

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>

git_branch devel

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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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* **datasets**

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