# TxDb.Cfamiliaris.UCSC.canFam6.refGene

October 9, 2024

---

`TxDb.Cfamiliaris.UCSC.canFam6.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 2023-03-16 20:08:08 +0000 (Thu, 16 Mar 2023) and based on the canFam6 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.canFam6.refGene)
## list the contents that are loaded into memory
ls('package:TxDb.Cfamiliaris.UCSC.canFam6.refGene')
## show the db object that is loaded by calling it's name
TxDb.Cfamiliaris.UCSC.canFam6.refGene
```

# Index

∗ **data**
TxDb.Cfamiliaris.UCSC.canFam6.refGene, 1
∗ **package**
TxDb.Cfamiliaris.UCSC.canFam6.refGene, 1

transcripts, *1*
transcriptsBy, *1*
TxDb.Cfamiliaris.UCSC.canFam6.refGene, 1
TxDb.Cfamiliaris.UCSC.canFam6.refGene-package (TxDb.Cfamiliaris.UCSC.canFam6.refGene), 1