

Resource Summary Report

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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 resource, software toolkit

Keywords: helps format Seurat objects, trajectory analysis, Monocle

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: 20260821T194220+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 40 mentions in open access literature.

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

Rahmawati M, et al. (2026) Tandem single-cell epigenetic and transcriptomic analysis reveals spatiotemporally distinct genomic features among testis cell lineages. *iScience*, 29(7), 116452.

Aggrey-Fynn JE, et al. (2026) Epigenetic context defines the transcriptional activity of canonical and noncanonical NF- κ B signaling in pancreatic cancer. *Cell death discovery*, 12(1).

Collot R, et al. (2026) IGSF11-VISTA is a critical and targetable immune checkpoint axis in diffuse midline glioma. *Cancer cell*, 44(3), 641.

Chen CY, et al. (2026) Transcriptome-guided development of a fibrosis-reversal compound reduces skin scarring and allows regeneration via mitochondrial uncoupling. *Cell reports. Medicine*, 7(5), 102821.

Muppirala AN, et al. (2026) Tachykinin signaling defines distinct populations of glia in the enteric nervous system. *Neuron*.

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

Canizo JR, et al. (2025) The guinea pig serves as an alternative model to study human preimplantation development. *Nature cell biology*, 27(4), 696.

Almaghrabi R, et al. (2025) A heterozygous CEBPA mutation disrupting the bZIP domain in a RUNX1 and SRSF2 mutational background causes MDS disease progression. *Nature communications*, 16(1), 5489.

Keramati F, et al. (2025) Systemic inflammation impairs myelopoiesis and interferon type I responses in humans. *Nature immunology*, 26(5), 737.

Fagiani F, et al. (2025) Spatially-restricted inflammation-induced senescent-like glia in multiple sclerosis and patient-derived organoids. *Nature communications*, 16(1), 8477.

Zhang G, et al. (2025) Dissecting the Spatial and Single-Cell Transcriptomic Architecture of Cancer Stem Cell Niche Driving Tumor Progression in Gastric Cancer. *Advanced science (Weinheim, Baden-Wurttemberg, Germany)*, 12(18), e2413019.

Ennis CS, et al. (2025) Plasma exosomes from individuals with type 2 diabetes drive breast cancer aggression in patient-derived organoids. *Communications biology*, 8(1), 1276.

Prummel KD, et al. (2025) Inflammatory stromal and T cells mediate human bone marrow niche remodeling in clonal hematopoiesis and myelodysplasia. *Nature communications*, 16(1), 10042.

Lall D, et al. (2025) An organ-chip model of sporadic ALS using iPSC-derived spinal cord motor neurons and an integrated blood-brain-like barrier. *Cell stem cell*, 32(7), 1139.

Jayasinghe RG, et al. (2025) Single-cell transcriptomic profiling reveals diversity in human iNKT cells across hematologic tissues. *Cell reports*, 44(5), 115587.

Sato MP, et al. (2025) Transcriptomic comparison of avian auditory and vestibular sensory epithelia. *iScience*, 28(11), 113780.

Okoye GD, et al. (2025) Single-cell map of innate-like lymphocyte response to *Francisella tularensis* infection reveals interleukin-17-dependent protection by MAIT cells. *iScience*, 28(3), 111810.

Hata Y, et al. (2025) snRNA-seq analysis of the moss *Physcomitrium patens* identifies a conserved cytokinin-ESR module promoting pluripotent stem cell identity. *Developmental cell*, 60(13), 1884.

Gao Y, et al. (2025) Multimodal analyses reveal genes driving electrophysiological maturation of neurons in the primate prefrontal cortex. *Neuron*, 113(15), 2490.

Köhne M, et al. (2025) *Satb1* directs the differentiation of TH17 cells through suppression of IL-2 expression. *Cell reports*, 44(7), 115866.