TxDb.Scerevisiae.UCSC.sacCer2.sgdGene

November 17, 2026

`TxDb.Scerevisiae.UCSC.sacCer2.sgdGene`
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 2015-10-07 18:19:57 +0000 (Wed, 07 Oct 2015) and based on the sacCer2 genome based on the sgdGene table

Author(s)

Marc Carlson, Bioconductor Package Maintainer <maintainer@bioconductor.org> [cre]

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 db object that is loaded by calling it's name
TxDb.Scerevisiae.UCSC.sacCer2.sgdGene
```

Index

∗ **data**
TxDb.Scerevisiae.UCSC.sacCer2.sgdGene, 1
∗ **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