Package: OSCA.multisample (via r-universe)

September 27, 2026

Title OSCA for Multiple Samples

Version 1.21.1

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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://tempbioc.r-universe.dev

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