# Package 'CohortDiagnostics'

March 13, 2020

```
Type Package
Title Diagnostics for OHDSI studies
Version 0.1.0
Date 2020-03-12
Description Diagnostics for studies that use the OMOP Common Data Model and the OHDSI tools.
Depends DatabaseConnector
Imports SqlRender,
     ggplot2,
     ROhdsiWebApi,
     ParallelLogger,
     readr,
     tibble,
     dplyr,
     RJSONIO,
     FeatureExtraction,
Suggests testthat,
     knitr,
     rmarkdown,
     DT,
     plotly,
     shiny,
     shinydashboard,
     RColorBrewer,
     VennDiagram,
     htmltools
License Apache License
VignetteBuilder knitr
URL https://ohdsi.github.io/CohortDiagnostics, https:
     //github.com/OHDSI/CohortDiagnostics
BugReports https://github.com/OHDSI/CohortDiagnostics/issues
RoxygenNote 7.0.2
Encoding UTF-8
```

2 breakDownIndexEvents

# R topics documented:

Index		28
	runCohortDiagnosticsUsingExternalCounts	25
	runCohortDiagnostics	
	preMergeDiagnosticsFiles	
	plotincidenceRate	21
	launchDiagnosticsExplorer	
	launchCohortExplorer	
	instantiateCohortSet	
	instantiateCohort	17
	getTimeDistributions	16
	getInclusionStatisticsFromFiles	15
	getInclusionStatistics	14
	getIncidenceRate	13
	getCohortCounts	12
	getCohortCharacteristics	11
	findOrphanConcepts	10
	findCohortOrphanConcepts	
	findCohortIncludedSourceConcepts	7
	createConceptCountsTable	6
	createCohortTable	
	computeCohortOverlap	
	compareCohortCharacteristics	
	CohortDiagnostics	
	breakDownIndexEvents	2

breakDownIndexEvents Break down index events

# Description

For the concepts included in the index event definition, count how often they are encountered at the cohort index date.

```
breakDownIndexEvents(
  connectionDetails = NULL,
  connection = NULL,
  cdmDatabaseSchema,
  oracleTempSchema = NULL,
  cohortDatabaseSchema = cdmDatabaseSchema,
  cohortTable = "cohort",
  baseUrl = NULL,
  webApiCohortId = NULL,
  cohortJson = NULL,
  cohortSql = NULL,
  cohortId = cohortId
```

CohortDiagnostics 3

### **Arguments**

connectionDetails

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package. Can be left NULL if connection

is provided.

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.

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'.

oracleTempSchema

Should be used in Oracle to specify a schema where the user has write privileges for storing temporary tables.

cohortDatabaseSchema

Schema name where your cohort table resides. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'.

cohortTable Name of the cohort table.

baseUrl The base URL for the WebApi instance, for example: "http://server.org:80/WebAPI".

Needn't be provided if cohortJson and cohortSql are provided.

 $web {\tt API instance}. \ Needn't be \ provided \ if \ cohort {\tt Json}$ 

and cohortSql are provided.

cohortJson A character string containing the JSON of a cohort definition. Needn't be pro-

vided if baseUrl and cohortId are provided.

cohortSql The OHDSI SQL representation of the same cohort definition. Needn't be pro-

vided if baseUrl and cohortId are provided.

cohortId The cohort definition ID used to reference the cohort in the cohort table.

### Value

A data frame with concepts, and per concept the count of how often the concept was encountered at the index date.

CohortDiagnostics CohortDiagnostics

# **Description**

CohortDiagnostics

compareCohortCharacteristics

Compare cohort characteristics

# Description

Compare the characteristics of two cohorts, computing the standardized difference of the mean.

# Usage

```
compare Cohort Characteristics (characteristics 1, characteristics 2)\\
```

# **Arguments**

characteristics1

Characteristics of the first cohort, as created using the getCohortCharacteristics function.

characteristics2

 $Characteristics \ of the second \ cohort, as \ created \ using \ the \ {\tt getCohortCharacteristics} \ function.$ 

# Value

A data frame comparing the characteristics of the two cohorts.

computeCohortOverlap Compute overlap between two cohorts

# Description

Computes the overlap between a target and a comparator cohort.

```
computeCohortOverlap(
  connectionDetails = NULL,
  connection = NULL,
  cohortDatabaseSchema,
  cohortTable = "cohort",
  targetCohortId,
  comparatorCohortId
)
```

createCohortTable 5

### **Arguments**

connectionDetails

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package. Can be left NULL if connection is provided.

