

# TxDb.Drerio.UCSC.danRer10.refGene

May 22, 2024

---

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](#)