

# TxDB.Rnorvegicus.BioMart.igis

September 11, 2024

---

TxDB.Rnorvegicus.BioMart.igis

*Annotation package for TxDb object(s)*

---

## 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 BioMart on 2014-10-13 11:47:50 -0700 (Mon, 13 Oct 2014) and based on the IGIS 2.3 (Ensembl Genes 67)

## Author(s)

Gabriel Becker

## See Also

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

## Examples

```
## load the library
library(TxDB.Rnorvegicus.BioMart.igis)
## list the contents that are loaded into memory
ls('package:TxDB.Rnorvegicus.BioMart.igis')
## show the db object that is loaded by calling it's name
TxDb.Rnorvegicus.BioMart.igis
```

# Index

\* **data**

    TxDb.Rnorvegicus.BioMart.igis, [1](#)

\* **package**

    TxDb.Rnorvegicus.BioMart.igis, [1](#)

transcripts, [1](#)

transcriptsBy, [1](#)

TxDb.Rnorvegicus.BioMart.igis, [1](#)

TxDb.Rnorvegicus.BioMart.igis-package  
    (TxDb.Rnorvegicus.BioMart.igis),  
    [1](#)