TxDb.Mmusculus.UCSC.mm10.knownGene

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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 2014-03-17 16:23:39 -0700 (Mon, 17 Mar 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
```

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