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.

cohortDatabaseSchema

Schema name where your cohort table resides. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'.

cohortTable Name of the cohort table.

targetCohortId The cohort definition ID used to reference the target cohort in the cohort table. comparatorCohortId

The cohort definition ID used to reference the comparator cohort in the cohort table.

### Value

A data frame with overlap statistics.

# **Description**

This function creates an empty cohort table. Optionally, additional empty tables are created to store statistics on the various inclusion criteria.

```
createCohortTable(
  connectionDetails = NULL,
  connection = NULL,
  cohortDatabaseSchema,
  cohortTable = "cohort",
  createInclusionStatsTables = FALSE,
  resultsDatabaseSchema = cohortDatabaseSchema,
  cohortInclusionTable = paste0(cohortTable, "_inclusion"),
  cohortInclusionResultTable = paste0(cohortTable, "_inclusion_result"),
  cohortInclusionStatsTable = paste0(cohortTable, "_inclusion_stats"),
  cohortSummaryStatsTable = paste0(cohortTable, "_summary_stats")
)
```

connectionDetails

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package. Can be left NULL if connection is provided.

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.

cohortDatabaseSchema

Schema name where your cohort table resides. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'.

cohortTable Name of the cohort table.

createInclusionStatsTables

Create the four additional tables for storing inclusion rule statistics?

resultsDatabaseSchema

Schema name where the statistics tables reside. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'.

cohortInclusionTable

Name of the inclusion table, one of the tables for storing inclusion rule statistics.

cohortInclusionResultTable

Name of the inclusion result table, one of the tables for storing inclusion rule statistics.

cohortInclusionStatsTable

Name of the inclusion stats table, one of the tables for storing inclusion rule statistics.

 ${\tt cohortSummaryStatsTable}$ 

Name of the summary stats table, one of the tables for storing inclusion rule statistics.

createConceptCountsTable

Create concept counts table

### **Description**

Create a table with counts of how often each concept ID occurs in the CDM.

```
createConceptCountsTable(
  connectionDetails = NULL,
  connection = NULL,
  cdmDatabaseSchema,
  conceptCountsDatabaseSchema = cdmDatabaseSchema,
  conceptCountsTable = "concept_counts",
  conceptCountsTableIsTemp = FALSE
)
```

### connectionDetails

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package. Can be left NULL if connection is provided.

### 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.

### 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'.

### ${\tt conceptCountsDatabaseSchema}$

Schema name where your concept counts table resides. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'. Ignored if conceptCountsTableIsTemp = TRUE.

### conceptCountsTable

Name of the concept counts table. This table can be created using the createConceptCountsTable.

### conceptCountsTableIsTemp

Is the concept counts table a temp table?

findCohortIncludedSourceConcepts

Check source codes used in a cohort definition

# **Description**

This function first extracts all concept sets used in a cohort definition. Then, for each concept set the concept found in the CDM database the contributing source codes are identified.

```
findCohortIncludedSourceConcepts(
  connectionDetails = NULL,
  connection = NULL,
  cdmDatabaseSchema,
  oracleTempSchema = NULL,
  baseUrl = NULL,
  webApiCohortId = NULL,
  cohortJson = NULL,
  cohortSql = NULL,
  byMonth = FALSE,
  useSourceValues = FALSE
)
```

connectionDetails

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package. Can be left NULL if connection is provided.

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.

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'.

oracleTempSchema

Should be used in Oracle to specify a schema where the user has write privileges for storing temporary tables.

baseUrl The base URL for the WebApi instance, for example: "http://server.org:80/WebAPI".

Needn't be provided if cohortJson and cohortSql are provided.

webApiCohortId The ID of the cohort in the WebAPI instance. Needn't be provided if cohortJson

and cohortSql are provided.

cohortJson A character string containing the JSON of a cohort definition. Needn't be pro-

vided if baseUrl and cohortId are provided.

cohortSql The OHDSI SQL representation of the same cohort definition. Needn't be pro-

vided if baseUrl and cohortId are provided.

byMonth Compute counts by month? If FALSE, only overall counts are computed.

useSourceValues

Use the source\_value fields to find the codes used in the data? If not, this analysis will rely entirely on the source\_concept\_id fields instead. Note that, depending on the source data and ETL, it might be possible for the source\_value fields to contain patient-identifiable information by accident.

### Value

A data frame with source codes, with counts per domain how often the code was encountered in the CDM.

findCohortOrphanConcepts

Find orphan concepts for all concept sets in a cohort

# Description

Searches for concepts that should belong to the concept sets in a cohort definition but don't, for example because of missing source-to-standard concept maps, or erroneous hierarchical relationships.

