

Package ‘omock’

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Type Package

Title Creation of Mock Observational Medical Outcomes Partnership
Common Data Model

Version 0.2.0

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Description Creates mock data for testing and package development for the
Observational Medical Outcomes Partnership common data model. The
package offers functions crafted with pipeline-friendly
implementation, enabling users to effortlessly include only the
necessary tables for their testing needs.

License Apache License (>= 2)

URL <https://oxford-pharmacoepi.github.io/omock/>,
<https://github.com/oxford-pharmacoepi/omock>

Depends R (>= 2.10)

Imports cli, dplyr, lubridate, methods, omopgenerics (>= 0.0.3),
purrr, rlang, snakecase, stats

Suggests ggplot2, knitr, rmarkdown, testthat (>= 3.0.0)

VignetteBuilder knitr

Config/testthat/edition 3

Encoding UTF-8

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BugReports <https://github.com/oxford-pharmacoepi/omock/issues>

NeedsCompilation no

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| | |
|-------------------|---|
| mockCdmFromTables | <i>Creates a mock cdm_reference from a preexisting cdm_reference object (it can be empty) and a list of tables. It makes sure that all the observations are within observation and all individuals are present in person table.</i> |
|-------------------|---|

Description

Creates a mock cdm_reference from a preexisting cdm_reference object (it can be empty) and a list of tables. It makes sure that all the observations are within observation and all individuals are present in person table.

Usage

```
mockCdmFromTables(cdm = mockCdmReference(), tables = list(), seed = NULL)
```

Arguments

| | |
|--------|---|
| cdm | A cdm_reference object |
| tables | List of named tables. They can be either omop standard tables or cohort tables. |
| seed | An integer to fix the seed of random numbers. If NULL seed is not set. |

Value

a cdm object

Examples

```
library(omock)
library(dplyr)

cohort <- tibble(
  cohort_definition_id = c(1, 1, 2, 2, 1, 3, 3, 3, 1, 3),
  subject_id = c(1, 4, 2, 3, 5, 5, 4, 3, 3, 1),
  cohort_start_date = as.Date(c(
    "2020-04-01", "2021-06-01", "2022-05-22", "2010-01-01", "2019-08-01",
    "2019-04-07", "2021-01-01", "2008-02-02", "2009-09-09", "2021-01-01"
  )),
  cohort_end_date = cohort_start_date
)

cdm <- mockCdmFromTables(tables = list(cohort = cohort))

cdm
cdm$cohort
cdm$person
```

mockCdmReference *Function to create empty cdm reference for mock database*

Description

Function to create empty cdm reference for mock database

Usage

```
mockCdmReference(cdmName = "mock database", vocabularySet = "mock")
```

Arguments

cdmName name of the mock cdm object
vocabularySet name of the vocabulary set

Value

an empty cdm object

Examples

```
library(omock)

cdm <- mockCdmReference()

cdm
```

mockCohort

Function to generate synthetic Cohort

Description

Function to generate synthetic Cohort

Usage

```
mockCohort(
  cdm,
  name = "cohort",
  numberCohorts = 1,
  cohortName = paste0("cohort_", seq_len(numberCohorts)),
  recordPerson = 1,
  seed = 1
)
```

Arguments

| | |
|---------------|---|
| cdm | the CDM reference into which the synthetic cohort will be added |
| name | The name of the table within the CDM where the cohort data will be stored. Defaults to "cohort". |
| numberCohorts | The number of different cohorts to create within the table. Defaults to 1. |
| cohortName | The name(s) of the cohorts to be created. This can be a single name or a vector of names if multiple cohorts are being created. Defaults to generating names based on a sequence. |
| recordPerson | The expected number of records per person within each cohort. This can help simulate the frequency of observations for individuals in the cohort. |
| seed | A random seed to ensure reproducibility of the generated data. |

Value

A cdm reference with the mock tables

Examples

```
library(omock)
cdm <- mockCdmReference() |>
  mockPerson(nPerson = 100) |>
  mockObservationPeriod() |>
  mockCohort(
    name = "omock_example",
    numberCohorts = 2,
    cohortName = c("omock_cohort_1", "omock_cohort_2")
  )

cdm
```

| | |
|--------------|--|
| mockConcepts | <i>function to add mockconcept et to concept table</i> |
|--------------|--|

Description

function to add mockconcept et to concept table

Usage

```
mockConcepts(cdm, conceptSet, domain = "Condition", seed = 1)
```

Arguments

| | |
|------------|------------------------------|
| cdm | Name of the cdm object |
| conceptSet | conceptset as a vector |
| domain | the domain of the conceptSet |
| seed | random seed |

Value

A cdm reference with the updated concept table

Examples

```
library(omock)
library(dplyr)

cdm <- mockCdmReference() |> mockConcepts(conceptSet = c(100,200), domain = "Condition")

cdm$concept |> filter(domain_id == "Condition")
```

mockConditionOccurrence

Function to generate condition occurrence table

Description

Function to generate condition occurrence table

Usage

```
mockConditionOccurrence(cdm, recordPerson = 1, seed = 1)
```

Arguments

| | |
|--------------|---|
| cdm | the CDM reference into which the synthetic cohort will be added |
| recordPerson | The expected number of records per person within each cohort. This can help simulate the frequency of observations for individuals in the cohort. |
| seed | A random seed to ensure reproducibility of the generated data. |

