## TxDb.Mmusculus.UCSC.mm9.knownGene

November 20, 2024

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 2015-10-07 18:13:02 +0000 (Wed, 07 Oct 2015) and based on the mm9 genome based on the knownGene table

#### Author(s)

Marc Carlson, Bioconductor Package Maintainer <maintainer@bioconductor.org> [cre]

#### 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
```

# **Index**

```
* data
    TxDb.Mmusculus.UCSC.mm9.knownGene,
    1
* package
    TxDb.Mmusculus.UCSC.mm9.knownGene,
    1

transcripts, I
transcriptsBy, I
TxDb.Mmusculus.UCSC.mm9.knownGene, 1
TxDb.Mmusculus.UCSC.mm9.knownGene-package
    (TxDb.Mmusculus.UCSC.mm9.knownGene),
    1
```