### **Usage**

```
findCohortOrphanConcepts(
  connectionDetails = NULL,
  connection = NULL,
  cdmDatabaseSchema,
  oracleTempSchema = NULL,
  baseUrl = NULL,
  webApiCohortId = NULL,
  cohortJson = NULL,
  conceptCountsDatabaseSchema = cdmDatabaseSchema,
  conceptCountsTable = "concept_counts",
  conceptCountsTableIsTemp = FALSE
)
```

### **Arguments**

connectionDetails

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package. Can be left NULL if connection is provided.

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.

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'.

oracleTempSchema

Should be used in Oracle to specify a schema where the user has write privileges for storing temporary tables.

baseUrl

The base URL for the WebApi instance, for example: "http://server.org:80/WebAPI". Needn't be provided if cohortJson is provided.

webApiCohortId The ID of the cohort in the WebAPI instance. Needn't be provided if cohortJson is provided.

cohortJson

A character string containing the JSON of a cohort definition. Needn't be provided if baseUrl and webApiCohortId are provided.

 ${\tt conceptCountsDatabaseSchema}$ 

Schema name where your concept counts table resides. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'. Ignored if conceptCountsTableIsTemp = TRUE.

conceptCountsTable

Name of the concept counts table. This table can be created using the createConceptCountsTable. conceptCountsTableIsTemp

Is the concept counts table a temp table?

### **Details**

Logically, this function performs the following steps for each concept set expression in the cohort definition:

10 findOrphanConcepts

- Given the concept set expression, find all included concepts.
- Find all names of the input concepts, including synonyms, and the names of source concepts that map to them.
- Search for concepts (standard and source) that contain any of those names as substring.
- Filter those concepts to those that are not in the original set of concepts (i.e. orphans).
- Restrict the set of orphan concepts to those that appear in the CDM database and across network concept prevalence (as either source concept or standard concept).

### Value

A data frame with orphan concepts, with counts how often the code was encountered in the CDM.

findOrphanConcepts

Find (source) concepts that do not roll up to their ancestor(s)

# Description

Searches for concepts that should belong to the set of concepts but don't, for example because of missing source-to-standard concept maps, or erroneous hierarchical relationships.

### Usage

```
findOrphanConcepts(
  connectionDetails = NULL,
  connection = NULL,
  cdmDatabaseSchema,
  oracleTempSchema = NULL,
  conceptIds,
  conceptCountsDatabaseSchema = cdmDatabaseSchema,
  conceptCountsTable = "concept_counts",
  conceptCountsTableIsTemp = FALSE
)
```

# Arguments

connection Details

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package. Can be left NULL if connection is provided.

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.

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'.

oracleTempSchema

Should be used in Oracle to specify a schema where the user has write privileges for storing temporary tables.

```
conceptIds A vector of concept IDs for which we want to find orphans. conceptCountsDatabaseSchema
```

Schema name where your concept counts table resides. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'. Ignored if conceptCountsTableIsTemp = TRUE.

conceptCountsTable

 $Name\ of\ the\ concept\ counts\ table.\ This\ table\ can\ be\ created\ using\ the\ create\ Concept\ Counts\ Table.$   $concept\ Counts\ Table\ Is\ Temp$ 

Is the concept counts table a temp table?

### **Details**

Logically, this function performs the following steps for the input set of concept IDs:

- Find all names of the input concepts, including synonyms, and the names of source concepts that map to them.
- Search for concepts (standard and source) that contain any of those names as substring.
- Filter those concepts to those that are not in the original set of concepts (i.e. orphans).
- Restrict the set of orphan concepts to those that appear in the CDM database and across network concept prevalence (as either source concept or standard concept).

### Value

A data frame with orphan concepts, with counts how often the code was encountered in the CDM.

```
getCohortCharacteristics
```

Create characterization of a cohort

# **Description**

Computes features using all drugs, conditions, procedures, etc. observed on or prior to the cohort index date.

```
getCohortCharacteristics(
  connectionDetails = NULL,
  connection = NULL,
  cdmDatabaseSchema,
  oracleTempSchema = NULL,
  cohortDatabaseSchema = cdmDatabaseSchema,
  cohortTable = "cohort",
  cohortId,
  covariateSettings = FeatureExtraction::createDefaultCovariateSettings()
)
```

12 getCohortCounts

### **Arguments**

connectionDetails

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package. Can be left NULL if connection is provided.

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.

