

Package ‘RTCGA.PANCAN12’

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Type Package

Title PanCan 12 from Genome Cancer Browser

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

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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>

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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/>

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