

# EpiTxDb.Hs.hg38

October 9, 2024

---

EpiTxDb.Hs.hg38

*Annotation package for EpiTxDb objects*

---

## Description

This package loads one or more EpiTxDb objects. Such EpiTxDb 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 `EpiTxDb.Hs.hg38.snoRNAdb` would be a EpiTxDb object of Homo sapiens data from snoRNAdb build based on the hg38 genome.

## Usage

```
EpiTxDb.Hs.hg38.RMBase(version = "1")  
EpiTxDb.Hs.hg38.snoRNAdb(version = "1")  
EpiTxDb.Hs.hg38.tRNAdb(version = "1")  
chain.rRNA.hg19Tohg38(version = "1")  
chain.rRNA.hg38Tohg19(version = "1")  
snoRNA.targets.hg38(version = "1")
```

## Arguments

`version` a character value defining a version. Versions available: "1".(default: version = "1")

## Value

a [EpiTxDb](#) object

**Author(s)**

Felix G M Ernst [aut]

**See Also**

- [modifications](#)
- [modificationsBy](#)
- [modifiedSeqsByTranscript](#)

**Examples**

`EpiTxDb.Hs.hg38.snoRNadb()`

# Index

chain.rRNA.hg19Tohg38  
    (EpiTxDb.Hs.hg38), [1](#)  
chain.rRNA.hg38Tohg19  
    (EpiTxDb.Hs.hg38), [1](#)

EpiTxDb, [1](#)  
EpiTxDb.Hs.hg38, [1](#)

modifications, [2](#)  
modificationsBy, [2](#)  
modifiedSeqsByTranscript, [2](#)

snoRNA.targets.hg38 (EpiTxDb.Hs.hg38), [1](#)