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'.

oracleTempSchema

Should be used in Oracle to specify a schema where the user has write privileges for storing temporary tables.

cohortDatabaseSchema

Schema name where your cohort table resides. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'.

cohortTable Name of the cohort table.

cohortId The cohort definition ID used to reference the cohort in the cohort table. covariateSettings

Either an object of type covariateSettings as created using one of the create-Covariate functions in the FeatureExtraction package, or a list of such objects.

### Value

A data frame with cohort characteristics.

# Description

Computes the subject and entry count per cohort

```
getCohortCounts(
  connectionDetails = NULL,
  connection = NULL,
  cohortDatabaseSchema,
  cohortTable = "cohort",
  cohortIds = c()
)
```

getIncidenceRate 13

### **Arguments**

connectionDetails

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package. Can be left NULL if connection

is provided.

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.

cohortDatabaseSchema

Schema name where your cohort table resides. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'.

cohortTable Name of the cohort table.

cohortIds The cohort definition ID(s0 used to reference the cohort in the cohort table. If

left empty, all cohorts in the table will be included.

### Value

A data frame with cohort counts

getIncidenceRate

Compute incidence rate for a cohort

### **Description**

Returns yearly incidence rate stratified by age and gender

### Usage

```
getIncidenceRate(
  connectionDetails = NULL,
  connection = NULL,
  cohortDatabaseSchema,
  cohortTable,
  cdmDatabaseSchema,
  oracleTempSchema = oracleTempSchema,
  firstOccurrenceOnly = TRUE,
  washoutPeriod = 365,
  cohortId
)
```

### **Arguments**

connectionDetails

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package. Can be left NULL if connection is provided.

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.

14 getInclusionStatistics

cohortDatabaseSchema

Schema name where your cohort table resides. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'.

cohortTable Name of the cohort table.

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'.

oracleTempSchema

Should be used in Oracle to specify a schema where the user has write privileges for storing temporary tables.

firstOccurrenceOnly

Use only the first occurrence of the cohort per person?

washoutPeriod The minimum amount of observation time required before the occurrence of a

cohort entry. This is also used to eliminate immortal time from the denominator.

cohortId The cohort definition ID used to reference the cohort in the cohort table.

### Value

Returns a data frame of cohort count, person year count, and incidence rate per 1000 persons years with the following stratifications: 1) no stratification, 2) gender stratification, 3) age (10-year) stratification, 4) calendar year and age (10-year) stratification, 5) calendar year and gender stratification, 6) calendar year, age (10-year), and gender stratification with option to save dataframes.

 ${\tt getInclusionStatistics}$ 

Get statistics on cohort inclusion criteria

# **Description**

Get statistics on cohort inclusion criteria

```
getInclusionStatistics(
  connectionDetails = NULL,
  connection = NULL,
  resultsDatabaseSchema,
  cohortId,
  simplify = TRUE,
  cohortTable = "cohort",
  cohortInclusionTable = paste0(cohortTable, "_inclusion"),
  cohortInclusionResultTable = paste0(cohortTable, "_inclusion_result"),
  cohortInclusionStatsTable = paste0(cohortTable, "_inclusion_stats"),
  cohortSummaryStatsTable = paste0(cohortTable, "_summary_stats")
)
```

connectionDetails

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package. Can be left NULL if connection

is provided.

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.

resultsDatabaseSchema

Schema name where the statistics tables reside. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'.

cohortId The cohort definition ID used to reference the cohort in the cohort table.

simplify Simply output the attrition table?

cohortTable Name of the cohort table. Used only to conveniently derive names of the four

rule statistics tables.

cohortInclusionTable

Name of the inclusion table, one of the tables for storing inclusion rule statistics.

cohortInclusionResultTable

Name of the inclusion result table, one of the tables for storing inclusion rule statistics.

cohortInclusionStatsTable

Name of the inclusion stats table, one of the tables for storing inclusion rule statistics.

 ${\tt cohortSummaryStatsTable}$ 

Name of the summary stats table, one of the tables for storing inclusion rule statistics.

### Value

If 'simplify = TRUE', this function returns a single data frame. Else a list of data frames is returned.

```
getInclusionStatisticsFromFiles
```

Get inclusion criteria statistics from files

### **Description**

Gets inclusion criteria statistics from files, as stored when using the ROhdsiWebApi::insertCohortDefinitionSetInPa function with generateStats = TRUE.

```
getInclusionStatisticsFromFiles(
  cohortId,
  folder,
  cohortInclusionFile = file.path(folder, "cohortInclusion.csv"),
  cohortInclusionResultFile = file.path(folder, "cohortIncResult.csv"),
  cohortInclusionStatsFile = file.path(folder, "cohortIncStats.csv"),
  cohortSummaryStatsFile = file.path(folder, "cohortSummaryStats.csv"),
  simplify = TRUE
)
```

16 getTimeDistributions

### **Arguments**

cohortId The cohort definition ID used to reference the cohort in the cohort table.

