

RmiR.hsa

September 24, 2025

RmiR.hsa_dbconn

Collect information from different miRNA targets databases

Description

A function to get a connection object to some miRNA targets databases.

Usage

```
RmiR.hsa_dbconn()  
RmiR.hsa_dbfile()
```

Details

RmiR.hsa_dbconn returns a connection object to some miRNA targets databases. **IMPORTANT:** Do not call [dbDisconnect](#) on the connection object returned by RmiR.hsa_dbconn or you will break all the [AnnDbObj](#) objects defined in this package!

RmiR.hsa_dbfile returns the path (character string) to the package annotation DB (this is an SQLite v3 file).

Value

RmiR.hsa_dbconn

A DBIConnection object representing an open connection to the miRNA targets database.

RmiR.hsa_dbfile

A character string with the path to the miRNA targets database.

See Also

[dbGetQuery](#), [dbConnect](#), [dbconn](#), [dbfile](#)

Examples

```
## List all the tables of the database:
dbListTables(RmiR.hsa_dbconn())

## Count the number of rows in the "targetscan" table:
dbGetQuery(RmiR.hsa_dbconn(), "SELECT COUNT(*) FROM targetscan")

## List the contents of the "mirbase" tables:
dbListFields(RmiR.hsa_dbconn(), "mirbase")

## List all targets of "hsa-miR-21" in miranda database:
dbGetQuery(RmiR.hsa_dbconn(),
  "SELECT * FROM tarbase where mature_miRNA='hsa-miR-21'")

## The connection object returned by RmiR.hsa_dbconn() was created with:
dbConnect(SQLite(), dbname=RmiR.hsa_dbfile(), cache_size=64000, synchronous=0)
```

Index

* **datasets**

RmiR.hsa_dbconn, [1](#)

* **utilities**

RmiR.hsa_dbconn, [1](#)

AnnDbObj, [1](#)

dbconn, [1](#)

dbConnect, [1](#)

dbDisconnect, [1](#)

dbfile, [1](#)

dbGetQuery, [1](#)

RmiR.hsa_dbconn, [1](#)

RmiR.hsa_dbfile (RmiR.hsa_dbconn), [1](#)