Package 'CaseCrossover'

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Description An R package for performing case-crossover and case-time-control analyses in an observational database in the OMOP Common Data Model.
VignetteBuilder knitr
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R topics documented:
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CaseCrossover

CaseCrossover

Description

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createMatchingCriteria

Create matching criteria

Description

Create matching criteria

Usage

```
createMatchingCriteria(controlsPerCase = 1, matchOnAge = TRUE,
   ageCaliper = 2, matchOnGender = TRUE, matchOnProvider = FALSE,
   matchOnCareSite = FALSE, matchOnVisitDate = FALSE,
   visitDateCaliper = 30, matchOnTimeInCohort = FALSE,
   daysInCohortCaliper = 30)
```

Arguments

controlsPerCase

Maximum number of controls to select per case.

matchOnAge Match on age?

ageCaliper Maximum difference (in years) in age when matching on age.

matchOnGender Match on gender?

matchOnProvider

Match on provider (as specified in the person table)?

 ${\tt matchOnCareSite}$

Match on care site (as specified in the person table)?

 ${\tt matchOnVisitDate}$

Should the index date of the control be changed to the nearest visit date?

visitDateCaliper

Maximum difference (in days) between the index date and the visit date when matching on visit date.

matchOnTimeInCohort

Match on time in nesting cohort? When not using nesting, this is interpreted as time observed prior to index.

daysInCohortCaliper

Maximum difference (in days) in time in cohort.

fitCaseCrossoverModel 3

fitCaseCrossoverModel Fit case-crossover model

Description

Fit case-crossover model

Usage

fitCaseCrossoverModel(exposureStatus)

Arguments

 ${\tt exposureStatus} \ \ A \ data \ frame \ as \ generated \ using \ the \ {\tt getExposureStatus} \ function.$

Details

Fits a conditional logistic regression on the case-crossover data.

getAttritionTable

Get the attrition table for a set of subjects

Description

Get the attrition table for a set of subjects

Usage

getAttritionTable(subjects)

Arguments

subjects

A data frame of subjects as generated by the function selectSubjectsToInclude.

Value

A data frame specifying the number of cases and events after various steps of filtering.

```
getDbCaseCrossoverData
```

Load case-crossover data from the database

Description

Load all data about the cases from the database.

Usage

```
getDbCaseCrossoverData(connectionDetails, cdmDatabaseSchema,
  oracleTempSchema = cdmDatabaseSchema,
  outcomeDatabaseSchema = cdmDatabaseSchema, outcomeTable = "condition_era",
  outcomeIds = c(), useNestingCohort = FALSE,
  nestingCohortDatabaseSchema = cdmDatabaseSchema,
  nestingCohortTable = "cohort", nestingCohortId = NULL,
  useObservationEndAsNestingEndDate = TRUE, getVisits = FALSE,
  exposureDatabaseSchema = cdmDatabaseSchema, exposureTable = "drug_era",
  exposureIds = c(), studyStartDate = "", studyEndDate = "",
  getTimeControlData = FALSE)
```

Arguments

connectionDetails

An R object of type ConnectionDetails created using the function createConnectionDetails in the DatabaseConnector package.

cdmDatabaseSchema

The name of the database schema that contains the OMOP CDM instance. Requires read permissions to this database. On SQL Server, this should specify both the database and the schema, so for example 'cdm_instance.dbo'.

oracleTempSchema

A schema where temp tables can be created in Oracle.

outcomeDatabaseSchema

The name of the database schema that is the location where the data used to define the outcome cohorts is available. If outcomeTable = CONDITION_ERA, outcomeDatabaseSchema is not used. Requires read permissions to this database.

outcomeTable

The tablename that contains the outcome cohorts. If outcomeTable is not CON-DITION_OCCURRENCE or CONDITION_ERA, then expectation is outcomeTable has format of COHORT table: COHORT_DEFINITION_ID, SUBJECT_ID, COHORT_START_DATE, COHORT_END_DATE.

outcomeIds

A list of ids used to define outcomes. If outcomeTable = CONDITION_OCCURRENCE, the list is a set of ancestor CONCEPT_IDs, and all occurrences of all descendant concepts will be selected. If outcomeTable <> CONDITION_OCCURRENCE, the list contains records found in COHORT_DEFINITION_ID field.

useNestingCohort

Should the study be nested in a cohort (e.g. people with a specific indication)? If not, the study will be nested in the general population.

nestingCohortDatabaseSchema

The name of the database schema that is the location where the nesting cohort is defined.

getExposureStatus 5

 $nesting {\tt CohortTable}$

Name of the table holding the nesting cohort. This table should have the same structure as the cohort table.

nestingCohortId

A cohort definition ID identifying the records in the nestingCohortTable to use as nesting cohort.

useObservationEndAsNestingEndDate

When using a nesting cohort, should the observation period end date be used instead of the cohort end date?

getVisits Get data on visits? This is needed when performing a time- case-control study and matching on visit date is requested later on.

exposureDatabaseSchema

The name of the database schema that is the location where the exposure data used to define the exposure cohorts is available. If exposureTable = DRUG_ERA, exposureDatabaseSchema is not used but assumed to be cdmSchema. Requires read permissions to this database.

