**TxDb.Mmulatta.UCSC.rheMac3.refGene**

December 9, 2016

---

**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 04:35:23 +0000 (Thu, 29 Sep 2016) and based on the rheMac3 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.rheMac3.refGene)
## list the contents that are loaded into memory
ls('package:TxDb.Mmulatta.UCSC.rheMac3.refGene')
## show the db object that is loaded by calling it's name
TxDb.Mmulatta.UCSC.rheMac3.refGene
```
Index

*Topic data
  TxDb.Mmulatta.UCSC.rheMac3.refGene,
  1

*Topic package
  TxDb.Mmulatta.UCSC.rheMac3.refGene,
  1

transcripts. /
transcriptsBy. /
TxDb.Mmulatta.UCSC.rheMac3.refGene, 1
TxDb.Mmulatta.UCSC.rheMac3.refGene-package
  (TxDb.Mmulatta.UCSC.rheMac3.refGene),
  1