**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**

```r
## 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
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