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

February 28, 2024

---

**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 2020-10-11 17:31:23 +0000 (Sun, 11 Oct 2020) and based on the susScr3 genome based on the refGene table

**Author(s)**

Bioconductor Core Team, Bioconductor Package Maintainer <maintainer@bioconductor.org> [cre]

**See Also**

transcripts, transcriptsBy

**Examples**

```r
## load the library
callTxDb('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
```
Index

* data
  TxDb.Sscrofa.UCSC.susScr3.refGene,
  1

* package
  TxDb.Sscrofa.UCSC.susScr3.refGene,
  1

transcripts, /
transcriptsBy, /
TxDb.Sscrofa.UCSC.susScr3.refGene, 1
TxDb.Sscrofa.UCSC.susScr3.refGene-package
  (TxDb.Sscrofa.UCSC.susScr3.refGene),
  1