

# TxDb.Celegans.UCSC.ce11.ensGene

May 1, 2024

---

TxDb.Celegans.UCSC.ce11.ensGene

*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 2022-04-15 17:21:59 +0000 (Fri, 15 Apr 2022) and based on the ce11 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.Celegans.UCSC.ce11.ensGene)
## list the contents that are loaded into memory
ls('package:TxDb.Celegans.UCSC.ce11.ensGene')
## show the db object that is loaded by calling it's name
TxDb.Celegans.UCSC.ce11.ensGene
```

# Index

\* **data**

`Txdb.Celegans.UCSC.ce11.ensGene`, [1](#)

\* **package**

`Txdb.Celegans.UCSC.ce11.ensGene`, [1](#)

`transcripts`, [1](#)

`transcriptsBy`, [1](#)

`Txdb.Celegans.UCSC.ce11.ensGene`, [1](#)

`Txdb.Celegans.UCSC.ce11.ensGene-package`  
(`Txdb.Celegans.UCSC.ce11.ensGene`),  
[1](#)