TxDb.Mmulatta.UCSC.rheMac8.refGene

February 28, 2017

TxDb.Mmulatta.UCSC.rheMac8.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 2016-09-29 03:44:33 +0000 (Thu, 29 Sep 2016) and based on the rheMac8 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.Mmulatta.UCSC.rheMac8.refGene)
## list the contents that are loaded into memory
ls('package:TxDb.Mmulatta.UCSC.rheMac8.refGene')
## show the db object that is loaded by calling it's name
TxDb.Mmulatta.UCSC.rheMac8.refGene
```

Index

```
*Topic data
TxDb.Mmulatta.UCSC.rheMac8.refGene,
1
*Topic package
TxDb.Mmulatta.UCSC.rheMac8.refGene,
1
transcripts, I
transcriptsBy, I
TxDb.Mmulatta.UCSC.rheMac8.refGene, 1
TxDb.Mmulatta.UCSC.rheMac8.refGene-package
(TxDb.Mmulatta.UCSC.rheMac8.refGene),
1
```