Value

A cdm reference with the mock tables

Examples

```
library(omock)

cdm <- mockCdmReference() |> mockPerson() |> mockObservationPeriod() |>
mockConditionOccurrence()

cdm$condition_occurrence
```

mockDeath

Function to creat mock death table

Description

Function to creat mock death table

Usage

```
mockDeath(cdm, recordPerson = 1, seed = 1)
```

Arguments

| | |
|--------------|---|
| cdm | the CDM reference into which the synthetic cohort will be added |
| recordPerson | The expected number of records per person within each cohort. This can help simulate the frequency of observations for individuals in the cohort. |
| seed | A random seed to ensure reproducibility of the generated data. |

Value

A cdm reference with the mock tables

Examples

```
library(omock)

cdm <- mockCdmReference() |> mockPerson() |> mockObservationPeriod() |>
mockDeath()

cdm$death
```

mockDrugExposure *Function to generate drug exposure table*

Description

Function to generate drug exposure table

Usage

```
mockDrugExposure(cdm, recordPerson = 1, seed = 1)
```

Arguments

| | |
|--------------|---|
| cdm | the CDM reference into which the synthetic cohort will be added |
| recordPerson | The expected number of records per person within each cohort. This can help simulate the frequency of observations for individuals in the cohort. |
| seed | A random seed to ensure reproducibility of the generated data. |

Value

A cdm reference with the mock tables

Examples

```
library(omock)

cdm <- mockCdmReference() |> mockPerson() |> mockObservationPeriod() |>
mockDrugExposure()

cdm$drug_exposure
```

| | |
|-----------------|---|
| mockMeasurement | <i>Function to generate measurement table</i> |
|-----------------|---|

Description

Function to generate measurement table

Usage

```
mockMeasurement(cdm, recordPerson = 1, seed = 1)
```

Arguments

| | |
|--------------|---|
| cdm | the CDM reference into which the mock measurement occurrence table will be added |
| recordPerson | The expected number of records per person within each cohort. This can help simulate the frequency of observations for individuals in the cohort. |
| seed | A random seed to ensure reproducibility of the generated data. |

Value

A cdm reference with the mock tables

Examples

```
library(omock)

cdm <- mockCdmReference() |> mockPerson() |> mockObservationPeriod() |>
mockMeasurement()

cdm$measurement
```

| | |
|-----------------|---|
| mockObservation | <i>Function to generate observation table</i> |
|-----------------|---|

Description

Function to generate observation table

Usage

```
mockObservation(cdm, recordPerson = 1, seed = 1)
```

Arguments

| | |
|--------------|---|
| cdm | the CDM reference into which the mock observation table will be added |
| recordPerson | The expected number of records per person within each cohort. This can help simulate the frequency of observations for individuals in the cohort. |
| seed | A random seed to ensure reproducibility of the generated data. |

Value

A cdm reference with the mock tables

Examples

```
library(omock)

cdm <- mockCdmReference() |> mockPerson() |> mockObservationPeriod() |>
mockObservation()

cdm$observation
```

| | |
|-----------------------|------------------------------|
| mockObservationPeriod | <i>mockObservationPeriod</i> |
|-----------------------|------------------------------|

Description

mockObservationPeriod

Usage

```
mockObservationPeriod(cdm, seed = 1)
```

Arguments

| | |
|------|------------------------|
| cdm | Name of the cdm object |
| seed | random seed |

Value

a cdm object with mock observation table

Examples

```
library(omock)
cdm <- mockCdmReference() |>
  mockPerson(nPerson = 100) |>
  mockObservationPeriod()
```

| | |
|------------|-------------------|
| mockPerson | <i>mockPerson</i> |
|------------|-------------------|

Description

mockPerson

Usage

```
mockPerson(
  cdm = mockCdmReference(),
  nPerson = 10,
  birthRange = as.Date(c("1950-01-01", "2000-12-31")),
  proportionFemale = 0.5,
  seed = 1
)
```

Arguments

| | |
|------------------|---|
| cdm | Name of the cdm object |
| nPerson | number of mock person to create in person table |
| birthRange | birthday range of the person in person table |
| proportionFemale | proportion of person in the person table are female |
| seed | random seed |

Value

A cdm reference with the mock person table

Examples

```
library(omock)
cdm <- mockPerson(cdm = mockCdmReference(), nPerson = 10)

cdm$person
```

mockVocabularyTables *It creates a mock database with the vocabulary.*

Description

It creates a mock database with the vocabulary.

Usage

```
mockVocabularyTables(
  cdm = mockCdmReference(),
  vocabularySet = "mock",
  cdmSource = NULL,
  concept = NULL,
  vocabulary = NULL,
  conceptRelationship = NULL,
  conceptSynonym = NULL,
  conceptAncestor = NULL,
  drugStrength = NULL
)
```

Arguments

| | |
|---------------------|-----------------------------|
| cdm | name of the cdm object |
| vocabularySet | name of the vocabulary set |
| cdmSource | cdm source table. |
| concept | Concept table. |
| vocabulary | Vocabulary table |
| conceptRelationship | Concept_relationship table. |
| conceptSynonym | Concept_synonym table. |
| conceptAncestor | Concept_ancestor table. |
| drugStrength | Drug_strength table. |

Value

A cdm reference with the vocabulary mock tables

Examples

```
library(omock)
```

```
cdm <- mockCdmReference() |> mockVocabularyTables(vocabularySet = "mock")
```

```
names(cdm)
```

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