TxDb.Dmelanogaster.UCSC.dm3.ensGene

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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 2012-09-10 13:00:23 -0700 (Mon, 10 Sep 2012) and based on the dm3 genome based on the ensGene table

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See Also

transcripts, transcriptsBy

Examples

```
## load the library
library(TxDb.Dmelanogaster.UCSC.dm3.ensGene)
## list the contents that are loaded into memory
ls('package:TxDb.Dmelanogaster.UCSC.dm3.ensGene')
## show the db object that is loaded by calling it's name
TxDb.Dmelanogaster.UCSC.dm3.ensGene
```

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