## Package 'SpatialDatasets'

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Version 1.7.3
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Title Collection of spatial omics datasets

**Description** This is a collection of publically available spatial omics datasets.

Where possible we have curated these datasets as either SpatialExperiments,

MoleculeExperiments or CytoImageLists and included annotations of the sample characteristics.

URL https://github.com/SydneyBioX/SpatialDatasets

BugReports https://github.com/SydneyBioX/SpatialDatasets/issues

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Encoding UTF-8

biocViews ExperimentData, ExperimentHub, ReproducibleResearch,

SingleCellData, SpatialData, Homo\_sapiens\_Data,

Mus\_musculus\_Data

Depends ExperimentHub, SpatialExperiment,

Imports utils

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Suggests BiocStyle, knitr, rmarkdown, cytomapper

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#### **Description**

This is a collection of publically available spatial omics datasets. Where possible we have curated these datasets as either SpatialExperiments, MoleculeExperiments or CytoImageLists and included annotations of the sample characteristics.

#### **Details**

The SpatialDatasets package contains a collection of spatially-resolved omics datasets for use in examples, demonstrations, and tutorials.

The datasets are from several different platforms including IMC, MIBI-TOF, CODEX, Xenium, CosMx and MERFISH. They have been sourced from various publicly available sources.

Additional examples and documentation are provided in the package vignette.

Datasets

The SpatialDatasets package contains the following datasets:

- spe\_Keren\_2018 (MIBI-TOF): A study on triple negative breast cancer containing 40 samples measured using MIBI-TOF published by [Keren et al. (2018)](https://doi.org/10.1016/j.cell.2018.08.039).
- Ferguson\_Images (IMC): A study on head and neck cutaneous squamous cell carcinoma containing 44 samples measured using IMC published by [Ferguson et al. (2022)](https://doi.org/10.1158/1078-0432.CCR-22-1332).
- spe\_Ferguson\_2022 (IMC): A study on head and neck cutaneous squamous cell carcinoma containing 44 samples measured using IMC published by [Ferguson et al. (2022)](https://doi.org/10.1158/1078-0432.CCR-22-1332).
- spe\_Schurch\_2020 (CODEX): A study on advanced colorectal cancer containing 140 samples measured using CODEX published by [Schurch et al. (2020)](https://doi.org/10.1016/j.cell.2020.07.005).
- spe\_Ali\_2020 (IMC): A study on breast cancer containing 483 samples measured using IMC published by [Ali et al. (2020)](https://doi.org/10.1038/s43018-020-0026-6).
- spe\_Amancherla\_2025 (Xenium): A study on heart transplant rejection containing 62 (49 adult and 13 pediatric) samples measured with Xenium published by [Amancherla et al. (2025)](https://doi.org/10.1101/2025.02.28.640852).
- spe\_Vannan\_2025 (Xenium): A study on pulmonary fibrosis containing 35 lung samples measured with Xenium published by [Vannan et al. (2025)](https://doi.org/10.1038/s41588-025-02080-x).

The datasets will be a SpatialExperiment or MoleculeExperiment.

# load using dataset name spe <- spe\_Keren\_2018() spe

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