

# Package: TxDb.Drerio.UCSC.danRer11.ensGene (via r-universe)

August 11, 2024

**Title** Annotation package for TxDb object(s)

**Description** Exposes an annotation databases generated from UCSC by exposing these as TxDb objects

**Version** 1.0.0

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**Depends** GenomicFeatures (>= 1.50.4)

**Imports** AnnotationDbi

**License** MIT + file LICENSE

**organism** Danio rerio

**species** Danio rerio

**provider** UCSC

**provider\_version** danRer11 genome based on the ensGene table

**release\_date** 2023-03-20 09:52:36 -0400 (Mon, 20 Mar 2023)

**resource\_url** <http://genome.ucsc.edu/>

**biocViews** AnnotationData, Genetics, TxDb, Danio\_rerio

**URL** <https://github.com/blaserlab/TxDb.Drerio.UCSC.danRer11.ensGene>

**BugReports** <https://github.com/blaserlab/TxDb.Drerio.UCSC.danRer11.ensGene/issues>

**Repository** <https://blaserlab.r-universe.dev>

**RemoteUrl** <https://github.com/blaserlab/TxDb.Drerio.UCSC.danRer11.ensGene>

**RemoteRef** HEAD

**RemoteSha** fa2a71b165971ceeba7a58a69a85b996a2adc71b

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TxDb.Drerio.UCSC.danRer11.ensGene

*Annotation package for TxDb object(s)*

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## 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 2023-03-20 09:52:36 -0400 (Mon, 20 Mar 2023) and based on the danRer11 genome based on the ensGene table

## Author(s)

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## See Also

[transcripts](#), [transcriptsBy](#)

## Examples

```
## load the library
library(TxDb.Drerio.UCSC.danRer11.ensGene)
## list the contents that are loaded into memory
ls('package:TxDb.Drerio.UCSC.danRer11.ensGene')
## show the db object that is loaded by calling it's name
TxDb.Drerio.UCSC.danRer11.ensGene
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