

Package ‘RTCGA.PANCAN12’

October 10, 2024

Type Package

Title PanCan 12 from Genome Cancer Browser

Version 1.33.0

Date 2015-11-10

Author Przemyslaw Biecek <przemyslaw.biecek@gmail.com>

Maintainer Marcin Kosinski <m.p.kosinski@gmail.com>

Description Package provides clinical, expression, cnv and mutation data from Genome Cancer Browser.

License GPL-2

LazyData FALSE

Roxygen list(wrap = TRUE)

BugReports <https://github.com/RTCGA/RTCGA/issues>

Depends R (>= 3.3.0), RTCGA

Suggests knitr, rmarkdown

biocViews AnnotationData

VignetteBuilder knitr

NeedsCompilation no

RoxygenNote 5.0.1

git_url <https://git.bioconductor.org/packages/RTCGA.PANCAN12>

git_branch devel

git_last_commit 263359a

git_last_commit_date 2024-04-30

Repository Bioconductor 3.20

Date/Publication 2024-10-10

Contents

pancan12	2
Index	3

pancan12

PANCAN12 datasets from Cancer Browser project

Description

Package provides clinical, expression, cnv and mutation data from Genome Cancer Browser for 12 cohorts types. Converting those datasets from data.frames to Bioconductor classes is explained here [convertTCGA](#).

Usage

clinical.cb

cnv.cb

mutation.cb

expression.cb1

expression.cb2

Value

Data frames PANCAN12 data.

Source

<https://genome-cancer.ucsc.edu/proj/site/hgHeatmap/>

Index

* datasets

pancan12, [2](#)

clinical.cb (pancan12), [2](#)

cnv.cb (pancan12), [2](#)

cnv.cb, (pancan12), [2](#)

convertTCGA, [2](#)

expression.cb, (pancan12), [2](#)

expression.cb1 (pancan12), [2](#)

expression.cb2 (pancan12), [2](#)

mutation.cb (pancan12), [2](#)

mutation.cb, (pancan12), [2](#)

pancan12, [2](#)