The path to the folder where the inclusion statistics are stored.

cohortInclusionFile

Name of the inclusion table, one of the tables for storing inclusion rule statistics.

cohortInclusionResultFile

Name of the inclusion result table, one of the tables for storing inclusion rule

statistics.

cohortInclusionStatsFile

Name of the inclusion stats table, one of the tables for storing inclusion rule

statistics.

cohortSummaryStatsFile

Name of the summary stats table, one of the tables for storing inclusion rule

statistics.

simplify Simply output the attrition table?

### Value

If 'simplify = TRUE', this function returns a single data frame. Else a list of data frames is returned.

getTimeDistributions Get time distributions of a cohort

# Description

Computes the distribution of the observation time before and after index, and time within a cohort.

# Usage

```
getTimeDistributions(
  connectionDetails = NULL,
  connection = NULL,
  cdmDatabaseSchema,
  oracleTempSchema = NULL,
  cohortDatabaseSchema = cdmDatabaseSchema,
  cohortTable = "cohort",
  cohortId
)
```

# **Arguments**

connectionDetails

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package. Can be left NULL if connection is provided.

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.

instantiateCohort 17

### 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'.

### oracleTempSchema

Should be used in Oracle to specify a schema where the user has write privileges for storing temporary tables.

### cohortDatabaseSchema

Schema name where your cohort table resides. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'.

cohortTable Name of the cohort table.

cohortId The cohort definition ID used to reference the cohort in the cohort table.

### Value

A data frame with time distributions

instantiateCohort

Instantiate a cohort

# **Description**

This function instantiates the cohort in the cohort table. Optionally, the inclusion rule statistics are computed and stored in the inclusion rule statistics tables described in createCohortTable).

```
instantiateCohort(
  connectionDetails = NULL,
  connection = NULL,
  cdmDatabaseSchema,
  oracleTempSchema = NULL,
  cohortDatabaseSchema = cdmDatabaseSchema,
  cohortTable = "cohort",
  baseUrl = NULL,
  webApiCohortId = NULL,
  cohortJson = NULL,
  cohortSql = NULL,
  cohortId = webApiCohortId,
  generateInclusionStats = FALSE,
  resultsDatabaseSchema = cohortDatabaseSchema,
  cohortInclusionTable = paste0(cohortTable, "_inclusion"),
  cohortInclusionResultTable = paste0(cohortTable, "_inclusion_result"),
cohortInclusionStatsTable = paste0(cohortTable, "_inclusion_stats"),
  cohortSummaryStatsTable = paste0(cohortTable, "_summary_stats")
```

18 instantiateCohort

### **Arguments**

connectionDetails

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package. Can be left NULL if connection is provided.

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.

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'.

oracleTempSchema

Should be used in Oracle to specify a schema where the user has write privileges for storing temporary tables.

cohortDatabaseSchema

Schema name where your cohort table resides. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'.

cohortTable Name of the cohort table.

baseUrl The base URL for the WebApi instance, for example: "http://server.org:80/WebAPI".

Needn't be provided if cohortJson and cohortSql are provided.

webApiCohortId The ID of the cohort in the WebAPI instance. Needn't be provided if cohortJson and cohortSql are provided.

cohortJson A character string containing the JSON of a cohort definition. Needn't be provided if baseUrl and cohortId are provided.

cohortSql The OHDSI SQL representation of the same cohort definition. Needn't be provided if baseUrl and cohortId are provided.

cohortId The cohort definition ID used to reference the cohort in the cohort table.

generateInclusionStats

Compute and store inclusion rule statistics?

resultsDatabaseSchema

Schema name where the statistics tables reside. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'.

cohortInclusionTable

Name of the inclusion table, one of the tables for storing inclusion rule statistics.

 ${\tt cohortInclusionResultTable}$ 

Name of the inclusion result table, one of the tables for storing inclusion rule statistics.

 ${\tt cohortInclusionStatsTable}$ 

Name of the inclusion stats table, one of the tables for storing inclusion rule statistics.

cohortSummaryStatsTable

Name of the summary stats table, one of the tables for storing inclusion rule statistics.

instantiateCohortSet 19

instantiateCohortSet Instantiate a set of cohort

# **Description**

This function instantiates a set of cohort in the cohort table, using definitions that are fetched from a WebApi interface. Optionally, the inclusion rule statistics are computed and stored in the inclusionStatisticsFolder.

