Name:** SeuratWrappers

**Resource ID:** SCR\_022555

**License:** GNU GPL

**Record Creation Time:** 20220713T050151+0000

**Record Last Update:** 20250522T061356+0000

### Ratings and Alerts

No rating or validation information has been found for SeuratWrappers.

No alerts have been found for SeuratWrappers.

## Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We found 21 mentions in open access literature.

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

Chai D, et al. (2025) Lipid nanoparticles deliver DNA-encoded biologics and induce potent protective immunity. *Molecular cancer*, 24(1), 12.

Borges KS, et al. (2024) Non-canonical Wnt signaling triggered by WNT2B drives adrenal aldosterone production. *bioRxiv : the preprint server for biology*.

Ismaeel A, et al. (2024) Division-Independent Differentiation of Muscle Stem Cells During a Growth Stimulus. *Stem cells (Dayton, Ohio)*, 42(3), 266.

Aihara G, et al. (2024) SEraster: a rasterization preprocessing framework for scalable spatial omics data analysis. *Bioinformatics (Oxford, England)*, 40(7).

Sun S, et al. (2024) Efficient generation of human NOTCH ligand-expressing haemogenic endothelial cells as infrastructure for in vitro haematopoiesis and lymphopoiesis. *Nature communications*, 15(1), 7698.

Yoo K, et al. (2024) Muscle-resident mesenchymal progenitors sense and repair peripheral nerve injury via the GDNF-BDNF axis. *eLife*, 13.

Geller E, et al. (2024) Massively parallel disruption of enhancers active in human neural stem cells. *Cell reports*, 43(2), 113693.

Iyer DP, et al. (2024) mTOR activity paces human blastocyst stage developmental progression. *Cell*, 187(23), 6566.

Esfahani SN, et al. (2024) Derivation of human primordial germ cell-like cells in an embryonic-like culture. *Nature communications*, 15(1), 167.

Ascenção C, et al. (2024) A TOPBP1 allele causing male infertility uncouples XY silencing dynamics from sex body formation. *eLife*, 12.

Cheung G, et al. (2024) Multipotent progenitors instruct ontogeny of the superior colliculus. *Neuron*, 112(2), 230.

Lee H, et al. (2024) Nuclear respiratory factor-1 (NRF1) induction as a powerful strategy to deter mitochondrial dysfunction and senescence in mesenchymal stem cells. *Aging cell*, e14446.

Vasudevan P, et al. (2023) CCR2 macrophage response determines the functional outcome following cardiomyocyte transplantation. *Genome medicine*, 15(1), 61.

Chen X, et al. (2023) Genetic background of idiopathic neurodevelopmental delay patients with significant brain deviation volume. *Chinese medical journal*, 136(7), 807.

Naas S, et al. (2023) Hypoxia controls expression of kidney-pathogenic MUC1 variants. *Life science alliance*, 6(9).

Ko KD, et al. (2023) Integrating single-cell transcriptomes, chromatin accessibility, and multiomics analysis of mesoderm-induced embryonic stem cells. *STAR protocols*, 4(2), 102307.

Vandenbon A, et al. (2023) A universal tool for predicting differentially active features in single-cell and spatial genomics data. *Scientific reports*, 13(1), 11830.

Delannoy E, et al. (2023) Cell specialization and coordination in *Arabidopsis* leaves upon pathogenic attack revealed by scRNA-seq. *Plant communications*, 4(5), 100676.

Fleck JS, et al. (2023) Inferring and perturbing cell fate regulomes in human brain organoids. *Nature*, 621(7978), 365.

Eid SA, et al. (2023) Single-cell RNA-seq uncovers novel metabolic functions of Schwann cells beyond myelination. *Journal of neurochemistry*, 166(2), 367.