

Package ‘TENxVisiumData’

April 30, 2026

Version 1.21.0

Title Visium spatial gene expression data by 10X Genomics

Description Collection of Visium spatial gene expression datasets by 10X Genomics, formatted into objects of class `SpatialExperiment`.

Data cover various organisms and tissues, and include: single- and multi-section experiments, as well as single sections subjected to both whole transcriptome and targeted panel analysis. Datasets may be used for testing of and as examples in packages, for tutorials and workflow demonstrations, or similar purposes.

URL <https://github.com/helenalc/TENxVisiumData>

BugReports <https://github.com/helenalc/TENxVisiumData/issues>

biocViews ExperimentHub, ExpressionData, ExperimentData, Homo_sapiens_Data, Mus_musculus_Data, ReproducibleResearch

Depends ExperimentHub, SpatialExperiment

Imports utils

Suggests BiocStyle, knitr, rmarkdown

VignetteBuilder knitr

License MIT + file LICENSE

Encoding UTF-8

RoxygenNote 7.1.1

git_url <https://git.bioconductor.org/packages/TENxVisiumData>

git_branch devel

git_last_commit 3fa35d9

git_last_commit_date 2026-04-28

Repository Bioconductor 3.24

Date/Publication 2026-04-30

Author Helena L. Crowell [aut, cre] (ORCID: <https://orcid.org/0000-0002-4801-1767>)

Maintainer Helena L. Crowell <helena@crowell.eu>

Contents

TENxVisiumData	2
Index	4

TENxVisiumData

TENxVisiumData

Description

Collection of Visium spatial gene expression datasets by 10X Genomics, formatted into objects of class `SpatialExperiment`. Data cover various organisms and tissues, and include: single- and multi-section experiments, as well as single sections subjected to both whole transcriptome and targeted panel analysis. Datasets may be used for testing of and as examples in packages, for tutorials and workflow demonstrations, or similar purposes.

Details

The following Visium Spatial Transcriptomics datasets by 10X Genomics are currently available (see the package vignette for links to details):

- `HumanBreastCancerIDC`
- `HumanBreastCancerILC`
- `HumanCerebellum`
- `HumanColorectalCancer`
- `HumanGlioblastoma`
- `HumanHeart`
- `HumanLymphNode`
- `HumanOvarianCancer`
- `HumanSpinalCord`
- `MouseBrainCoronal`
- `MouseBrainSagittalPosterior`
- `MouseBrainSagittalAnterior`
- `MouseKidneyCoronal`

Value

a `SpatialExperiment` with rows corresponding to samples (spots) and columns to features (transcripts). If, in addition to whole transcriptome analysis, the specified dataset contains measurements from targeted panel analysis/es, these will be stored as `altExps`.

Author(s)

Helena L. Crowell

Examples

```
# initialize hub instance
eh <- ExperimentHub()

# query for TENxVisium datasets
(q <- query(eh, "TENxVisium"))

# retrieve dataset of interest
```

```
pat <- "HumanCerebellum$"  
idx <- grep(pat, q$title)  
id <- q$ah_id[idx]  
(spe <- eh[[id]])  
  
# access targeted panel  
altExpNames(spe)  
altExp(spe, "TargetedNeuroscience")
```

Index

HumanBreastCancerIDC (TENxVisiumData), 2
HumanBreastCancerILC (TENxVisiumData), 2
HumanCerebellum (TENxVisiumData), 2
HumanColorectalCancer (TENxVisiumData),
2
HumanGlioblastoma (TENxVisiumData), 2
HumanHeart (TENxVisiumData), 2
HumanLymphNode (TENxVisiumData), 2
HumanOvarianCancer (TENxVisiumData), 2
HumanSpinalCord (TENxVisiumData), 2

MouseBrainCoronal (TENxVisiumData), 2
MouseBrainSagittalAnterior
(TENxVisiumData), 2
MouseBrainSagittalPosterior
(TENxVisiumData), 2
MouseKidneyCoronal (TENxVisiumData), 2

SpatialExperiment, 2

TENxVisiumData, 2