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 2023-09-20 17:26:31 +0000 (Wed, 20 Sep 2023) and based on the hg38 genome based on the refGene table.

Author(s)

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

See Also

transcripts, transcriptsBy

Examples

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

* data
  - TxDb.Hsapiens.UCSC.hg38.refGene

* package
  - TxDb.Hsapiens.UCSC.hg38.refGene

transcripts, /
transcriptsBy, /
TxDb.Hsapiens.UCSC.hg38.refGene
TxDb.Hsapiens.UCSC.hg38.refGene-package
  (TxDb.Hsapiens.UCSC.hg38.refGene),
  1