### Usage

```
instantiateCohortSet(
  connectionDetails = NULL,
  connection = NULL,
  cdmDatabaseSchema,
  oracleTempSchema = NULL,
  cohortDatabaseSchema = cdmDatabaseSchema,
  cohortTable = "cohort",
  baseUrl,
  cohortSetReference,
  generateInclusionStats = FALSE,
  inclusionStatisticsFolder = NULL
)
```

# **Arguments**

connectionDetails

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package. Can be left NULL if connection is provided.

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.

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'.

oracleTempSchema

Should be used in Oracle to specify a schema where the user has write privileges for storing temporary tables.

cohortDatabaseSchema

Schema name where your cohort table resides. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'.

cohortTable Name of the cohort table.

baseUrl The base URL for the WebApi instance, for example: "http://server.org:80/WebAPI". cohortSetReference

A data frame with four columns, as described in the details.

generateInclusionStats

Compute and store inclusion rule statistics?

inclusionStatisticsFolder

The folder where the inclusion rule statistics are stored. Can be left NULL if generateInclusionStats = FALSE.

### **Details**

The cohortSetReference argument must be a data frame with the following columns:

atlasId The cohort ID in ATLAS.

atlasName The full name of the cohort. This will be shown in the Shiny app.

**cohortId** The cohort ID to use in the package. USually the same as the cohort ID in ATLAS.

**name** A short name for the cohort, to use to create file names. do not use special characters.

launchCohortExplorer Launch the CohortExplorer Shiny app

### **Description**

Launch the CohortExplorer Shiny app

### Usage

```
launchCohortExplorer(
  connectionDetails,
  cdmDatabaseSchema,
  cohortDatabaseSchema,
  cohortTable,
  cohortId,
  sampleSize = 100,
  subjectIds = NULL
)
```

### **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'.

cohortDatabaseSchema

Schema name where your cohort table resides. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'.

cohortTable Name of the cohort table. cohortId The ID of the cohort.

sampleSize Number of subjects to sample from the cohort. Ignored if subjectIds is specified.

subjectIds A vector of subject IDs to view.

### Details

Launches a Shiny app that allows the user to explore a cohort of interest.

 $launch {\tt DiagnosticsExplorer}$ 

Launch the Diagnostics Explorer Shiny app

# **Description**

Launch the Diagnostics Explorer Shiny app

# Usage

launchDiagnosticsExplorer(dataFolder, launch.browser = FALSE)

# **Arguments**

dataFolder A folder where the exported zip files for the diagnostics are stored. Use the

runCohortDiagnostics function to generate these zip files. Zip files containing

results from multiple databases can be placed in the same folder.

launch.browser Should the app be launched in your default browser, or in a Shiny window. Note:

copying to clipboard will not work in a Shiny window.

### **Details**

Launches a Shiny app that allows the user to explore the diagnostics

plotincidenceRate

Plot incidence rate by year, age, and/or gender

# Description

Characterizes the incidence rate of a cohort definition.

```
plotincidenceRate(
  incidenceRate,
  minPersonYears = 1000,
  stratifyByAge = TRUE,
  stratifyByGender = TRUE,
  stratifyByCalendarYear = TRUE,
  fileName = NULL
)
```

incidenceRate Incidence rate time series data for plotting generated using getIncidenceRate

function.

minPersonYears Estimates get very unstable with low background counts, so removing them

makes for cleaner plots.

stratifyByAge Should the plot be stratified by age?

stratifyByGender

Should the plot be stratified by gender?

stratifyByCalendarYear

Should the plot be stratified by calendar year?

fileName Optional: name of the file where the plot should be saved, for example 'plot.png'.

See the function ggsave in the ggplot2 package for supported file formats.

### **Details**

Generates time series plots of the incidence rate per 1000 person years of cohort entry by year, age, and/or gender.

### Value

A ggplot object. Use the ggsave function to save to file in a different format.

preMergeDiagnosticsFiles

Premerge Shiny diagnostics files

# **Description**

If there are many diagnostics files, starting the Shiny app may take a very long time. This function already does most of the preprocessing, increasing loadding speed.

The merged data will be stored in the same folder, and will automatically be recognized by the Shiny app.

### Usage

preMergeDiagnosticsFiles(dataFolder)

# **Arguments**

dataFolder

folder where the exported zip files for the diagnostics are stored. Use the runCohortDiagnostics function to generate these zip files. Zip files containing results from multiple

databases can be placed in the same folder.

runCohortDiagnostics 23

runCohortDiagnostics Run cohort diagnostics

# Description

Runs the cohort diagnostics on all (or a subset of) the cohorts instantiated using the ROhdsiWebApi::insertCohortDefin function. Assumes the cohorts have already been instantiated.

