EpiTxDb.Hs.hg38

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`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)

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See Also

- `modifications`
- `modificationsBy`
- `modifiedSeqsByTranscript`

Examples

```
EpiTxDb.Hs.hg38.snoRNAdb()
```

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