

Package ‘SomatiCAData’

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

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

Author Mengjie Chen

Maintainer Mengjie Chen <mengjie.chen@yale.edu>

Depends R (>= 2.14)

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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)

Mengjie Chen <mengjie.chen@yale.edu>

Maintainer: Mengjie Chen <mengjie.chen@yale.edu>

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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