TxDb.Mmulatta.UCSC.rheMac10.refGene

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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 $2021-09-16\ 17:16:42+0000$ (Thu, $16\ Sep\ 2021$) and based on the rheMac10 genome based on the refGene table

Author(s)

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See Also

transcripts, transcriptsBy

Examples

```
## load the library
library(TxDb.Mmulatta.UCSC.rheMac10.refGene)
## list the contents that are loaded into memory
ls('package:TxDb.Mmulatta.UCSC.rheMac10.refGene')
## show the db object that is loaded by calling it's name
TxDb.Mmulatta.UCSC.rheMac10.refGene
```

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