TxDb.Hsapiens.UCSC.hg19.knownGene

September 21, 2022

TxDb.Hsapiens.UCSC.hg19.knownGene

Annotation package for TxDb object(s)

Description

This package loads one or more TxDb objects. Such TxDb objects are an R interface to prefabricated databases contained by this package.

The names of any objects exposed by this package indicate the origin and resources exposed. So for example TxDb.Hsapiens.UCSC.hg19.knownGene would be a TxDb object, of Homo sapiens data from UCSC build hg19 based on the knownGene Track.

Note

This data package was made from resources at UCSC on 2015-10-07 18:11:28 +0000 (Wed, 07 Oct 2015) and based on the hg19 genome based on the knownGene table

Author(s)

Marc Carlson, Bioconductor Package Maintainer <maintainer@bioconductor.org> [cre]

See Also

transcripts, transcriptsBy

Examples

load the library library(TxDb.Hsapiens.UCSC.hg19.knownGene) ## list the contents that are loaded into memory ls('package:TxDb.Hsapiens.UCSC.hg19.knownGene') ## show the db object that is loaded by calling it's name TxDb.Hsapiens.UCSC.hg19.knownGene

Index

* data
 TxDb.Hsapiens.UCSC.hg19.knownGene,
 1
* package
 TxDb.Hsapiens.UCSC.hg19.knownGene,
 1
transcripts, 1
transcriptsBy, 1
TxDb.Hsapiens.UCSC.hg19.knownGene, 1
TxDb.Hsapiens.UCSC.hg19.knownGene-package
 (TxDb.Hsapiens.UCSC.hg19.knownGene),
 1