

Package: SomatiCAData (via r-universe)

October 10, 2026

Version 1.50.0

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

Repository Bioconductor 3.23

Date/Publication 2026-04-28 12:25:20 UTC

RemoteUrl <https://github.com/bioc/SomatiCAData>

RemoteRef RELEASE_3_23

RemoteSha 15fb95966002668736513338878d25758203f026

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SomaticData-package	<i>An example cancer whole genome sequencing data for the SomaticData package</i>
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Description

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

Author(s)

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Examples

```
data(glio)
```

glio	<i>Sequencing data for a tumor sample with control.</i>
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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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