Package 'PhenotypeLibrary'

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Type Package

Title The OHDSI Phenotype library

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Description A repository to store the content of the OHDSI Phenotype library.

Imports readr,
    dplyr,
    rlang,
    checkmate

License Apache License

RoxygenNote 7.2.0

Roxygen list(markdown = TRUE)

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Language en-US

URL https://ohdsi.github.io/PhenotypeLibrary/, https:
    //github.com/OHDSI/PhenotypeLibrary

BugReports https://github.com/OHDSI/PhenotypeLibrary/issues

R topics documented:

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getPlCohortDefinitionSet

Get a cohort definition set

Description

Get a cohort definition set

Usage

```
getPlCohortDefinitionSet(cohortIds)
```

Arguments

cohortIds   IDs of cohorts to extraction from the library.

Value

A tibble with the cohort ID, name, sql, and JSON for the provided cohort IDs. Can be used by the `CohortGenerator` package.

Examples

```
cohorts <- listPhenotypes()
subsetIds <- cohorts$cohortId[1:3]
getPlCohortDefinitionSet(subsetIds)
```

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listPhenotypes   *List all phenotypes in the library*

Description

List all phenotypes in the library

Usage

```
listPhenotypes()
```

Value

A tibble with the cohort ID and name.

Examples

```
listPhenotypes()
```

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