

# TxDb.Scerevisiae.UCSC.sacCer2.ensGene

February 8, 2012

---

Scerevisiae\_UCSC\_sacCer2\_ensGene\_TxDb

*Annotation package for the Scerevisiae\_UCSC\_sacCer2\_ensGene\_TxDb object*

---

## Description

Contains the Scerevisiae\_UCSC\_sacCer2\_ensGene\_TxDb object annotation database as generated from UCSC

This package loads the above mentioned object. This TranscriptDb object is an R interface to the prefabricated database contained by this package.

## Note

This Transcript data package was made from resources at UCSC on 2011-05-03 15:19:18 -0700 (Tue, 03 May 2011) and based on the sacCer2 genome base on the ensGene table

## Author(s)

Marc Carlson

## See Also

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

## Examples

```
## load the library
library(TxDb.Scerevisiae.UCSC.sacCer2.ensGene)
## list the contents that are loaded into memory
ls('package:TxDb.Scerevisiae.UCSC.sacCer2.ensGene')
## show the txdb object that is loaded by calling it's name
TxDb.Scerevisiae.UCSC.sacCer2.ensGene
```

# Index

## \*Topic **data**

Scerevisiae\_UCSC\_sacCer2\_ensGene\_TxDb,  
[1](#)

## \*Topic **package**

Scerevisiae\_UCSC\_sacCer2\_ensGene\_TxDb,  
[1](#)

Scerevisiae\_UCSC\_sacCer2\_ensGene\_TxDb,  
[1](#)

transcripts, [1](#)

transcriptsBy, [1](#)

TxDb.Scerevisiae.UCSC.sacCer2.ensGene  
(*Scerevisiae\_UCSC\_sacCer2\_ensGene\_TxDb*),  
[1](#)

TxDb.Scerevisiae.UCSC.sacCer2.ensGene-package  
(*Scerevisiae\_UCSC\_sacCer2\_ensGene\_TxDb*),  
[1](#)