

# Package ‘ResultModelManager’

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ConnectionHandler	<i>ConnectionHandler</i>
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## Description

Class for handling DatabaseConnector:connection objects with consistent R6 interfaces for pooled and non-pooled connections. Allows a connection to cleanly be opened and closed and stored within class/object variables

## Value

DatabaseConnector Connection instance close Connection

boolean TRUE if connection is valid queryDb

boolean TRUE if connection is valid executeSql

## Public fields

connectionDetails DatabaseConnector connectionDetails object

con DatabaseConnector connection object

isActive Is connection active or not #'

snakeCaseToCamelCase (Optional) Boolean. return the results columns in camel case (default)

## Methods

### Public methods:

- [ConnectionHandler\\$new\(\)](#)
- [ConnectionHandler\\$dbms\(\)](#)
- [ConnectionHandler\\$tbl\(\)](#)
- [ConnectionHandler\\$renderTranslateSql\(\)](#)
- [ConnectionHandler\\$initConnection\(\)](#)
- [ConnectionHandler\\$getConnection\(\)](#)
- [ConnectionHandler\\$closeConnection\(\)](#)
- [ConnectionHandler\\$finalize\(\)](#)
- [ConnectionHandler\\$dbIsValid\(\)](#)
- [ConnectionHandler\\$queryDb\(\)](#)
- [ConnectionHandler\\$executeSql\(\)](#)
- [ConnectionHandler\\$queryFunction\(\)](#)
- [ConnectionHandler\\$executeFunction\(\)](#)
- [ConnectionHandler\\$clone\(\)](#)

**Method new():***Usage:*

```
ConnectionHandler$new(
  connectionDetails,
  loadConnection = TRUE,
  snakeCaseToCamelCase = TRUE
)
```

*Arguments:*

**connectionDetails** DatabaseConnector::connectionDetails class  
**loadConnection** Boolean option to load connection right away  
**snakeCaseToCamelCase** (Optional) Boolean. return the results columns in camel case (default) get dbms

**Method dbms():** Get the dbms type of the connection get table*Usage:*

```
ConnectionHandler$dbms()
```

**Method tbl():** get a dplyr table object (i.e. lazy loaded)*Usage:*

```
ConnectionHandler$tbl(table, databaseSchema = NULL)
```

*Arguments:*

**table** table name  
**databaseSchema** databaseSchema to which table belongs Render Translate Sql.

**Method renderTranslateSql():** Masked call to SqlRender*Usage:*

```
ConnectionHandler$renderTranslateSql(sql, ...)
```

*Arguments:*

**sql** Sql query string  
**...** Elipsis initConnection

**Method initConnection():** Load connection Get Connection*Usage:*

```
ConnectionHandler$initConnection()
```

**Method getConnection():** Returns connection for use with standard DatabaseConnector calls. Connects automatically if it isn't yet loaded*Usage:*

```
ConnectionHandler$getConnection()
```

**Method closeConnection():** Closes connection (if active) close Connection*Usage:*

```
ConnectionHandler$closeConnection()
```

**Method finalize():** Closes connection (if active) db Is Valid*Usage:*

```
ConnectionHandler$finalize()
```

**Method dbIsValid():** Masks call to DBI::dbIsValid. Returns False if connection is NULL

*Usage:*

```
ConnectionHandler$dbIsValid()
```

**Method queryDb():** query database and return the resulting data.frame

If environment variable LIMIT\_ROW\_COUNT is set Returned rows are limited to this value (no default) Limit row count is intended for web applications that may cause a denial of service if they consume too many resources.

*Usage:*

```
ConnectionHandler$queryDb(
  sql,
  snakeCaseToCamelCase = self$snakeCaseToCamelCase,
  overrideRowLimit = FALSE,
  ...
)
```

*Arguments:*

sql sql query string

snakeCaseToCamelCase (Optional) Boolean. return the results columns in camel case (default)

overrideRowLimit (Optional) Boolean. In some cases, where row limit is enforced on the system You may wish to ignore it.

... Additional query parameters

**Method executeSql():** execute set of database queries

*Usage:*

```
ConnectionHandler$executeSql(sql, ...)
```

*Arguments:*

sql sql query string

... Additional query parameters query Function

**Method queryFunction():** queryFunction that can be overridden with subclasses (e.g. use different base function or intercept query) Does not translate or render sql.

