

Package ‘SomatiCData’

March 28, 2024

Version 1.40.0

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)

License Artistic-2.0

LazyData yes

biocViews ExperimentData, Genome, CancerData, SequencingData

InstallableEverywhere yes

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

git_branch RELEASE_3_18

git_last_commit cee61d4

git_last_commit_date 2023-10-24

Repository Bioconductor 3.18

Date/Publication 2024-03-28

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