

TxDB.Scerevisiae.UCSC.sacCer3.sgdGene

December 12, 2014

TxDB.Scerevisiae.UCSC.sacCer3.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 2014-09-26 11:28:15 -0700 (Fri, 26 Sep 2014) and based on the sacCer3 genome based on the sgdGene table

Author(s)

Marc Carlson

See Also

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

Examples

```
## load the library
library(TxDB.Scerevisiae.UCSC.sacCer3.sgdGene)
## list the contents that are loaded into memory
ls(package:TxDB.Scerevisiae.UCSC.sacCer3.sgdGene)
## show the db object that is loaded by calling its name
TxDB.Scerevisiae.UCSC.sacCer3.sgdGene
```

Index

*Topic **data**

`TxDb.Scerevisiae.UCSC.sacCer3.sgdGene,`

[1](#)

*Topic **package**

`TxDb.Scerevisiae.UCSC.sacCer3.sgdGene,`

[1](#)

`transcripts, 1`

`transcriptsBy, 1`

`TxDb.Scerevisiae.UCSC.sacCer3.sgdGene,`

[1](#)

`TxDb.Scerevisiae.UCSC.sacCer3.sgdGene-package`

`(TxDb.Scerevisiae.UCSC.sacCer3.sgdGene),`

[1](#)