

Package: RTCGA.RPPA (via r-universe)

October 10, 2026

Type Package

Title RPPA datasets from The Cancer Genome Atlas Project

Version 1.40.0

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Description Package provides RPPA datasets from The Cancer Genome Atlas Project for all available cohorts types from <http://gdac.broadinstitute.org/>. Data format is explained here <https://wiki.nci.nih.gov/display/TCGA/Protein+Array+Data+Format+Specification?src=search>

License GPL-2

LazyData 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

Config/pak/sysreqs cmake make libicu-dev libjpeg-dev libpng-dev libuv1-dev libxml2-dev libssl-dev

Repository Bioconductor 3.23

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RemoteUrl <https://github.com/bioc/RTCGA.RPPA>

RemoteRef RELEASE_3_23

RemoteSha 78a528543859a694c55f292fd920cf02dc6dae53

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Description

Package provides RPPA datasets from The Cancer Genome Atlas Project for all available 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://rtcg.github.io/RTCGA/>. Use cases, examples and information about datasets in **RTCGA.data** family can be found here: `browseVignettes("RTCGA")`. RPPA data format is explained here <https://wiki.nci.nih.gov/display/TCGA/Protein+Array+Data+Format+Specification?src=search>. Converting **RTCGA.RPPA** to Bioconductor datasets' format is explained here [convertTCGA](#).

Usage

`ACC.RPPA``BLCA.RPPA``BRCA.RPPA``CESC.RPPA``CHOL.RPPA``COAD.RPPA``COADREAD.RPPA``DLBC.RPPA``ESCA.RPPA``GBM.RPPA``GBMLGG.RPPA``HNSC.RPPA``KICH.RPPA``KIPAN.RPPA``KIRC.RPPA``KIRP.RPPA`

LGG.RPPA

LIHC.RPPA

LUAD.RPPA

LUSC.RPPA

MESO.RPPA

OV.RPPA

PAAD.RPPA

PCPG.RPPA

PRAD.RPPA

READ.RPPA

SARC.RPPA

SKCM.RPPA

STAD.RPPA

STES.RPPA

TGCT.RPPA

THCA.RPPA

THYM.RPPA

UCEC.RPPA

UCS.RPPA

UVM.RPPA

Value

Data frames with RPPA data.

Source

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

Examples

```
## Not run:  
browseVignettes("RTCGA")  
browseVignettes("RTCGA.RPPA")  
  
## End(Not run)
```

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