**TxDb.Mmulatta.UCSC.rheMac8.refGene**

*July 17, 2024*

---

**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 2020-10-11 17:32:17 +0000 (Sun, 11 Oct 2020) 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**

```r
## 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

* data
  
  TxDb.Mmulatta.UCSC.rheMac8.refGene,

* package
  
  TxDb.Mmulatta.UCSC.rheMac8.refGene,

transcripts, /
transcriptsBy, /
TxDb.Mmulatta.UCSC.rheMac8.refGene, 
TxDb.Mmulatta.UCSC.rheMac8.refGene-package
  
  (TxDb.Mmulatta.UCSC.rheMac8.refGene),
