

# TxDb.Hsapiens.UCSC.hg19.lincRNAsTranscripts

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TxDb.Hsapiens.UCSC.hg19.lincRNAsTranscripts  
*Annotation package for TxDb object(s)*

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## Description

This package loads one or more TxDb objects. Such TxDb 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 TxDb 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 2015-10-07 18:11:57 +0000 (Wed, 07 Oct 2015) and based on the hg19 genome based on the lincRNAsTranscripts table

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## See Also

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

## Examples

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