FDb.UCSC.tRNAs

November 11, 2026

FDb.UCSC.tRNAs *Annotation package for FeatureDb object(s)*

Description

This package loads one or more FeatureDb objects. Such FeatureDb 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 13:15:01 -0700 (Mon, 10 Sep 2012) and based on the hg19 genome based on the tRNAs table

Author(s)

Marc Carlson

See Also

transcripts, transcriptsBy

Examples

```
## load the library
library(FDb.UCSC.tRNAs)
## list the contents that are loaded into memory
ls('package:FDb.UCSC.tRNAs')
## show the db object that is loaded by calling it's name
FDb.Hsapiens.UCSC.hg19.tRNAs
```

Index

∗ **data**
FDb.UCSC.tRNAs, 1
∗ **package**
FDb.UCSC.tRNAs, 1

FDb.Hsapiens.UCSC.hg18.tRNAs
*(*FDb.UCSC.tRNAs*)*, 1
FDb.Hsapiens.UCSC.hg19.tRNAs
*(*FDb.UCSC.tRNAs*)*, 1
FDb.Mmusculus.UCSC.mm9.tRNAs
*(*FDb.UCSC.tRNAs*)*, 1
FDb.Rnorvegicus.UCSC.rn4.tRNAs
*(*FDb.UCSC.tRNAs*)*, 1
FDb.UCSC.tRNAs, 1
FDb.UCSC.tRNAs-package
*(*FDb.UCSC.tRNAs*)*, 1

transcripts, *1*
transcriptsBy, *1*