

Package ‘RTCGA.mutations’

April 13, 2017

Type Package

Title Mutations datasets from The Cancer Genome Atlas Project

Version 20151101.4.0

Date 2016-04-26

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Description Package provides mutations datasets from The Cancer Genome
Atlas Project for all cohorts types from
<http://gdac.broadinstitute.org/>. Mutations data format is
explained here
[https://wiki.nci.nih.gov/display/TCGA/Mutation+Annotation+Format+\(MAF\)+Specification](https://wiki.nci.nih.gov/display/TCGA/Mutation+Annotation+Format+(MAF)+Specification).
There is extra one column with patients' barcodes. Data from 2015-11-01 snapshot.

License GPL-2

LazyData TRUE

Repository Bioconductor

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

Depends R (>= 3.2.0), RTCGA

Suggests knitr

biocViews Annotation Data

VignetteBuilder knitr

NeedsCompilation no

R topics documented:

mutations 2

Index 4

<code>mutations</code>	<i>Mutations datasets from TCGA project</i>
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Description

Package provides mutations 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")` Mutations data format is explained here <https://wiki.nci.nih.gov/display/TCGA/RTCGA+Data+Format>. There is extra one column with patients' barcodes.

Usage

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

```
LUAD.mutations  
LUSC.mutations  
OV.mutations  
PAAD.mutations  
PCPG.mutations  
PRAD.mutations  
READ.mutations  
SKCM.mutations  
STAD.mutations  
STES.mutations  
SARC.mutations  
TGCT.mutations  
THCA.mutations  
UCEC.mutations  
UCS.mutations  
UVM.mutations
```

Details

```
browseVignettes("RTCGA")
```

Value

Data frames with mutations data.

Source

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

Index

*Topic **datasets**
 mutations, 2

ACC.mutations (mutations), 2

BLCA.mutations (mutations), 2

BRCA.mutations (mutations), 2

CESC.mutations (mutations), 2

CHOL.mutations (mutations), 2

COAD.mutations (mutations), 2

COADREAD.mutations (mutations), 2

DLBC.mutations (mutations), 2

ESCA.mutations (mutations), 2

GBM.mutations (mutations), 2

GBMLGG.mutations (mutations), 2

HNSC.mutations (mutations), 2

KICH.mutations (mutations), 2

KIPAN.mutations (mutations), 2

KIRC.mutations (mutations), 2

KIRP.mutations (mutations), 2

LAML.mutations (mutations), 2

LGG.mutations (mutations), 2

LIHC.mutations (mutations), 2

LUAD.mutations (mutations), 2

LUSC.mutations (mutations), 2

mutations, 2

OV.mutations (mutations), 2

PAAD.mutations (mutations), 2

PCPG.mutations (mutations), 2

PRAD.mutations (mutations), 2

READ.mutations (mutations), 2

RTCGA-package, 2

SARC.mutations (mutations), 2

SKCM.mutations (mutations), 2

STAD.mutations (mutations), 2

STES.mutations (mutations), 2

TGCT.mutations (mutations), 2

THCA.mutations (mutations), 2

UCEC.mutations (mutations), 2

UCS.mutations (mutations), 2

UVM.mutations (mutations), 2