

# Package ‘AlendronateVsRaloxifene’

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**Type** Package

**Title** Alendronate versus Raloxifene and the risk of Hip Fracture

**Version** 0.2.0

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**Description** More about what it does (maybe more than one line).

**Depends** DatabaseConnector (>= 1.10.0)

**Imports** SqlRender (>= 1.3.0),  
RJDBC,  
FeatureExtraction (>= 1.0.3),  
CohortMethod (>= 2.2.2),  
EmpiricalCalibration (>= 1.2.0),  
OhdsiSharing (>= 0.1.1),  
Cyclops (>= 1.2.2),  
rmarkdown,  
ggplot2,  
ff,  
ffbase

**Suggests** OhdsiRTools (>= 1.3.0)

**License** Apache License 2.0

**LazyData** TRUE

**RoxygenNote** 6.0.1

## R topics documented:

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AlendronateVsRaloxifene  
*AlendronateVsRaloxifene*

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## Description

AlendronateVsRaloxifene

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assessFeasibility	<i>Execute OHDSI Alendronate Vs Raloxifene study feasibility assessment</i>
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## Description

Execute OHDSI Alendronate Vs Raloxifene study feasibility assessment

## Usage

```
assessFeasibility(connectionDetails, cdmDatabaseSchema,
  workDatabaseSchema = cdmDatabaseSchema,
  studyCohortTable = "ohdsi_alendronate_raloxifene",
  oracleTempSchema = workDatabaseSchema, outputFolder)
```

## Arguments

connectionDetails	An object of type connectionDetails as created using the <a href="#">createConnectionDetails</a> function in the DatabaseConnector package.
cdmDatabaseSchema	Schema name where your patient-level data in OMOP CDM format resides. Note that for SQL Server, this should include both the database and schema name, for example 'cdm_data.dbo'.
workDatabaseSchema	Schema name where intermediate data can be stored. You will need to have write privileges in this schema. Note that for SQL Server, this should include both the database and schema name, for example 'cdm_data.dbo'.
studyCohortTable	The name of the table that will be created in the work database schema. This table will hold the exposure and outcome cohorts used in this study.
oracleTempSchema	Should be used in Oracle to specify a schema where the user has write privileges for storing temporary tables.
outputFolder	Name of local folder to place results; make sure to use forward slashes (/). Do not use a folder on a network drive since this greatly impacts performance.

## Details

This function executes the OHDSI Alendronate Vs Raloxifene study feasibility assessment.

## Examples

```
## Not run:
connectionDetails <- createConnectionDetails(dbms = "postgresql",
                                             user = "joe",
                                             password = "secret",
                                             server = "myserver")

assessFeasibility(connectionDetails,
                  cdmDatabaseSchema = "cdm_data",
                  workDatabaseSchema = "results",
                  studyCohortTable = "ohdsi_alendronate_raloxifene",
                  oracleTempSchema = NULL,
                  outputFolder = "c:/temp/feasibility_results")

## End(Not run)
```

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createAnalysesDetails *Create the analyses details*

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## Description

Create the analyses details

## Usage

```
createAnalysesDetails(connectionDetails, cdmDatabaseSchema, workFolder)
```

## Arguments

connectionDetails	An object of type connectionDetails as created using the <a href="#">createConnectionDetails</a> function in the DatabaseConnector package.
cdmDatabaseSchema	Schema name where your patient-level data in OMOP CDM format resides. Note that for SQL Server, this should include both the database and schema name, for example 'cdm_data.dbo'.
workFolder	Name of local folder to place results; make sure to use forward slashes (/)

## Details

This function creates files specifying the analyses that will be performed.

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createCohorts	<i>Create the exposure and outcome cohorts</i>
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### Description

Create the exposure and outcome cohorts

### Usage

```
createCohorts(connectionDetails, cdmDatabaseSchema, workDatabaseSchema,
  studyCohortTable = "ohdsi_alendronate_raloxifene", oracleTempSchema,
  outputFolder)
```

### Arguments

connectionDetails	An object of type connectionDetails as created using the <a href="#">createConnectionDetails</a> function in the DatabaseConnector package.
cdmDatabaseSchema	Schema name where your patient-level data in OMOP CDM format resides. Note that for SQL Server, this should include both the database and schema name, for example 'cdm_data.dbo'.
workDatabaseSchema	Schema name where intermediate data can be stored. You will need to have write privileges in this schema. Note that for SQL Server, this should include both the database and schema name, for example 'cdm_data.dbo'.
studyCohortTable	The name of the table that will be created in the work database schema. This table will hold the exposure and outcome cohorts used in this study.
oracleTempSchema	Should be used in Oracle to specify a schema where the user has write privileges for storing temporary tables.
outputFolder	Name of local folder to place results; make sure to use forward slashes (/)

### Details

This function will create the exposure and outcome cohorts following the definitions included in this package.

