

TxDB.Mmusculus.UCSC.mm10.knownGene

September 24, 2013

TxDB.Mmusculus.UCSC.mm10.knownGene

Annotation package for TranscriptDb object(s)

Description

This package loads one or more TranscriptDb objects. Such TranscriptDb 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 TranscriptDb 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 2013-03-08 09:47:19 -0800 (Fri, 08 Mar 2013) 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 it's 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](#)