**TxDb.Sscrofa.UCSC.susScr3.refGene**

April 26, 2017

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**TxDb.Sscrofa.UCSC.susScr3.refGene**  
*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 2016-09-29 04:28:10 +0000 (Thu, 29 Sep 2016) and based on the susScr3 genome based on the refGene table

**Author(s)**

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**See Also**

transcripts, transcriptsBy

**Examples**

```r
## load the library
library(TxDb.Sscrofa.UCSC.susScr3.refGene)
## list the contents that are loaded into memory
ls(package:TxDb.Sscrofa.UCSC.susScr3.refGene)
## show the db object that is loaded by calling it's name
TxDb.Sscrofa.UCSC.susScr3.refGene
```
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