*Usage:*

```
ConnectionHandler$queryFunction(
  sql,
  snakeCaseToCamelCase = self$snakeCaseToCamelCase
)
```

*Arguments:*

sql sql query string

snakeCaseToCamelCase (Optional) Boolean. return the results columns in camel case (default) execute Function

**Method executeFunction():** exec query Function that can be overridden with subclasses (e.g. use different base function or intercept query) Does not translate or render sql.

*Usage:*

```
ConnectionHandler$executeFunction(sql)
```

*Arguments:*

sql sql query string

**Method clone():** The objects of this class are cloneable with this method.

*Usage:*

```
ConnectionHandler$clone(deep = FALSE)
```

*Arguments:*

deep Whether to make a deep clone.

---

createQueryNamespace    *Create query namespace*

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## Description

Create a QueryNamespace instance from either a connection handler or a connectionDetails object Allows construction with various options not handled by QueryNamespace\$new

Note - currently not supported is having multiple table prefixes for multiple table namespaces

## Usage

```
createQueryNamespace(
  connectionDetails = NULL,
  connectionHandler = NULL,
  usePooledConnection = FALSE,
  tableSpecification = NULL,
  resultModelSpecificationPath = NULL,
  tablePrefix = "",
  snakeCaseToCamelCase = TRUE,
  ...
)
```

## Arguments

resultModelSpecificationPath  
(optional) csv file or files for tableSpecifications - must conform to table spec format.

---

DataMigrationManager    *DataMigrationManager (DMM)*

---

## Description

R6 class for management of database migration

## Value

data frame all migrations, including file name, order and execution status Get connection handler

**Public fields**

migrationPath Path migrations exist in  
 databaseSchema Path migrations exist in  
 packageName packageName, can be null  
 tablePrefix packageName, can be null

**Methods****Public methods:**

- `DataMigrationManager$new()`
- `DataMigrationManager$migrationTableExists()`
- `DataMigrationManager$getMigrationsPath()`
- `DataMigrationManager$getStatus()`
- `DataMigrationManager$getConnectionHandler()`
- `DataMigrationManager$check()`
- `DataMigrationManager$executeMigrations()`
- `DataMigrationManager$isPackage()`
- `DataMigrationManager$finalize()`
- `DataMigrationManager$clone()`

**Method new():***Usage:*

```
DataMigrationManager$new(
  connectionDetails,
  databaseSchema,
  tablePrefix = "",
  migrationPath,
  packageName = NULL,
  migrationRegexp = .defaultMigrationRegexp
)
```

*Arguments:*

connectionDetails DatabaseConnector connection details object  
 databaseSchema Database Schema to execute on  
 tablePrefix Optional table prefix for all tables (e.g. plp, cm, cd etc)  
 migrationPath Path to location of migration sql files. If in package mode, this should just be a folder (e.g. "migrations") that lives in the location "sql/sql\_server" (and) other database platforms. If in folder model, the folder must include "sql\_server" in the relative path, (e.g if migrationPath = 'migrations' then the folder 'migrations/sql\_server' should exists)  
 packageName If in package mode, the name of the R package  
 migrationRegexp (Optional) regular expression pattern default is `(Migration_[0-9]+)-(.+)\.sql`  
 Migration table exists

**Method migrationTableExists():** Check if migration table is present in schema

*Usage:*

```
DataMigrationManager$migrationTableExists()
```

*Returns:* boolean Get path of migrations

**Method** getMigrationsPath(): Get path to sql migration files

*Usage:*

```
DataMigrationManager$getMigrationsPath(dbms = "sql server")
```

*Arguments:*

dbms Optionally specify the dbms that the migration fits under Get status of result model

**Method** getStatus(): Get status of all migrations (executed or not)

*Usage:*

```
DataMigrationManager$getStatus()
```

**Method** getConnectionHandler(): Return connection handler instance

*Usage:*

```
DataMigrationManager$getConnectionHandler()
```

*Returns:* ConnectionHandler instance Check migrations in folder

**Method** check(): Check if file names are valid for migrations Execute Migrations

*Usage:*

```
DataMigrationManager$check()
```

**Method** executeMigrations(): Execute any unexecuted migrations

*Usage:*

```
DataMigrationManager$executeMigrations(stopMigrationVersion = NULL)
```

*Arguments:*

stopMigrationVersion (Optional) Migrate to a specific migration number isPackage

**Method** isPackage(): is a package folder structure or not finalize

*Usage:*

```
DataMigrationManager$isPackage()
```

**Method** finalize(): close database connection

*Usage:*

```
DataMigrationManager$finalize()
```

**Method** clone(): The objects of this class are cloneable with this method.

*Usage:*

```
DataMigrationManager$clone(deep = FALSE)
```

*Arguments:*

deep Whether to make a deep clone.