exposureTable The tablename that contains the exposure cohorts. If exposureTable <> DRUG_ERA,

then expectation is exposure Table has format of COHORT table: cohort_concept_id, SUBJECT_ID, COHORT_START_DATE, COHORT_END_DATE.

exposureIds are provided, all drugs or cohorts in the exposureTable are included

exposureIds A list of identifiers to define the exposures of interest. If exposureTable =

DRUG_ERA, exposureIds should be CONCEPT_ID. If exposureTable <> DRUG_ERA, exposureIds is used to select the cohort_concept_id in the cohort-like table. If no

as exposures.

studyStartDate A calendar date specifying the minimum date where data is used. Date format

is 'yyyymmdd'.

studyEndDate A calendar date specifying the maximum date where data is used. Date format

is 'yyyymmdd'.

getTimeControlData

Should data for time controls be fetched? (needed for case-time-control analyses.

Value

Returns an object of type caseCrossoverData, containing information on the cases, the nesting cohort, exposures, and optionally visits. Information about multiple outcomes can be captured at once for efficiency reasons. The generic summary() function has been implemented for this object.

getExposureStatus

Get the exposure status for cases (and controls).

Description

Get the exposure status for cases (and controls).

Usage

```
getExposureStatus(subjects, caseCrossoverData, exposureId,
  firstExposureOnly = FALSE, riskWindowStart = -30, riskWindowEnd = 0,
  controlWindowOffsets = c(-60))
```

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Arguments

subjects A data frame as generated using the selectSubjectsToInclude function.

caseCrossoverData

 $An \,object \,of \,type \,case Crossover Data \,as \,generated \,using \,the \,get Db Case Crossover Data \,and \,an extended a contraction of the contract$

function.

exposureId The identifier of the exposure.

firstExposureOnly

Should only the first exposure per subject be included?

riskWindowStart

The start of the risk window (in days) relative to the index date. This number

should be non-positive.

riskWindowEnd The end of the risk window (in days) relative to the index date. This number

should be non-positive.

controlWindowOffsets

Offsets in days of the control windows relative to the case window.

Details

This function determines the exposure status for a give, exposure ID in various windows relative to the index date.

Value

A data frame with these columns:

personId The person ID

indexDate The index date

isCase Is the person a case or a control?

stratumId The ID linking cases and controls in a matched set

isCaseWindow Is this a case window (as opposed to a control window)?

exposed Was the person exposed during the window?

loadCaseCrossoverData Load the case data from a folder

Description

loadCaseCrossoverData loads an object of type caseCrossoverData from a folder in the file system.

Usage

loadCaseCrossoverData(folder, readOnly = TRUE)

Arguments

folder The name of the folder containing the data.

readOnly If true, the data is opened read only.

saveCaseCrossoverData 7

Details

The data will be written to a set of files in the folder specified by the user.

Value

An object of class caseCrossoverData.

saveCaseCrossoverData Save the case-crossover data to folder

Description

saveCaseCrossoverData saves an object of type caseCrossoverData to folder.

Usage

saveCaseCrossoverData(caseCrossoverData, folder)

Arguments

caseCrossoverData

An object of type caseCrossoverData as generated using getDbCaseCrossoverData.

folder

The name of the folder where the data will be written. The folder should not yet exist.

Details

The data will be written to a set of files in the specified folder.

selectSubjectsToInclude

Select subjects to include

Description

Select subjects to include

Usage

```
selectSubjectsToInclude(caseCrossoverData, outcomeId, firstOutcomeOnly = TRUE,
  washoutPeriod = 180, matchingCriteria = NULL, minAge = NULL,
  maxAge = NULL)
```

Arguments

caseCrossoverData

 $An object of type \ case {\tt CrossoverData} \ as \ generated \ using \ the \ {\tt getDbCaseCrossoverData} \ as \ generated \ using \ the \ {\tt getDbCaseCrossoverData} \ as \ {\tt generated} \ as \ {\tt ge$

function.

outcomeId The outcome ID of the cases for which we need to pick controls.

firstOutcomeOnly

Use the first outcome per person?

washoutPeriod Minimum required numbers of days of observation for inclusion as either case

or control.

matchingCriteria

If provided, a case-time-control analysis will be performed and controls will be

matched based on these criteria.

minAge Minimum age at which patient time will be included in the analysis. Note that

information prior to the min age is still used to determine exposure status after the minimum age (e.g. when a prescription was started just prior to reaching the minimum age). Also, outcomes occurring before the minimum age is reached will be considered as prior outcomes when using first outcomes only. Age should be specified in years, but non-integer values are allowed. If not

specified, no age restriction will be applied.

maxAge Maximum age at which patient time will be included in the analysis. Age should

be specified in years, but non-integer values are allowed. If not specified, no age

restriction will be applied.

Details

Subject to include in the study are selected for a specific outcome, optionally filtering using a washout period, restricting to first occurrences of the outcome only, and restricting on age. If matching criteria are provided controls will be selected for each case. These controls will be used to adjust for time trends in exposure, turning the analysis into a case-time-control analysis (Suissa, 1995).

Value

A data frame with these columns:

personId The person ID

indexDate The index date

isCase Is the person a case or a control?

stratumId The ID linking cases and controls in a matched set

observationPeriodStartDate The observation period start date

References

Suissa S (1995) The case-time-control design. Epidemiology; 6:248-253.

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