Package 'ResultModelManager'

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Description Database data model management utilities for OHDSI packages.

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   ParallelLogger,
   checkmate,
   DBI,
   pool,
   readr,
   zip,
   dplyr,
   dbplyr,
   rlang,
   lubridate,
   fastmap

Suggests testthat (>= 3.0.0),
   RSQLite,
   duckdb,
   RPostgres,
   withr,
   knitr,
   rmarkdown,
   keyring,
   devtools,
   pkgdown,
   remotes,
   styler,
   Andromeda

Config/testthat/edition 3

R topics documented:

ConnectionHandler   *ConnectionHandler*

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 overriden 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 overriden 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*

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 names-
 paces

Usage

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

Arguments

`connectionDetails`

   An object of type `connectionDetails` as created using the `createConnectionDetails`
   function in the DatabaseConnector package.

`connectionHandler`

   ResultModelManager ConnectionHandler or PooledConnectionHandler in-
   stance

`usePooledConnection`

   Use Pooled database connection instead of standard DatabaseConnector
   single connection.

```
tableSpecification
```
                    Table specfication data.frame
```
resultModelSpecificationPath
```
                    (optional) csv file or files for tableSpecifications - must conform to table
                    spec format.

```
tablePrefix
```
          String to prefix table names with - default is empty string
```
snakeCaseToCamelCase
```
                    convert snakecase results to camelCase field names (TRUE by default)

`...`   Elipsis - use for any additional string keys to replace

---

```
createResultExportManager
```
Create Result Export Manager

Description

For a give table specification file, create an export manager instance for creating results
data sets that conform to the data model.

This checks that, at export time, internal validity is assured for the data (e.g. primary keys
are valid, data types are compatible

In addition this utility will create a manifest object that can be used to maintain the validity
of data.

If an instance of a DataMigrationManager is present and available a packageVersion refer-
ence (where applicable) and migration set will be referenced. Allowing data to be imported
into a database schema at a specific version.

Usage

```
createResultExportManager(
  tableSpecification,
  exportDir,
  minCellCount = getOption("ohdsi.minCellCount", default = 5),
  databaseId = NULL
)
```

Arguments

```
tableSpecification
```
                    Table specification data.frame

```
exportDir
```
            Directory files are being exported to

```
minCellCount
```
         Minimum cell count - reccomended that you set with options("ohdsi.minCellCount"
                    = count) in all R projects. Default is 5

```
databaseId
```
           database identifier - required when exporting according to many specs

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 tablePrefix, can be empty character vector

packageTablePrefix packageTablePrefix, can be empty character vector

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 = "",
  packageTablePrefix = "",
  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)

packageTablePrefix A table prefix when used in conjunction with other package results
schema, e.g. "cd_", "sccs_", "plp_", "cm_"

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 'migra-
tions/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

`deleteAllRowsForDatabaseId`
Delete all rows for database id

Description

Delete all rows for database id

Usage

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

Arguments

connection	DatabaseConnector connection instance
schema	The schema on the postgres server where the results table exists
tableName	Database table name
databaseId	Results source database identifier
idIsInt	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

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

Details

Only PostgreSQL servers are supported.

---

generateSqlSchema   *Schema generator*

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 schemaDefintiion 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

---

grantTablePermissions *Grant Table Permissions*

Description

Grant a given permission for all tables on a given tableSpecification

Very useful if you're hosting studies on data.ohdsi.org or other postgresql instances

NOTE: only tested on postgresql, users' of other platforms may have Sql translation issues

Usage

```
grantTablePermissions(
  connectionDetails = NULL,
  connection = NULL,
  tableSpecification,
  databaseSchema,
  tablePrefix = "",
  permissions = "SELECT",
  user
)
```

Arguments

connectionDetails

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package.

connection   DatabaseConnector connection instance

tableSpecification

data.frame conforming to table spec (must contain tableName field)

databaseSchema   database schema to run this on

tablePrefix   String to prefix table names with - default is empty string

permissions   permissions to generate must be one of SELECT, INSERT, DELETE or UPDATE

user   database user to grant permissions to

---

loadResultsDataModelSpecifications

Get specifications from a given file path

Description

Get specifications from a given file path

Usage

loadResultsDataModelSpecifications(filePath)

Arguments

filePath   path to a valid csv file

Value

A tibble data frame object with specifications

PooledConnectionHandler

Pooled Connection Handler

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(..., poolArgs = NULL)

    Arguments:

... Elisis @seealsoConnectionHandler

poolArgs Optional arguments to call pool::dbPool overrides default usage of connectionDetails 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.

QueryNamespace   *QueryNamespace*

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 striaghtforward 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$getVars()
- QueryNamespace$finalize()
- QueryNamespace$clone()

Method new(): initialize class

Usage:

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

Arguments:

connectionHandler ConnectionHandler instance @seealsoConnectionHandler

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 obeject 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 Destruct object

Usage:

`QueryNamespace$getVars()`

Method `finalize()`: Close connections etc

Usage:

`QueryNamespace$finalize()`

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),
  dataType = c("int", "varchar", "varchar", "varchar")
)

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

## End(Not run)
```

ResultExportManager   *Result Set Export Manager*

Description

EXPERIMENTAL - this feature is still in design stage and it is not reccomended that you implement this for your package at this stage. Utility for simplifying export of results to files from sql queries

Note that this utility is not strictly thread safe though seperate processes can export separate tables without issue. When exporting a the same table across multiple threads primary key checks may create issues.

Public fields

exportDir direcotry path to export files to Init

Methods

Public methods:

- ResultExportManager$new()
- ResultExportManager$getTableSpec()
- ResultExportManager$getMinColValues()
- ResultExportManager$checkRowTypes()
- ResultExportManager$listTables()
- ResultExportManager$checkPrimaryKeys()
- ResultExportManager$exportDataFrame()
- ResultExportManager$exportQuery()
- ResultExportManager$getManifestList()
- ResultExportManager$writeManifest()
- ResultExportManager$clone()

Method new(): Create a class for exporting results from a study in a standard, consistend manner

Usage:

