# Package 'DataQualityDashboard'

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```
Type Package
Title Execute and View Data Quality Checks on OMOP CDM Database
Version 2.4.0
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Maintainer Katy Sadowski <sadowski@ohdsi.org>
Description An R package for assessing data quality in standardized OMOP Com-
     mon Data Model data sources.
License Apache License (>= 2)
Config/build/clean-inst-doc FALSE
VignetteBuilder knitr
Config/testthat/edition 3
URL https://github.com/OHDSI/DataQualityDashboard
\pmb{BugReports} \ \text{https://github.com/OHDSI/DataQualityDashboard/issues}
Depends R (>= 3.2.2),
     DatabaseConnector (>= 2.0.2)
Imports magrittr,
     ParallelLogger,
     dplyr,
     jsonlite,
     rJava,
     SqlRender (>= 1.10.1),
     plyr,
     stringr,
     rlang,
     tidyselect,
     readr
Suggests testthat,
     knitr,
     rmarkdown,
     markdown,
     shiny,
     ggplot2,
     Eunomia,
     R.utils
```

Remotes ohdsi/Eunomia

RoxygenNote 7.2.2 Encoding UTF-8

## **R** topics documented:

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convertJsonResultsFileCase

Convert JSON results file case

## Description

Convert a DQD JSON results file between camelcase and (all-caps) snakecase. Enables viewing of pre-v.2.1.0 results files in later DQD versions, and vice versa

#### Usage

```
convertJsonResultsFileCase(
   jsonFilePath,
   writeToFile,
   outputFolder = NA,
   outputFile = "",
   targetCase
)
```

## **Arguments**

jsonFilePath Path to the JSON results file to be converted
writeToFile Whether or not to write the converted results back to a file (must be either TRUE or FALSE)

outputFolder The folder to output the converted JSON results file to

outputFile (OPTIONAL) File to write converted results JSON object to. Default is name of input file with a "\_camel" or "\_snake" postfix

targetCase Case into which the results file parameters should be converted (must be either "camel" or "snake")

#### Value

DQD results object (a named list)

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executeDqChecks

Execute DQ checks

## **Description**

This function will connect to the database, generate the sql scripts, and run the data quality checks against the database.

#### Usage

