

TxDB.Cfamiliaris.UCSC.canFam3.refGene

November 23, 2022

TxDB.Cfamiliaris.UCSC.canFam3.refGene

Annotation package for TxDb object(s)

Description

This package loads one or more TxDb objects. Such TxDb 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 TxDb 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 2020-04-28 14:19:25 +0000 (Tue, 28 Apr 2020) and based on the canFam3 genome based on the refGene table

Author(s)

Bioconductor Core Team, Bioconductor Package Maintainer <maintainer@bioconductor.org> [cre]

See Also

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

Examples

```
## load the library
library(TxDB.Cfamiliaris.UCSC.canFam3.refGene)
## list the contents that are loaded into memory
ls('package:TxDB.Cfamiliaris.UCSC.canFam3.refGene')
## show the db object that is loaded by calling it's name
TxDb.Cfamiliaris.UCSC.canFam3.refGene
```

Index

*** data**

`Txdb.Cfamiliaris.UCSC.canFam3.refGene,`

[1](#)

*** package**

`Txdb.Cfamiliaris.UCSC.canFam3.refGene,`

[1](#)

`transcripts,` [1](#)

`transcriptsBy,` [1](#)

`Txdb.Cfamiliaris.UCSC.canFam3.refGene,`

[1](#)

`Txdb.Cfamiliaris.UCSC.canFam3.refGene-package`

`(Txdb.Cfamiliaris.UCSC.canFam3.refGene),`

[1](#)