

TxDB.Rnorvegicus.UCSC.rn4.ensGene

April 2, 2013

TxDB.Rnorvegicus.UCSC.rn4.ensGene

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 13:03:11 -0700 (Mon, 10 Sep 2012) and based on the rn4 genome based on the ensGene table

Author(s)

Marc Carlson

See Also

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

Examples

```
## load the library
library(TxDB.Rnorvegicus.UCSC.rn4.ensGene)
## list the contents that are loaded into memory
ls('package:TxDB.Rnorvegicus.UCSC.rn4.ensGene')
## show the db object that is loaded by calling it's name
TxDb.Rnorvegicus.UCSC.rn4.ensGene
```

Index

*Topic **data**

TxDb.Rnorvegicus.UCSC.rn4.ensGene,
[1](#)

*Topic **package**

TxDb.Rnorvegicus.UCSC.rn4.ensGene,
[1](#)

transcripts, [1](#)

transcriptsBy, [1](#)

TxDb.Rnorvegicus.UCSC.rn4.ensGene, [1](#)

TxDb.Rnorvegicus.UCSC.rn4.ensGene-
package
(TxDb.Rnorvegicus.UCSC.rn4.ensGene),
[1](#)