**TxDb.Hsapiens.UCSC.hg38.knownGene**

February 28, 2017

---

**TxDb.Hsapiens.UCSC.hg38.knownGene**

*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 2016-09-29 13:02:09 +0000 (Thu, 29 Sep 2016) and based on the hg38 genome based on the knownGene table.

**Author(s)**

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

**See Also**

transcripts, transcriptsBy

**Examples**

```r
## load the library
library(TxDB.Hsapiens.UCSC.hg38.knownGene)
## list the contents that are loaded into memory
ls('\package:TxDb.Hsapiens.UCSC.hg38.knownGene\')
## show the db object that is loaded by calling it's name
TxDb.Hsapiens.UCSC.hg38.knownGene
```
Index

*Topic data
  TxDB.Hsapiens.UCSC.hg38.knownGene, 1

*Topic package
  TxDB.Hsapiens.UCSC.hg38.knownGene, 1

transcripts, /
  transcriptsBy, /
  TxDB.Hsapiens.UCSC.hg38.knownGene, 1
  TxDB.Hsapiens.UCSC.hg38.knownGene-package
    (TxDB.Hsapiens.UCSC.hg38.knownGene), 1