# TxDb.Scerevisiae.UCSC.sacCer3.sgdGene

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`TxDb.Scerevisiae.UCSC.sacCer3.sgdGene`
*Annotation package for TxDb object(s)*

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## 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 18:20:42 +0000 (Wed, 07 Oct 2015) and based on the sacCer3 genome based on the sgdGene table

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## 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 it's name
TxDb.Scerevisiae.UCSC.sacCer3.sgdGene
```

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