# TxDb.Dmelanogaster.UCSC.dm6.ensGene

July 3, 2024

 $\label{thm:constraint} {\sf TxDb.Dmelanogaster.UCSC.dm6.ensGene} \\ {\it Annotation\ package\ for\ TxDb\ object(s)}$ 

## **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 2020-10-11 17:28:24 +0000 (Sun, 11 Oct 2020) and based on the dm6 genome based on the ensGene table

### Author(s)

Bioconductor Core Team, Bioconductor Package Maintainer <maintainer@bioconductor.org> [cre]

#### See Also

transcripts, transcriptsBy

#### **Examples**

```
## load the library
library(TxDb.Dmelanogaster.UCSC.dm6.ensGene)
## list the contents that are loaded into memory
ls('package:TxDb.Dmelanogaster.UCSC.dm6.ensGene')
## show the db object that is loaded by calling it's name
TxDb.Dmelanogaster.UCSC.dm6.ensGene
```

# **Index**