

Package ‘RTCGA.rnaseq’

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Title Rna-seq datasets from The Cancer Genome Atlas Project

Version 1.0.0

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Description Package provides rna-seq datasets from The Cancer Genome Atlas Project for all cohorts types from <http://gdac.broadinstitute.org/>. Rna-seq data format is explained here <https://wiki.nci.nih.gov/display/TCGA/RNASeq+Version+2>.

License GPL-2

LazyData TRUE

Repository Bioconductor

BugReports <https://github.com/mi2-warsaw/RTCGA.data/issues>

Depends R (>= 3.2.0), RTCGA

Suggests knitr

biocViews Annotation Data

VignetteBuilder knitr

NeedsCompilation no

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rnaseq

rnaseq datasets from TCGA project

Description

rnaseq data format is explained here <https://wiki.nci.nih.gov/display/TCGA/RNASeq+Version+2>. Package provides rnaseq 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: 2015-08-21. The process is described here: <http://mi2-warsaw.github.io/RTCGA.data/>. Use cases, examples and information about datasets in **RTCGA.data** family can be found here: `browseVignettes("RTCGA")`

Usage

ACC.rnaseq

BLCA.rnaseq

BRCA.rnaseq

CESC.rnaseq

CHOL.rnaseq

COAD.rnaseq

DLBC.rnaseq

ESCA.rnaseq

GBM.rnaseq

HNSC.rnaseq

KICH.rnaseq

KIPAN.rnaseq

KIRC.rnaseq

KIRP.rnaseq

LAML.rnaseq

LGG.rnaseq

LIHC.rnaseq

LUAD.rnaseq

LUSC.rnaseq

MESO.rnaseq

OV.rnaseq

PAAD.rnaseq

PCPG.rnaseq

PRAD.rnaseq

READ.rnaseq

SKCM.rnaseq

STES.rnaseq

TGCT.rnaseq

THCA.rnaseq

THYM.rnaseq

UCEC.rnaseq

UCS.rnaseq

UVM.rnaseq

Details

`browseVignettes("RTCGA")`

Value

Data frames with rnaseq data.

Source

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

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