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 04:15:25 +0000 (Thu, 29 Sep 2016) and based on the mm10 genome based on the ensGene table

Author(s)

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

See Also

transcripts, transcriptsBy

Examples

```r
## load the library
library(TxDb.Mmusculus.UCSC.mm10.ensGene)
## list the contents that are loaded into memory
ls('package:TxDb.Mmusculus.UCSC.mm10.ensGene')
## show the db object that is loaded by calling it's name
TxDb.Mmusculus.UCSC.mm10.ensGene
```
Index

*Topic data
  Txdb.Mmusculus.UCSC.mm10.ensGene, 1

*Topic package
  Txdb.Mmusculus.UCSC.mm10.ensGene-package
  (Txdb.Mmusculus.UCSC.mm10.ensGene), 1

transcripts./
transcriptsBy./
Txdb.Mmusculus.UCSC.mm10.ensGene, 1
Txdb.Mmusculus.UCSC.mm10.ensGene-package
  (Txdb.Mmusculus.UCSC.mm10.ensGene), 1