

Package: OSCA.multisample (via r-universe)

October 10, 2026

Title OSCA for Multiple Samples

Version 1.21.1

Date 2026-08-28

Description Deploys the multi-sample analysis chapters for the
``Orchestrating Single Cell Analysis with Bioconductor" book.
This describes the handling of multiple samples in a
single-cell RNA-seq analysis, starting with integration of
multiple datasets into a common space for consistent analyses,
differential expression comparisons between conditions based on
pseudo-bulk samples, and differential abundance analyses for
cell subpopulations. It is intended for readers who are already
familiar with basic single-cell analyses, possibly after
reading some of the prior books in this collection.

Depends scrapper

Suggests rmarkdown, BiocStyle, knitr

VignetteBuilder knitr

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URL <https://github.com/OSCA-source/OSCA.multisample>

biocViews ImmunoOncologyWorkflow, Workflow, SingleCellWorkflow

Config/pak/sysreqs zlib1g-dev

Repository <https://bioc.r-universe.dev>

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