

# FDb.UCSC.tRNAs

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FDb.UCSC.tRNAs

*Annotation package for FeatureDb object(s)*

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## Description

This package loads one or more FeatureDb objects. Such FeatureDb 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 TxDb.Hsapiens.UCSC.hg19.knownGene would be a TranscriptDb object, of Homo Sapiens data from UCSC build hg19 based on the knownGene Track.

## Note

This data package was made from resources at UCSC on 2012-09-10 13:15:01 -0700 (Mon, 10 Sep 2012) and based on the hg19 genome based on the tRNAs table

## Author(s)

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## See Also

[transcripts](#), [transcriptsBy](#)

## Examples

```
## load the library
library(FDb.UCSC.tRNAs)
## list the contents that are loaded into memory
ls('package:FDb.UCSC.tRNAs')
## show the db object that is loaded by calling it's name
FDb.Hsapiens.UCSC.hg19.tRNAs
```

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