TxDb.Mmulatta.UCSC.rheMac8.refGene

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 $\label{thm:continuous} {\it TxDb.Mmulatta.UCSC.rheMac8.refGene} \\ {\it Annotation\ package\ for\ TxDb\ object(s)}$

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 rheMac8 refGene on 2017-10-02 12:20:15 -0400 (Mon, 02 Oct 2017) and based on the UCSC rheMac8 refGene

Author(s)

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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
```

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