

# EpiTxDb.Mm.mm10

December 4, 2024

---

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](#)