

TxDB.Drerio.UCSC.danRer10.refGene

January 19, 2022

TxDB.Drerio.UCSC.danRer10.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 2019-04-25 15:14:04 +0000 (Thu, 25 Apr 2019) and based on the danRer10 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.Drerio.UCSC.danRer10.refGene)
## list the contents that are loaded into memory
ls('package:TxDB.Drerio.UCSC.danRer10.refGene')
## show the db object that is loaded by calling it's name
TxDb.Drerio.UCSC.danRer10.refGene
```

Index

*** data**

[TxDb.Drerio.UCSC.danRer10.refGene,](#)
[1](#)

*** package**

[TxDb.Drerio.UCSC.danRer10.refGene,](#)
[1](#)

[transcripts,](#) [1](#)

[transcriptsBy,](#) [1](#)

[TxDb.Drerio.UCSC.danRer10.refGene,](#) [1](#)

[TxDb.Drerio.UCSC.danRer10.refGene-package](#)
[\(TxDb.Drerio.UCSC.danRer10.refGene\),](#)
[1](#)