## See Also

[ConnectionHandler](#) for information on returned class

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`deleteAllRowsForDatabaseId`*Delete all rows for database id*

---

**Description**

Delete all rows for database id

**Usage**

```
deleteAllRowsForDatabaseId(  
    connection,  
    schema,  
    tableName,  
    databaseId,  
    idIsInt = TRUE  
)
```

**Arguments**

<code>connection</code>	DatabaseConnector connection instance
<code>schema</code>	The schema on the postgres server where the results table exists
<code>tableName</code>	Database table name
<code>databaseId</code>	Results source database identifier
<code>idIsInt</code>	Identified is a numeric type? If not character is used

**Details**

Only PostgreSQL servers are supported.

---

`deleteAllRowsForPrimaryKey`*Delete results rows for primary key values from database server tables*

---

**Description**

Delete results rows for primary key values from database server tables

**Usage**

```
deleteAllRowsForPrimaryKey(connection, schema, tableName, keyValues)
```

**Arguments**

<code>connection</code>	DatabaseConnector connection instance
<code>schema</code>	The schema on the postgres server where the results table exists
<code>tableName</code>	Database table name
<code>keyValues</code>	Key values of results rows to be deleted



**Details**

Only PostgreSQL servers are supported.

---

generateSqlSchema	<i>Schema generator</i>
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**Description**

Take a csv schema definition and create a basic sql script with it.

**Usage**

```
generateSqlSchema(
  csvFilepath = NULL,
  schemaDefinition = NULL,
  sqlOutputPath = NULL,
  overwrite = FALSE
)
```

**Arguments**

csvFilepath	Path to schema file. Csv file must have the columns: "table_name", "column_name", "data_type", "is_required", "primary_key"
schemaDefinition	A schemaDefintion data.frame' with the columns: tableName, columnName, dataType, isRequired, primaryKey
sqlOutputPath	File to write sql to.
overwrite	Boolean - overwrite existing file?

**Value**

string containing the sql for the table

---

loadResultsDataModelSpecifications	<i>Get specifications from a given file path</i>
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**Description**

Get specifications from a given file path

**Usage**

```
loadResultsDataModelSpecifications(filePath)
```

**Arguments**

filePath	path to a valid csv file
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**Value**

A tibble data frame object with specifications

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PooledConnectionHandler

*Pooled Connection Handler*


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## Description

Transparently works the same way as a standard connection handler but stores pooled connections. Useful for long running applications that serve multiple concurrent requests.

## Super class

`ResultModelManager::ConnectionHandler` -> PooledConnectionHandler

## Methods

### Public methods:

- `PooledConnectionHandler$new()`
- `PooledConnectionHandler$initConnection()`
- `PooledConnectionHandler$dbms()`
- `PooledConnectionHandler$closeConnection()`
- `PooledConnectionHandler$queryFunction()`
- `PooledConnectionHandler$executeFunction()`
- `PooledConnectionHandler$clone()`

### Method new():

*Usage:*

`PooledConnectionHandler$new(...)`

*Arguments:*

... Elisis @seealso `ConnectionHandler` initialize pooled db connection

**Method initConnection():** Overrides ConnectionHandler Call get dbms

*Usage:*

`PooledConnectionHandler$initConnection()`

**Method dbms():** Get the dbms type of the connection Close Connection

*Usage:*

`PooledConnectionHandler$dbms()`

**Method closeConnection():** Overrides ConnectionHandler Call query Function

*Usage:*

`PooledConnectionHandler$closeConnection()`

**Method queryFunction():** Overrides ConnectionHandler Call. Does not translate or render sql.

*Usage:*

```
PooledConnectionHandler$queryFunction(
  sql,
  snakeCaseToCamelCase = self$snakeCaseToCamelCase
)
```

*Arguments:*

sql sql query string

snakeCaseToCamelCase (Optional) Boolean. return the results columns in camel case (default) query Function

**Method** executeFunction(): Overrides ConnectionHandler Call. Does not translate or render sql.

*Usage:*

PooledConnectionHandler\$executeFunction(sql)

*Arguments:*

sql sql query string

**Method** clone(): The objects of this class are cloneable with this method.

*Usage:*

PooledConnectionHandler\$clone(deep = FALSE)

*Arguments:*

deep Whether to make a deep clone.

