

Package ‘scTHI.data’

October 28, 2025

Type Package
Title The package contains examples of single cell data used in vignettes and examples of the scTHI package; data contain both tumor cells and immune cells from public dataset of glioma
Version 1.21.0
Description Data for the vignette and tutorial of the package scTHI.
License GPL-2
Depends R (>= 4.0)
Encoding UTF-8
LazyData false
RoxygenNote 6.1.1.9000
biocViews ExperimentData, SingleCellData
git_url <https://git.bioconductor.org/packages/scTHI.data>
git_branch devel
git_last_commit abce09f
git_last_commit_date 2025-04-15
Repository Bioconductor 3.22
Date/Publication 2025-10-28
Author Francesca Pia Caruso [aut],
Michele Ceccarelli [aut, cre]
Maintainer Michele Ceccarelli <m.ceccarelli@gmail.com>

Contents

H3K27	2
H3K27.meta	2
MGH45	3
MGH45.annotation	3
scExample	3
Index	4

H3K27

Example expression matrix for scTHI.

Description

A 21673 x 5527 matrix patient PatientBCH836 form Filbin, M. G., Tirosh, I., Hovestadt, V., Shaw, M. L., Escalante, L. E., Mathewson, N. D., ... & Haberler, C. (2018). Developmental and oncogenic programs in H3K27M gliomas dissected by single-cell RNA-seq. Science, 360(6386), 331-335. H3K27

Usage

H3K27

Format

An object of class `matrix` with 21673 rows and 527 columns.

H3K27.meta

Annotation for the H3K27 e expression matrix for scTHI.

Description

A dataframe 527x9 for the matrix of patient PatientBCH836 form Filbin, M. G., Tirosh, I., Hovestadt, V., Shaw, M. L., Escalante, L. E., Mathewson, N. D., ... & Haberler, C. (2018). Developmental and oncogenic programs in H3K27M gliomas dissected by single-cell RNA-seq. Science, 360(6386), 331-335. H3K27.annotation

Usage

H3K27.meta

Format

An object of class `data.frame` with 527 rows and 9 columns.

`MGH45`*Example expression matrix for scTHI.*

Description

A 17584 x 608 matrix patient PatientBCH836 form Venteicher AS, Tirosh I, Hebert C, Yizhak K et al. Decoupling genetics, lineages, and microenvironment in IDH-mutant gliomas by single-cell RNA-seq. Science 2017 Mar 31;355(6332) MGH45

Usage`MGH45`**Format**

An object of class `matrix` with 17584 rows and 608 columns.

`MGH45.annotation`*Annotation for Example expression matrix for scTHI.*

Description

A 608 x 2 dataframe with the annotation for the MGH4 matrix Venteicher AS, Tirosh I, Hebert C, Yizhak K et al. Decoupling genetics, lineages, and microenvironment in IDH-mutant gliomas by single-cell RNA-seq. Science 2017 Mar 31;355(6332) MGH45.annotation

Usage`MGH45.annotation`**Format**

An object of class `data.frame` with 608 rows and 2 columns.

`scExample`*Example expression matrix for scTHI.*

Description

A 2000 x 100 matrix from the wiki manual to showcase the use of scTHI `scExample`

Usage`scExample`**Format**

An object of class `matrix` with 2000 rows and 100 columns.

Index

* **datasets**

H3K27, [2](#)

H3K27.meta, [2](#)

MGH45, [3](#)

MGH45.annotation, [3](#)

scExample, [3](#)

H3K27, [2](#)

H3K27.meta, [2](#)

MGH45, [3](#)

MGH45.annotation, [3](#)

scExample, [3](#)