

Package ‘SomatiCAData’

January 16, 2025

Version 1.44.0

Title An example cancer whole genome sequencing data for the SomatiCA package

Description An example cancer whole genome sequencing data for the SomatiCA package

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Depends R (>= 2.14)

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LazyData yes

biocViews ExperimentData, Genome, CancerData, SequencingData

InstallableEverywhere yes

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

git_branch RELEASE_3_20

git_last_commit c84ee6c

git_last_commit_date 2024-10-29

Repository Bioconductor 3.20

Date/Publication 2025-01-16

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SomatiCAData-package *An example cancer whole genome sequencing data for the SomatiCA package*

Description

A dataset of a paired tumor-normal sequencing sample from Complete Genomics.

Author(s)

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Examples

```
data(glio)
```

glio

Sequencing data for a tumor sample with control.

Description

A dataset of a paired tumor-normal sequencing sample from Complete Genomics.

Usage

```
data(glio)
```

Format

A data frame with 3458745 rows on the following 7 variables.

seqnames a character vector

start a character vector

zygosity a character vector

tCount an integer vector

LAF a numeric vector

tCountN an integer vector

germLAF a numeric vector

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