**TxDb.Cfamiliaris.UCSC.canFam5.refGene**

**January 10, 2024**

---

**TxDb.Cfamiliaris.UCSC.canFam5.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 2021-09-16 17:16:23 +0000 (Thu, 16 Sep 2021) and based on the canFam5 genome based on the refGene table

**Author(s)**

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

**See Also**

transcripts, transcriptsBy

**Examples**

```r
## load the library
library(TxDb.Cfamiliaris.UCSC.canFam5.refGene)
## list the contents that are loaded into memory
ls('package:TxDb.Cfamiliaris.UCSC.canFam5.refGene')
## show the db object that is loaded by calling it's name
TxDb.Cfamiliaris.UCSC.canFam5.refGene
```
Index

* data
  TxDb.Cfamiliaris.UCSC.canFam5.refGene,
  |
* package
  TxDb.Cfamiliaris.UCSC.canFam5.refGene,
  |

transcripts, /
transcriptsBy, /
TxDb.Cfamiliaris.UCSC.canFam5.refGene,
  |
TxDb.Cfamiliaris.UCSC.canFam5.refGene-package
  (TxDb.Cfamiliaris.UCSC.canFam5.refGene),
  |