# Usage

```
runCohortDiagnostics(
  packageName = NULL,
  cohortToCreateFile = "settings/CohortsToCreate.csv",
 baseUrl = NULL,
  cohortSetReference = NULL,
  connectionDetails = NULL,
  connection = NULL,
  cdmDatabaseSchema,
 oracleTempSchema = NULL,
  cohortDatabaseSchema,
  cohortTable = "cohort",
  cohortIds = NULL,
  inclusionStatisticsFolder = NULL,
  exportFolder,
  databaseId,
 databaseName = databaseId,
  databaseDescription = "",
  runInclusionStatistics = TRUE,
  runIncludedSourceConcepts = TRUE,
  runOrphanConcepts = TRUE,
  runTimeDistributions = TRUE,
  runBreakdownIndexEvents = TRUE,
  runIncidenceRate = TRUE,
  runCohortOverlap = TRUE,
  runCohortCharacterization = TRUE,
 minCellCount = 5
)
```

### **Arguments**

packageName The name of the package containing the cohort definitions. Can be left NULL if

baseUrl and cohortSetReference have been specified.

cohortToCreateFile

The location of the cohortToCreate file within the package. Is ignored if baseUrl

and cohortSetReference have been specified.

baseUrl The base URL for the WebApi instance, for example: "http://server.org:80/WebAPI".

Can be left NULL if packageName and cohortToCreateFile have been speci-

fied.

cohortSetReference

A data frame with four columns, as described in the details. Can be left NULL if packageName and cohortToCreateFile have been specified.

24 runCohortDiagnostics

connectionDetails

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package. Can be left NULL if connection is provided.

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.

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'.

oracleTempSchema

Should be used in Oracle to specify a schema where the user has write privileges for storing temporary tables.

cohortDatabaseSchema

Schema name where your cohort table resides. Note that for SQL Server, this should include both the database and schema name, for example 'scratch.dbo'.

cohortTable Name of the cohort table.

cohortIds Optionally, provide a subset of cohort IDs to restrict the diagnostics to.

inclusionStatisticsFolder

The folder where the inclusion rule statistics are stored. Can be left NULL if runInclusionStatistics = FALSE.

exportFolder The folder where the output will be exported to. If this folder does not exist it

will be created.

databaseId A short string for identifying the database (e.g. 'Synpuf').

databaseName The full name of the database.

databaseDescription

A short description (several sentences) of the database.

runInclusionStatistics

Generate and export statistic on the cohort incusion rules?

 $\verb"runIncludedSourceConcepts"$ 

Generate and export the source concepts included in the cohorts?

runOrphanConcepts

Generate and export potential orphan concepts?

runTimeDistributions

Generate and export cohort time distributions?

runBreakdownIndexEvents

Generate and export the breakdown of index events?

runIncidenceRate

Generate and export the cohort incidence rates?

runCohortOverlap

Generate and export the cohort overlap?

runCohortCharacterization

Generate and export the cohort characterization?

minCellCount The minimum cell count for fields contains person counts or fractions.

#### **Details**

Currently two ways of executing this function are supported, either (1) embedded in a study package, assuming the cohort definitions are stored in that package using the ROhdsiWebApi::insertCohortDefinitionSet or (2) by using a WebApi interface to retrieve the cohort definitions.

When using this function from within a study package, use the packageName and cohortToCreateFile to specify the name of the study package, and the name of the cohortToCreate file within that package, respectivly

When using this function using a WebApi interface, use the baseUrl and cohortSetReference to specify how to connect to the WebApi, and which cohorts to fetch, respectively.

The cohortSetReference argument must be a data frame with the following columns:

```
atlasId The cohort ID in ATLAS.
```

atlasName The full name of the cohort. This will be shown in the Shiny app.

cohortId The cohort ID to use in the package. USually the same as the cohort ID in ATLAS.

name A short name for the cohort, to use to create file names. do not use special characters.

