

# Package: OSCA (via r-universe)

June 3, 2026

**Title** Orchestrating Single Cell Analysis with Bioconductor

**Version** 1.23.0

**Date** 2022-09-05

**Description** Online book for orchestrating single cell analysis with Bioconductor. Contains plenty of worked examples, theoretical background and some philosophical discussions on the nature of single-cell data analysis.

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**Encoding** UTF-8

**URL** <https://github.com/Bioconductor/OrchestratingSingleCellAnalysis>

**biocViews** ImmunoOncologyWorkflow, Workflow, SingleCellWorkflow

**Workflow** True

**VignetteBuilder** knitr

**Depends** BiocStyle, knitr, rebook, rmarkdown

**Suggests** OSCA.intro, OSCA.basic, OSCA.advanced, OSCA.workflows, OSCA.multisample

**Config/pak/sysreqs** cmake make libuv1-dev libxml2-dev

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

**Date/Publication** 2026-04-28 12:53:30 UTC

**RemoteUrl** <https://github.com/bioc/OSCA>

**RemoteRef** HEAD

**RemoteSha** 73bbe7a3f5e47eeb4b7835dee2bac642e1895976