TxDb.Mmusculus.UCSC.mm9.knownGene

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 ${\it TxDb.Mmusculus.UCSC.mm9.knownGene} \\ Annotation\ package\ for\ TxDb\ object(s)$

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 2015-03-19 13:58:21 -0700 (Thu, 19 Mar 2015) and based on the mm9 genome based on the knownGene table

Author(s)

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See Also

transcripts, transcriptsBy

Examples

```
## load the library
library(TxDb.Mmusculus.UCSC.mm9.knownGene)
## list the contents that are loaded into memory
ls('package:TxDb.Mmusculus.UCSC.mm9.knownGene')
## show the db object that is loaded by calling it's name
TxDb.Mmusculus.UCSC.mm9.knownGene
```

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