

# Package: sceasy (via r-universe)

May 21, 2026

**Type** Package

**Title** A package to help convert different single-cell data formats to each other

**Version** 0.0.7

**Description** Convert Seurat, SingleCellExperiment, Loom object to AnnData object. Convert SingleCellExperiment object to and from Loom object.

**Depends** R (>= 3.5.1), reticulate

**Suggests** Seurat (>= 3.0.1), SingleCellExperiment (>= 1.4.0), LoomExperiment (>= 1.1.5), monocle3 (>= 0.2.0)

**SystemRequirements** anndata, loompy

**License** GPL-3

**LazyDate** true

**Config/pak/sysreqs** libpng-dev python3

**Repository** <https://blaserlab.r-universe.dev>

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**RemoteUrl** <https://github.com/cellgeni/sceasy>

**RemoteRef** HEAD

**RemoteSha** 3d0b5155752a61a5c42bd14fe60bd351400cbc51