

# TxDB.Hsapiens.UCSC.hg38.refGene

August 27, 2025

---

TxDB.Hsapiens.UCSC.hg38.refGene

*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 2024-03-23 22:26:48 +0000 (Sat, 23 Mar 2024) and based on the hg38 genome based on the refGene table

## Author(s)

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

## See Also

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

## Examples

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

# Index

## \* **data**

`TxDb.Hsapiens.UCSC.hg38.refGene`, [1](#)

## \* **package**

`TxDb.Hsapiens.UCSC.hg38.refGene`, [1](#)

`transcripts`, [1](#)

`transcriptsBy`, [1](#)

`TxDb.Hsapiens.UCSC.hg38.refGene`, [1](#)

`TxDb.Hsapiens.UCSC.hg38.refGene-package`  
(`TxDb.Hsapiens.UCSC.hg38.refGene`),  
[1](#)