

EpiTxDb.Mm.mm10

April 2, 2025

EpiTxDb.Mm.mm10

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.Mm.mm10.tRNAdb would be a EpiTxDb object of Mus musculus data from tRNAdb build based on the mm10 genome.

Usage

```
EpiTxDb.Mm.mm10.RMBase(version = "1")
```

```
EpiTxDb.Mm.mm10.snoRNAdb(version = "1")
```

```
EpiTxDb.Mm.mm10.tRNAdb(version = "1")
```

```
snoRNA.targets.mm10(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.Mm.mm10.tRNAdb()`

Index

EpiTxDb, [/](#)

EpiTxDb.Mm.mm10, [1](#)

modifications, [/](#)

modificationsBy, [/](#)

modifiedSeqsByTranscript, [/](#)

snoRNA.targets.mm10 (EpiTxDb.Mm.mm10), [1](#)