

# TxDB.Hsapiens.UCSC.hg19.knownGene

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TxDB.Hsapiens.UCSC.hg19.knownGene

*Annotation package for TranscriptDb object(s)*

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## Description

This package loads one or more TranscriptDb objects. Such TranscriptDb 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 TranscriptDb 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 2012-03-12 21:45:23 -0700 (Mon, 12 Mar 2012) and based on the hg19 genome based on the knownGene table

## Author(s)

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## See Also

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

## Examples

```
## load the library
library(TxDB.Hsapiens.UCSC.hg19.knownGene)
## list the contents that are loaded into memory
ls('package:TxDB.Hsapiens.UCSC.hg19.knownGene')
## show the db object that is loaded by calling it's name
TxDb.Hsapiens.UCSC.hg19.knownGene
```

# Index

\*Topic **data**

`TxDb.Hsapiens.UCSC.hg19.knownGene,`

[1](#)

\*Topic **package**

`TxDb.Hsapiens.UCSC.hg19.knownGene,`

[1](#)

`transcripts,` [1](#)

`transcriptsBy,` [1](#)

`TxDb.Hsapiens.UCSC.hg19.knownGene,` [1](#)

`TxDb.Hsapiens.UCSC.hg19.knownGene-package`  
`(TxDb.Hsapiens.UCSC.hg19.knownGene),`

[1](#)