```
ResultExportManager$new(
  tableSpecification,
  exportDir,
  minCellCount = getOption("ohdsi.minCellCount", default = 5),
  databaseId = NULL
)
```

Arguments:

tableSpecification Table specification data.frame

exportDir Directory files are being exported to

minCellCount Minimum cell count - reccomended that you set with options("ohdsi.minCellCount" = count) in all R projects. Default is 5

databaseId database identifier - required when exporting according to many specs get table spec

Method getTableSpec(): Get specification of table

Usage:

```
ResultExportManager$getTableSpec(exportTableName)
```

Arguments:

exportTableName table name Get min col values

Method getMinColValues(): Columns to convert to minimum for a given table name

Usage:

```
ResultExportManager$getMinColValues(rows, exportTableName)
```

Arguments:

rows data.frame of rows

exportTableName stering table name - must be defined in spec Check row types

Method checkRowTypes(): Check types of rows before exporting

Usage:

```
ResultExportManager$checkRowTypes(rows, exportTableName)
```

Arguments:

rows data.frame of rows to export

exportTableName table name List tables

Method listTables(): list all tables in schema Check primary keys of exported data

Usage:

```
ResultExportManager$listTables()
```

Method checkPrimaryKeys(): Checks to see if the rows conform to the valid primary keys If the same table has already been checked in the life of this object set "invalidateCache" to TRUE as the keys will be cached in a temporary file on disk.

Usage:

```
ResultExportManager$checkPrimaryKeys(
    rows,
    exportTableName,
    invalidateCache = FALSE
)
```

Arguments:

rows data.frame to export

exportTableName Table name (must be in spec)

invalidateCache logical - if starting a fresh export use this to delete cache of primary
    keys Export data frame

Method exportDataFrame(): This method is intended for use where exporting a data.frame and not a query from a rdbms table For example, if you perform a transformation in R this method will check primary keys, min cell counts and data types before writing the file to according to the table spec

Usage:

```
ResultExportManager$exportDataFrame(rows, exportTableName, append = FALSE)
```

Arguments:

rows Rows to export

exportTableName Table name

append logical - if true will append the result to a file, otherwise the file will be over-written Export Data table with sql query

Method `exportQuery()`:   Writes files in batch to stop overflowing system memory Checks primary keys on write Checks minimum cell count

Usage:
```
ResultExportManager$exportQuery(
  connection,
  sql,
  exportTableName,
  transformFunction = NULL,
  transformFunctionArgs = list(),
  append = FALSE,
  ...
)
```

Arguments:

connection DatabaseConnector connection instance

sql OHDSI sql string to export tables

exportTableName Name of table to export (in snake_case format)

transformFunction (optional) transformation of the data set callback. must take two paramters - rows and pos

```
    Following this transformation callback, results will be verified against data model,
     Primary keys will be checked and minCellValue rules will be enforced
```

transformFunctionArgs arguments to be passed to the transformation function

append Logical add results to existing file, if FALSE (default) creates a new file and removes primary key validation cache

... extra parameters passed to sql get manifest list

Method `getManifestList()`:   Create a meta data set for each collection of result files with sha256 has for all files

Usage:
```
ResultExportManager$getManifestList(
  packageName = NULL,
  packageVersion = NULL,
  migrationsPath = NULL,
  migrationRegexp = .defaultMigrationRegexp
)
```

Arguments:

packageName if an R analysis package, specify the name

packageVersion if an analysis package, specify the version

migrationsPath path to sql migrations (use top level folder (e.g. sql/sql_server/migrations)

migrationRegexp (optional) regular expression to search for sql files. It is not rec-comended to change the default. Write manifest

Method `writeManifest()`:   Write manifest json

Usage:
```
ResultExportManager$writeManifest(...)
```

Arguments:

... @seealso getManifestList

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

Usage:

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

Arguments:

deep Whether to make a deep clone.

---

unzipResults	*Unzips a results.zip file and enforces standards required by* `uploadResults`

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	*Upload results to the database server.*

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,
  purgeDataModel = FALSE,
  specifications
)
```

Arguments

connection	An object of type `connection` as created using the `connect` function in the DatabaseConnector package. Can be left NULL if `connectionDetails` is provided, in which case a new connection will be opened at the start of the function, and closed when the function finishes.
connectionDetails	
	An object of type `connectionDetails` as created using the `createConnectionDetails` function in the DatabaseConnector package.
schema	The schema on the postgres server where the tables have been created.
resultsFolder	The path to the folder containing the results to upload. See `unzipResults` for more information.
tablePrefix	String to prefix table names with - default is empty string
forceOverWriteOfSpecifications	
	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.
purgeSiteDataBeforeUploading	
	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.
databaseIdentifierFile	
	File contained that references databaseId field (used when purgeSite-DataBeforeUploading == TRUE). You may specify a relative path for the cdmSourceFile and the function will assume it resides in the results-Folder. Alternatively, you can provide a path outside of the resultsFolder for this file.
runCheckAndFixCommands	
	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.
warnOnMissingTable	
	Boolean, print a warning if a table file is missing.

purgeDataModel   This function will purge all data from the tables in the specification prior
                 to upload. Use with care. If interactive this will require further input.

specifications   A tibble data frame object with specifications.