Package 'PhenotypeLibrary'

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Title The OHDSI Phenotype library Version 3.0.0 Date 2022-07-06 Author Gowtham Rao [aut, cre] Maintainer Gowtham Rao <pre>rao@ohdsi.org> Description A repository to store the content of the OHDSI Phenotype library. Imports readr,</pre>	Type Package
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```
{\tt 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)</pre>
```

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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