

# Package ‘RTCGA.clinical’

April 10, 2025

**Type** Package

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

**Version** 20151101.37.0

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**Author** Marcin Kosinski <m.p.kosinski@gmail.com>

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

**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** devel

**git\_last\_commit** 706d355

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

**Repository** Bioconductor 3.21

**Date/Publication** 2025-04-10

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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/>

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