## TxDb.Mmulatta.UCSC.rheMac3.refGene

September 27, 2024

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