# TxDb.Mmulatta.UCSC.rheMac8.refGene

December 18, 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**

```
## 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**