

# Package ‘SpatialDatasets’

April 17, 2025

**Version** 1.6.0

**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>

**License** GPL-3

**Encoding** UTF-8

**biocViews** ExperimentData, ExperimentHub, ReproducibleResearch,  
SingleCellData, SpatialData, Homo\_sapiens\_Data,  
Mus\_musculus\_Data

**Depends** ExperimentHub, SpatialExperiment,

**Imports** utils

**VignetteBuilder** knitr

**Suggests** BiocStyle, knitr, rmarkdown

**RoxygenNote** 7.3.2

**git\_url** <https://git.bioconductor.org/packages/SpatialDatasets>

**git\_branch** RELEASE\_3\_21

**git\_last\_commit** a1f057d

**git\_last\_commit\_date** 2025-04-15

**Repository** Bioconductor 3.21

**Date/Publication** 2025-04-17

**Author** Farhan Ameen [aut],  
Nick Robertson [aut],  
Alex Qin [aut],  
Elijah Willie [aut],  
Ellis Patrick [aut, cre]

**Maintainer** Ellis Patrick <ellis.patrick@sydney.edu.au>

## Contents

SpatialDatasets . . . . .	2
<b>Index</b>	<b>3</b>

---

SpatialDatasets	<i>A collection of publically available spatial omics datasets.</i>
-----------------	---

---

### 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>).

The datasets will be a SpatialExperiment or MoleculeExperiment.

```
# load using dataset name spe <- spe_Keren_2018() spe
```

# Index

Ferguson\_Images (SpatialDatasets), [2](#)

SpatialDatasets, [2](#)

spe\_Ali\_2020 (SpatialDatasets), [2](#)

spe\_Ferguson\_2022 (SpatialDatasets), [2](#)

spe\_Keren\_2018 (SpatialDatasets), [2](#)

spe\_Schurch\_2020 (SpatialDatasets), [2](#)