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 2022-04-07 16:51:36 +0000 (Thu, 07 Apr 2022) and based on the rn7 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.Rnorvegicus.UCSC.rn7.refGene)
## list the contents that are loaded into memory
ls('package:TxDb.Rnorvegicus.UCSC.rn7.refGene')
## show the db object that is loaded by calling it's name
TxDb.Rnorvegicus.UCSC.rn7.refGene
```
Index

* data
  TxDb.Rnorvegicus.UCSC.rn7.refGene,

* package
  TxDb.Rnorvegicus.UCSC.rn7.refGene,

transcripts, /
transcriptsBy, /
TxDb.Rnorvegicus.UCSC.rn7.refGene, l
TxDb.Rnorvegicus.UCSC.rn7.refGene-package
  (TxDb.Rnorvegicus.UCSC.rn7.refGene),
  l