

# Package ‘RTCGA.clinical’

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**Type** Package

**Title** Clinical datasets from The Cancer Genome Atlas Project

**Version** 20151101.36.0

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**Description** Package provides clinical datasets from The Cancer Genome Atlas Project for all cohorts types from <http://gdac.broadinstitute.org/>. Clinical data format is explained here <https://wiki.nci.nih.gov/display/TCGA/Clinical+Data+Overview>. Data from 2015-11-01 snapshot.

**License** GPL-2

**Suggests** rmarkdown, knitr

**VignetteBuilder** knitr

**LazyData** TRUE

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

**Depends** R (>= 3.2.0), RTCGA

**biocViews** Annotation Data

**NeedsCompilation** no

**git\_url** <https://git.bioconductor.org/packages/RTCGA.clinical>

**git\_branch** RELEASE\_3\_20

**git\_last\_commit** b02d84e

**git\_last\_commit\_date** 2024-10-29

**Repository** Bioconductor 3.20

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`clinical`*Clinical datasets from TCGA project*

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## Description

Package provides clinical datasets from The Cancer Genome Atlas Project for all cohorts types from <http://gdac.broadinstitute.org/>. Data were downloaded using [RTCGA-package](#) and contain snapshots for the date: 2015-11-01. The process is described here: <http://rtcga.github.io/RTCGA/>. Use cases, examples and information about datasets in **RTCGA.data** family can be found here: `browseVignettes("RTCGA")`. Clinical data format is explained here <https://wiki.nci.nih.gov/display/TCGA/Clinical+Data+Overview>.

## Usage

```
ACC.clinical
BLCA.clinical
BRCA.clinical
CESC.clinical
CHOL.clinical
COAD.clinical
COADREAD.clinical
DLBC.clinical
ESCA.clinical
FPPP.clinical
GBM.clinical
GBMLGG.clinical
HNSC.clinical
KICH.clinical
KIPAN.clinical
KIRC.clinical
KIRP.clinical
LAML.clinical
LGG.clinical
```

LIHC.clinical

LUAD.clinical

LUSC.clinical

MESO.clinical

OV.clinical

PAAD.clinical

PCPG.clinical

PRAD.clinical

READ.clinical

SARC.clinical

SKCM.clinical

STAD.clinical

STES.clinical

TGCT.clinical

THCA.clinical

THYM.clinical

UCEC.clinical

UCS.clinical

UVM.clinical

**Details**

`browseVignettes("RTCGA")`

**Value**

Data frames with clinical data.

**Source**

<http://gdac.broadinstitute.org/>