runCohortDiagnosticsUsingExternalCounts

Run cohort diagnostics using external concept counts

### **Description**

Runs cohort diagnostics on all (or a subset of) the cohorts, but using external concept counts. The external counts must have the following columns:

```
concept_id The source or target concept ID.concept_count The number of records having the concept.concept_subjects The number of unique persons having the concept.
```

```
runCohortDiagnosticsUsingExternalCounts(
 packageName = NULL,
 cohortToCreateFile = "settings/CohortsToCreate.csv",
 baseUrl = NULL,
 cohortSetReference = NULL,
 connectionDetails = NULL,
 connection = NULL,
 cdmDatabaseSchema,
 oracleTempSchema = NULL,
 cohortIds = NULL,
 conceptCountsDatabaseSchema = cdmDatabaseSchema,
 conceptCountsTable = "concept_counts",
 conceptCountsTableIsTemp = FALSE,
 exportFolder,
 databaseId,
 databaseName = databaseId,
 databaseDescription = "",
```

```
runIncludedSourceConcepts = TRUE,
runOrphanConcepts = TRUE,
minCellCount = 5
)
```

packageName

The name of the package containing the cohort definitions. Can be left NULL if

baseUrl and cohortSetReference have been specified.

cohortToCreateFile

The location of the cohortToCreate file within the package. Is ignored if baseUrl

and cohortSetReference have been specified.

baseUrl The base URL for the WebApi instance, for example: "http://server.org:80/WebAPI".

 $Can \ be \ left \ NULL \ if \ package Name \ and \ cohort To Create File \ have \ been \ speciment \ and \ cohort \ and \ cohort \ and \ and \ cohort \ and \ cohort \ and \ a$ 

fied.

cohortSetReference

A data frame with four columns, as described in the details. Can be left NULL

if packageName and cohortToCreateFile have been specified.

connectionDetails

An object of type connectionDetails as created using the createConnectionDetails

function in the DatabaseConnector package. Can be left NULL if connection

is provided.

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.

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'.

oracleTempSchema

Should be used in Oracle to specify a schema where the user has write privileges

for storing temporary tables.

cohortIds Optionally, provide a subset of cohort IDs to restrict the diagnostics to.

 ${\tt conceptCountsDatabaseSchema}$ 

Schema name where your concept counts table resides. Note that for SQL Server, this should include both the database and schema name, for example

'scratch.dbo'. Ignored if conceptCountsTableIsTemp = TRUE.

conceptCountsTable

Name of the concept counts table. This table can be created using the createConceptCountsTable.

conceptCountsTableIsTemp

Is the concept counts table a temp table?

exportFolder The folder where the output will be exported to. If this folder does not exist it

will be created.

databaseId A short string for identifying the database (e.g. 'Synpuf').

databaseName The full name of the database.

databaseDescription

A short description (several sentences) of the database.

runIncludedSourceConcepts

Generate and export the source concepts included in the cohorts?

runOrphanConcepts

Generate and export potential orphan concepts?

minCellCount The minimum cell count for fields contains person counts or fractions.

### **Details**

Currently two ways of executing this function are supported, either (1) embedded in a study package, assuming the cohort definitions are stored in that package using the ROhdsiWebApi::insertCohortDefinitionSet or (2) by using a WebApi interface to retrieve the cohort definitions.

When using this function from within a study package, use the packageName and cohortToCreateFile to specify the name of the study package, and the name of the cohortToCreate file within that package, respectivly

When using this function using a WebApi interface, use the baseUrl and cohortSetReference to specify how to connect to the WebApi, and which cohorts to fetch, respectively.

The cohortSetReference argument must be a data frame with the following columns:

atlasId The cohort ID in ATLAS.

atlasName The full name of the cohort. This will be shown in the Shiny app.

cohortId The cohort ID to use in the package. USually the same as the cohort ID in ATLAS.

**name** A short name for the cohort, to use to create file names. do not use special characters.

# **Index**

```
breakDownIndexEvents, 2
CohortDiagnostics, 3
compareCohortCharacteristics, 4
computeCohortOverlap, 4
connect, 3, 5-10, 12, 13, 15, 16, 18, 19, 24, 26
createCohortTable, 5, 17
createConceptCountsTable, 6, 7, 9, 11, 26
createConnectionDetails, 3, 5-10, 12, 13,
         15, 16, 18–20, 24, 26
findCohortIncludedSourceConcepts, 7
findCohortOrphanConcepts, 8
findOrphanConcepts, 10
getCohortCharacteristics, 4, 11
getCohortCounts, 12
getIncidenceRate, 13, 22
getInclusionStatistics, 14
getInclusionStatisticsFromFiles, 15
getTimeDistributions, 16
ggsave, 22
instantiateCohort, 17
\verb|instantiateCohortSet|, 19
launchCohortExplorer, 20
launchDiagnosticsExplorer, 21
plotincidenceRate, 21
preMergeDiagnosticsFiles, 22
runCohortDiagnostics, 21, 22, 23
run Cohort {\tt Diagnostics Using External Counts},
        25
```