# TxDb.Mmulatta.UCSC.rheMac3.refGene

January 29, 2025

TxDb.Mmulatta.UCSC.rheMac3.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 17:32:41 +0000 (Sun, 11 Oct 2020) 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**

```
## 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**

```
* data
    TxDb.Mmulatta.UCSC.rheMac3.refGene,
    1
* package
    TxDb.Mmulatta.UCSC.rheMac3.refGene,
    1

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