

# TxDB.Mmusculus.UCSC.mm10.knownGene

April 10, 2015

---

TxDB.Mmusculus.UCSC.mm10.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 2014-09-26 11:21:18 -0700 (Fri, 26 Sep 2014) and based on the mm10 genome based on the knownGene table

## Author(s)

Marc Carlson

## See Also

[transcripts](#), [transcriptsBy](#)

## Examples

```
## load the library
library(TxDB.Mmusculus.UCSC.mm10.knownGene)
## list the contents that are loaded into memory
ls(package:TxDB.Mmusculus.UCSC.mm10.knownGene)
## show the db object that is loaded by calling its name
TxDB.Mmusculus.UCSC.mm10.knownGene
```

# Index

\*Topic **data**

`TxDb.Mmusculus.UCSC.mm10.knownGene,`

[1](#)

\*Topic **package**

`TxDb.Mmusculus.UCSC.mm10.knownGene,`

[1](#)

`transcripts, 1`

`transcriptsBy, 1`

`TxDb.Mmusculus.UCSC.mm10.knownGene, 1`

`TxDb.Mmusculus.UCSC.mm10.knownGene-package`  
`(TxDb.Mmusculus.UCSC.mm10.knownGene),`

[1](#)