TxDb.Hsapiens.UCSC.hg38.knownGene

June 4, 2025

TxDb.Hsapiens.UCSC.hg38.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 2025-03-02 02:45:03 +0000 (Sun, 02 Mar 2025) and based on the hg38 genome based on the knownGene table

Author(s)

Bioconductor Core Team, Bioconductor Package Maintainer <maintainer@bioconductor.org> [cre]

See Also

transcripts, transcriptsBy

Examples

load the library library(TxDb.Hsapiens.UCSC.hg38.knownGene) ## list the contents that are loaded into memory ls('package:TxDb.Hsapiens.UCSC.hg38.knownGene') ## show the db object that is loaded by calling it's name TxDb.Hsapiens.UCSC.hg38.knownGene

Index

* data
 TxDb.Hsapiens.UCSC.hg38.knownGene,
 1
* package
 TxDb.Hsapiens.UCSC.hg38.knownGene,
 1
transcripts, 1
transcriptsBy, 1
TxDb.Hsapiens.UCSC.hg38.knownGene, 1
TxDb.Hsapiens.UCSC.hg38.knownGene-package
 (TxDb.Hsapiens.UCSC.hg38.knownGene),
 1