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QueryNamespace	<i>QueryNamespace</i>
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## Description

Given a results specification and ConnectionHandler instance - this class allow queries to be namespaced within any tables specified within a list of pre-determined tables. This allows the encapsulation of queries, using specific table names in a consistent manner that is straightforward to maintain over time.

## Public fields

tablePrefix tablePrefix to use

## Methods

### Public methods:

- [QueryNamespace\\$new\(\)](#)
- [QueryNamespace\\$setConnectionHandler\(\)](#)
- [QueryNamespace\\$getConnectionHandler\(\)](#)
- [QueryNamespace\\$addReplacementVariable\(\)](#)
- [QueryNamespace\\$addTableSpecification\(\)](#)
- [QueryNamespace\\$render\(\)](#)
- [QueryNamespace\\$queryDb\(\)](#)
- [QueryNamespace\\$executeSql\(\)](#)
- [QueryNamespace\\$getVars\(\)](#)
- [QueryNamespace\\$clone\(\)](#)

**Method** new(): initialize class

*Usage:*

```
QueryNamespace$new(
  connectionHandler = NULL,
  tableSpecification = NULL,
  tablePrefix = "",
  ...
)
```

*Arguments:*

**connectionHandler** ConnectionHandler instance @seealso [ConnectionHandler](#)  
**tableSpecification** tableSpecification data.frame  
**tablePrefix** constant string to prefix all tables with  
 ... additional replacement variables e.g. database\_schema, vocabulary\_schema etc  
 Set Connection Handler

**Method** `setConnectionHandler()`: set connection handler object for object

*Usage:*

```
QueryNamespace$setConnectionHandler(connectionHandler)
```

*Arguments:*

**connectionHandler** ConnectionHandler instance Get connection handler

**Method** `getConnectionHandler()`: get connection handler object or throw error if not set

*Usage:*

```
QueryNamespace$getConnectionHandler()
```

**Method** `addReplacementVariable()`: add a variable to automatically be replaced in query strings (e.g. @database\_schema.@table\_name becomes 'database\_schema.table\_1')

*Usage:*

```
QueryNamespace$addReplacementVariable(key, value, replace = FALSE)
```

*Arguments:*

**key** variable name string (without @) to be replaced, eg. "table\_name"  
**value** atomic value for replacement  
**replace** if a variable of the same key is found, overwrite it add table specification

**Method** `addTableSpecification()`: add a variable to automatically be replaced in query strings (e.g. @database\_schema.@table\_name becomes 'database\_schema.table\_1')

*Usage:*

```
QueryNamespace$addTableSpecification(
  tableSpecification,
  useTablePrefix = TRUE,
  tablePrefix = self$tablePrefix,
  replace = TRUE
)
```

*Arguments:*

**tableSpecification** table specification data.frame conforming to column names table-Name, columnName, dataType and primaryKey  
**useTablePrefix** prefix the results with the tablePrefix (TRUE)  
**tablePrefix** prefix string - defaults to class variable set during initialization

replace replace existing variables of the same name Render

**Method** render(): Call to SqlRender::render replacing names stored in this class

*Usage:*

QueryNamespace\$render(sql, ...)

*Arguments:*

sql query string

... additional variables to be passed to SqlRender::render - will overwrite anything in namespace query Sql

**Method** queryDb(): Call to

*Usage:*

QueryNamespace\$queryDb(sql, ...)

*Arguments:*

sql query string

... additional variables to send to SqlRender::render execute Sql

**Method** executeSql(): Call to execute sql within namespaced queries

*Usage:*

QueryNamespace\$executeSql(sql, ...)

*Arguments:*

sql query string

... additional variables to send to SqlRender::render get vars

**Method** getVars(): returns full list of variables that will be replaced

*Usage:*

QueryNamespace\$getVars()

**Method** clone(): The objects of this class are cloneable with this method.

*Usage:*

QueryNamespace\$clone(deep = FALSE)

*Arguments:*

deep Whether to make a deep clone.

