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 04:30:42 +0000 (Thu, 29 Sep 2016) and based on the panTro4 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.Ptroglodytes.UCSC.panTro4.refGene)
## list the contents that are loaded into memory
ls('package:TxDb.Ptroglodytes.UCSC.panTro4.refGene')
## show the db object that is loaded by calling it's name
TxDb.Ptroglodytes.UCSC.panTro4.refGene
```
Index

*Topic data
  TxDB.Ptroglodytes.UCSC.panTro4.refGene,

*Topic package
  TxDB.Ptroglodytes.UCSC.panTro4.refGene,

transcripts, /
transcriptsBy, /
TxDB.Ptroglodytes.UCSC.panTro4.refGene,

TxDB.Ptroglodytes.UCSC.panTro4.refGene-package
  (TxDB.Ptroglodytes.UCSC.panTro4.refGene),