

# TxDB.Dmelanogaster.UCSC.dm3.ensGene

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TxDB.Dmelanogaster.UCSC.dm3.ensGene

*Annotation package for TranscriptDb object(s)*

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## Description

This package loads one or more TranscriptDb objects. Such TranscriptDb 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 TranscriptDb 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 2013-09-19 00:24:08 -0700 (Thu, 19 Sep 2013) and based on the dm3 genome based on the ensGene table

## Author(s)

Marc Carlson

## See Also

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

## Examples

```
## load the library
library(TxDB.Dmelanogaster.UCSC.dm3.ensGene)
## list the contents that are loaded into memory
ls('package:TxDB.Dmelanogaster.UCSC.dm3.ensGene')
## show the db object that is loaded by calling it's name
TxDb.Dmelanogaster.UCSC.dm3.ensGene
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