EpiTxDb.Mm.mm10

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`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)

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See Also

- modifications
- modificationsBy
- modifiedSeqsByTranscript

Examples

```
EpiTxDb.Mm.mm10.tRNAdb()
```

Index

EpiTxDb, *1*
EpiTxDb.Mm.mm10, 1

modifications, *2*
modificationsBy, *2*
modifiedSeqsByTranscript, *2*

snoRNA.targets.mm10 *(*EpiTxDb.Mm.mm10*)*, 1