

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

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Depends R (>= 2.14)

License Artistic-2.0

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Contents

SomatiCAData-package	1
glio	2
Index	3

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

Index

- * **datasets**

- [glio](#), [2](#)

- * **package**

- [SomatiCData-package](#), [1](#)

[glio](#), [2](#)

[SomatiCData \(SomatiCData-package\)](#), [1](#)

[SomatiCData-package](#), [1](#)