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createMetaData	<i>Create metadata file</i>
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### Description

Create metadata file

### Usage

```
createMetaData(connectionDetails, cdmDatabaseSchema, exportFolder)
```

Arguments

- connectionDetails

An object of type connectionDetails as created using the [createConnectionDetails](#) function in the DatabaseConnector package.
- cdmDatabaseSchema

Schema name where your patient-level data in OMOP CDM format resides. Note that for SQL Server, this should include both the database and schema name, for example 'cdm\_data.dbo'.
- exportFolder

The name of the folder where the metadata file should be created.

Details

Creates a file containing metadata about the source data (taken from the cdm\_source table) and R package versions.

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createTableAndFigures	Create tables and figures
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Description

Create tables and figures

Usage

```
createTableAndFigures(exportFolder)
```

Arguments

- exportFolder

The path to the export folder containing the results.

Details

Creates tables and figures for viewing and interpreting the results. Requires that the [execute](#) function has completed first.

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execute	Execute OHDSI Keppra and the Risk of Angioedema study
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Description

Execute OHDSI Keppra and the Risk of Angioedema study

Usage

```
execute(connectionDetails, cdmDatabaseSchema,
workDatabaseSchema = cdmDatabaseSchema,
studyCohortTable = "ohdsi_alendronate_raloxifene",
oracleTempSchema = workDatabaseSchema, outputFolder, createCohorts = TRUE,
runAnalyses = TRUE, packageResults = TRUE, maxCores = 4)
```

## Arguments

connectionDetails	An object of type connectionDetails as created using the <a href="#">createConnectionDetails</a> function in the DatabaseConnector package.
cdmDatabaseSchema	Schema name where your patient-level data in OMOP CDM format resides. Note that for SQL Server, this should include both the database and schema name, for example 'cdm_data.dbo'.
workDatabaseSchema	Schema name where intermediate data can be stored. You will need to have write privileges in this schema. Note that for SQL Server, this should include both the database and schema name, for example 'cdm_data.dbo'.
studyCohortTable	The name of the table that will be created in the work database schema. This table will hold the exposure and outcome cohorts used in this study.
oracleTempSchema	Should be used in Oracle to specify a schema where the user has write privileges for storing temporary tables.
outputFolder	Name of local folder to place results; make sure to use forward slashes (/). Do not use a folder on a network drive since this greatly impacts performance.
createCohorts	Create the studyCohortTable table with the exposure and outcome cohorts?
runAnalyses	Perform the cohort method analyses?
packageResults	Package the results for sharing?
maxCores	How many parallel cores should be used? If more cores are made available this can speed up the analyses.

## Details

This function executes the OHDSI Keppra and the Risk of Angioedema study.

## Examples

```
## Not run:
connectionDetails <- createConnectionDetails(dbms = "postgresql",
                                             user = "joe",
                                             password = "secret",
                                             server = "myserver")

execute(connectionDetails,
         cdmDatabaseSchema = "cdm_data",
         workDatabaseSchema = "results",
         studyCohortTable = "ohdsi_alendronate_raloxifene",
         oracleTempSchema = NULL,
         outputFolder = "c:/temp/study_results",
         maxCores = 4)

## End(Not run)
```

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packageResults	<i>Package the results for sharing with OHDSI researchers</i>
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**Description**

Package the results for sharing with OHDSI researchers

**Usage**

```
packageResults(connectionDetails, cdmDatabaseSchema, outputFolder,
  minCellCount = 5)
```

**Arguments**

connectionDetails	An object of type connectionDetails as created using the <a href="#">createConnectionDetails</a> function in the DatabaseConnector package.
cdmDatabaseSchema	Schema name where your patient-level data in OMOP CDM format resides. Note that for SQL Server, this should include both the database and schema name, for example 'cdm_data.dbo'.
outputFolder	Name of local folder to place results; make sure to use forward slashes (/)
minCellCount	The minimum number of subjects contributing to a count before it can be included in the results.

**Details**

This function packages the results.

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submitResults	<i>Submit the study results to the study coordinating center</i>
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**Description**

Submit the study results to the study coordinating center

**Usage**

```
submitResults(exportFolder, key, secret)
```

**Arguments**

exportFolder	The path to the folder containing the StudyResults.zip file.
key	The key string as provided by the study coordinator
secret	The secret string as provided by the study coordinator

**Details**

This will upload the file StudyResults.zip to the study coordinating center using Amazon S3. This requires an active internet connection.

**Value**

TRUE if the upload was successful.

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writeReport	<i>Write a report summarizing all the results for a single database</i>
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**Description**

Write a report summarizing all the results for a single database

**Usage**

```
writeReport(exportFolder, outputFile)
```

**Arguments**

exportFolder	The path to the export folder containing the results.
outputFile	The name of the .docx file that will be created.

**Details**

Requires that the [createTableAndFigures](#) has been executed first.



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