TxDb.Hsapiens.UCSC.hg19.refGene

November 17, 2026

`TxDb.Hsapiens.UCSC.hg19.refGene`
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 2025-11-01 23:04:26 -0700 (Sat, 01 Nov 2025) and based on the hg19 genome based on the refGene 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.hg19.refGene)
## list the contents that are loaded into memory
ls('package:TxDb.Hsapiens.UCSC.hg19.refGene')
## show the db object that is loaded by calling it's name
TxDb.Hsapiens.UCSC.hg19.refGene
```

Index

∗ **data**
TxDb.Hsapiens.UCSC.hg19.refGene, 1
∗ **package**
TxDb.Hsapiens.UCSC.hg19.refGene, 1

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