TxDb.Hsapiens.UCSC.hg38.knownGene

February 28, 2017

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 2016-09-29 13:02:09 +0000 (Thu, 29 Sep 2016) 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

```
*Topic data
TxDb.Hsapiens.UCSC.hg38.knownGene,

1
*Topic package
TxDb.Hsapiens.UCSC.hg38.knownGene,

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

1
```