RxDb.Ggallus.UCSC.galGal4.refGene

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An annotation package for RxDb objects

Description

This package loads one or more RxDb objects. Such RxDb 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 RxDb.Hsapiens.UCSC.hg19.knownGene would be a RxDb 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 2020-10-11 17:30:49 +0000 (Sun, 11 Oct 2020) and based on the galGal4 genome based on the refGene table

Author(s)

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See Also

transcripts, transcriptsBy

Examples

```r
## load the library
library(RxDB.Ggallus.UCSC.galGal4.refGene)
## list the contents that are loaded into memory
ls('package:RxDB.Ggallus.UCSC.galGal4.refGene')
## show the db object that is loaded by calling it's name
RxDB.Ggallus.UCSC.galGal4.refGene
```
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