

# TxDB.Drerio.UCSC.danRer11.refGene

July 24, 2024

---

TxDB.Drerio.UCSC.danRer11.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 15:14:29 +0000 (Thu, 25 Apr 2019) and based on the danRer11 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.danRer11.refGene)
## list the contents that are loaded into memory
ls('package:TxDB.Drerio.UCSC.danRer11.refGene')
## show the db object that is loaded by calling it's name
TxDb.Drerio.UCSC.danRer11.refGene
```

# Index

**\* data**

`Txdb.Drerio.UCSC.danRer11.refGene,`  
[1](#)

**\* package**

`Txdb.Drerio.UCSC.danRer11.refGene,`  
[1](#)

`transcripts,` [1](#)

`transcriptsBy,` [1](#)

`Txdb.Drerio.UCSC.danRer11.refGene,` [1](#)

`Txdb.Drerio.UCSC.danRer11.refGene-package`  
`(Txdb.Drerio.UCSC.danRer11.refGene),`  
[1](#)