EpiTxDb.Hs.hg38

August 27, 2020

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.hg28Tohg19(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]

1

See Also

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

Examples

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

Index