## Examples

```
## Not run:
library(ResultModelManager)
connectionHandler <- ConnectionHandler$new(connectionDetails = )

tableSpecification <- data.frame(
  tableName = "cohort",
  columnName = c(
    "cohort_definition_id",
    "cohort_name",
    "json",
    "sql"
  ),
),
primaryKey = c(TRUE, FALSE, FALSE, FALSE),
dataTypes = c("int", "varchar", "varchar", "varchar")
```

```
)

cohortNamespace <- QueryNamespace$new(
  connectionHandler = connectionHandler,
  tableSpecification = tableSpecification,
  result_schema = "main",
  tablePrefix = "cd_"
)
sql <- "SELECT * FROM @result_schema.@cohort WHERE cohort_id = @cohort_id"
# Returns : "SELECT * FROM main.cd_cohort WHERE cohort_id = @cohort_id"
print(cohortNamespace$render(sql))
# Returns query result
result <- cohortNamespace$querySql(sql, cohort_id = 1)

## End(Not run)
```

---

unzipResults	<i>Unzips a results.zip file and enforces standards required by uploadResults</i>
--------------	---

---

**Description**

This function will unzip the zipFile to the resultsFolder and assert that the file results-DataModelSpecification.csv exists in the resultsFolder to ensure that it will work with uploadResults

**Usage**

```
unzipResults(zipFile, resultsFolder)
```

**Arguments**

- zipFile            The location of the .zip file that holds the results to upload
- resultsFolder    The folder to use when unzipping the .zip file. If this folder does not exist, this function will attempt to create the folder.

---

uploadResults	<i>Upload results to the database server.</i>
---------------	---

---

**Description**

Requires the results data model tables have been created using following the specifications, @seealso [generateSqlSchema](#) function.

Results files should be in the snake\_case format for table headers and not camelCase

Set the POSTGRES\_PATH environmental variable to the path to the folder containing the psql executable to enable bulk upload (recommended).

**Usage**

```
uploadResults(
  connection = NULL,
  connectionDetails = NULL,
  schema,
  resultsFolder,
  tablePrefix = "",
  forceOverWriteOfSpecifications = FALSE,
  purgeSiteDataBeforeUploading = TRUE,
  databaseIdentifierFile = "cdm_source_info.csv",
  runCheckAndFixCommands = FALSE,
  warnOnMissingTable = TRUE,
  specifications
)
```

**Arguments**

- |                                       |   |
|---------------------------------------|---|
| <b>connection</b>                     | An object of type <code>connection</code> as created using the <a href="#">connect</a> function in the DatabaseConnector package. Can be left NULL if <code>connectionDetails</code> is provided, in which case a new connection will be opened at the start of the function, and closed when the function finishes.  |
| <b>connectionDetails</b>              | An object of type <code>connectionDetails</code> as created using the <a href="#">createConnectionDetails</a> function in the DatabaseConnector package.  |
| <b>schema</b>                         | The schema on the postgres server where the tables have been created.   |
| <b>resultsFolder</b>                  | The path to the folder containing the results to upload. See <a href="#">unzipResults</a> for more information.   |
| <b>tablePrefix</b>                    | String to prefix table names with - default is empty string   |
| <b>forceOverWriteOfSpecifications</b> | If TRUE, specifications of the phenotypes, cohort definitions, and analysis will be overwritten if they already exist on the database. Only use this if these specifications have changed since the last upload.  |
| <b>purgeSiteDataBeforeUploading</b>   | If TRUE, before inserting data for a specific databaseId all the data for that site will be dropped. This assumes the results folder contains the full data for that data site.   |
| <b>databaseIdentifierFile</b>         | File contained that references databaseId field (used when <code>purgeSiteDataBeforeUploading == TRUE</code> ). You may specify a relative path for the <code>cdmSourceFile</code> and the function will assume it resides in the <code>resultsFolder</code> . Alternatively, you can provide a path outside of the <code>resultsFolder</code> for this file. |
| <b>runCheckAndFixCommands</b>         | If TRUE, the upload code will attempt to fix column names, data types and duplicate rows. This parameter is kept for legacy reasons - it is strongly recommended that you correct errors in your results where those results are assembled instead of relying on this option to try and fix it during upload.   |
| <b>warnOnMissingTable</b>             | Boolean, print a warning if a table file is missing.  |
| <b>specifications</b>                 | A tibble data frame object with specifications.   |