

TxDb.Scerevisiae.UCSC.sacCer2.sgdGene

February 8, 2012

```
TxDb.Scerevisiae.UCSC.sacCer2.sgdGene
```

Annotation package for the Scerevisiae_UCSC_sacCer2_sgdGene_TxDb object

Description

Contains the Scerevisiae_UCSC_sacCer2_sgdGene_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 2011-09-29 18:28:29 -0700 (Thu, 29 Sep 2011) and based on the sacCer2 genome base on the sgdGene table

Author(s)

Marc Carlson

See Also

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

Examples

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

Index

*Topic **data**

`TxDb.Scerevisiae.UCSC.sacCer2.sgdGene,`

[1](#)

*Topic **package**

`TxDb.Scerevisiae.UCSC.sacCer2.sgdGene,`

[1](#)

`transcripts,` [1](#)

`transcriptsBy,` [1](#)

`TxDb.Scerevisiae.UCSC.sacCer2.sgdGene,`

[1](#)

`TxDb.Scerevisiae.UCSC.sacCer2.sgdGene-package`

`(TxDb.Scerevisiae.UCSC.sacCer2.sgdGene),`

[1](#)