```
executeDqChecks(
  connectionDetails,
  cdmDatabaseSchema.
  resultsDatabaseSchema,
  vocabDatabaseSchema = cdmDatabaseSchema,
  cdmSourceName,
 numThreads = 1,
  sqlOnly = FALSE,
  sqlOnlyUnionCount = 1,
  sqlOnlyIncrementalInsert = FALSE,
 outputFolder,
 outputFile = "",
 verboseMode = FALSE,
 writeToTable = TRUE,
 writeTableName = "dqdashboard_results",
 writeToCsv = FALSE,
  csvFile = "",
  checkLevels = c("TABLE", "FIELD", "CONCEPT"),
 checkNames = c(),
  cohortDefinitionId = c(),
  cohortDatabaseSchema = resultsDatabaseSchema,
  cohortTableName = "cohort",
  tablesToExclude = c("CONCEPT", "VOCABULARY", "CONCEPT_ANCESTOR",
  "CONCEPT_RELATIONSHIP", "CONCEPT_CLASS", "CONCEPT_SYNONYM", "RELATIONSHIP", "DOMAIN"),
  cdmVersion = "5.3",
  tableCheckThresholdLoc = "default",
  fieldCheckThresholdLoc = "default",
  conceptCheckThresholdLoc = "default"
)
```

## **Arguments**

connectionDetails

A connectionDetails object for connecting to the CDM database

cdmDatabaseSchema

The fully qualified database name of the CDM schema

resultsDatabaseSchema

The fully qualified database name of the results schema

vocabDatabaseSchema

The fully qualified database name of the vocabulary schema (default is to set it as the cdmDatabaseSchema)

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cdmSourceName The name of the CDM data source

numThreads The number of concurrent threads to use to execute the queries sqlonly Should the SQLs be executed (FALSE) or just returned (TRUE)?

sq10nlyUnionCount

(OPTIONAL) In sqlOnlyIncrementalInsert mode, how many SQL commands to union in each query to insert check results into results table (can speed processing when queries done in parallel). Default is 1.

sqlOnlyIncrementalInsert

(OPTIONAL) In sqlOnly mode, boolean to determine whether to generate SQL queries that insert check results and associated metadata into results table. Default is FALSE (for backwards compatibility to <= v2.2.0)

outputFolder The folder to output logs, SQL files, and JSON results file to

outputFile (OPTIONAL) File to write results JSON object

verboseMode Boolean to determine if the console will show all execution steps. Default is

**FALSE** 

writeToTable Boolean to indicate if the check results will be written to the dqdashboard\_results

table in the resultsDatabaseSchema. Default is TRUE

writeTableName The name of the results table. Defaults to 'dqdashboard\_results'. Used when

sqlOnly or writeToTable is True.

writeToCsv Boolean to indicate if the check results will be written to a csv file. Default is

**FALSE** 

csvFile (OPTIONAL) CSV file to write results

checkLevels Choose which DQ check levels to execute. Default is all 3 (TABLE, FIELD,

CONCEPT)

checkNames (OPTIONAL) Choose which check names to execute. Names can be found

in inst/csv/OMOP\_CDM\_v[cdmVersion]\_Check\_Descriptions.csv. Note that "cdmTable", "cdmField" and "measureValueCompleteness" are always executed.

cohortDefinitionId

The cohort definition id for the cohort you wish to run the DQD on. The package assumes a standard OHDSI cohort table with the fields cohort\_definition\_id and subject\_id.

cohortDatabaseSchema

The schema where the cohort table is located.

cohortTableName

The name of the cohort table. Defaults to 'cohort'.

tablesToExclude

(OPTIONAL) Choose which CDM tables to exclude from the execution.

cdmVersion The CDM version to target for the data source. Options are "5.2", "5.3", or

"5.4". By default, "5.3" is used.

tableCheckThresholdLoc

The location of the threshold file for evaluating the table checks. If not specified the default thresholds will be applied.

fieldCheckThresholdLoc

The location of the threshold file for evaluating the field checks. If not specified the default thresholds will be applied.

conceptCheckThresholdLoc

The location of the threshold file for evaluating the concept checks. If not specified the default thresholds will be applied.

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#### Value

If sqlOnly = FALSE, a list object of results

listDqChecks

List DQ checks

## **Description**

Details on all checks defined by the DataQualityDashboard Package.

## Usage

```
listDqChecks(
  cdmVersion = "5.3",
  tableCheckThresholdLoc = "default",
  fieldCheckThresholdLoc = "default",
  conceptCheckThresholdLoc = "default")
```

## Arguments

cdmVersion The CDM version to target for the data source. By default, 5.3 is used. tableCheckThresholdLoc

The location of the threshold file for evaluating the table checks. If not specified the default thresholds will be applied.

fieldCheckThresholdLoc

The location of the threshold file for evaluating the field checks. If not specified the default thresholds will be applied.

conceptCheckThresholdLoc

The location of the threshold file for evaluating the concept checks. If not specified the default thresholds will be applied.

reEvaluateThresholds Re

Re-evaluate Thresholds

#### **Description**

Re-evaluate an existing DQD result against an updated thresholds file.

## Usage

```
reEvaluateThresholds(
   jsonFilePath,
   outputFolder,
   outputFile,
   tableCheckThresholdLoc = "default",
   fieldCheckThresholdLoc = "default",
   conceptCheckThresholdLoc = "default",
   cdmVersion = "5.3"
)
```

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#### **Arguments**

jsonFilePath Path to the JSON results file generated using the execute function

outputFolder The folder to output new JSON result file to

outputFile File to write results JSON object to

tableCheckThresholdLoc

The location of the threshold file for evaluating the table checks. If not specified

the default thresholds will be applied.

fieldCheckThresholdLoc

The location of the threshold file for evaluating the field checks. If not specified

the default thresholds will be applied.

conceptCheckThresholdLoc

The location of the threshold file for evaluating the concept checks. If not spec-

ified the default thresholds will be applied.

cdmVersion The CDM version to target for the data source. By default, 5.3 is used.

viewDqDashboard View DQ Dashboard

## **Description**

View DQ Dashboard

## Usage

```
viewDqDashboard(jsonPath, launch.browser = NULL, display.mode = NULL, ...)
```

## **Arguments**

jsonPath The path to the JSON file produced by executeDqChecks

launch.browser Passed on to shiny::runApp
display.mode Passed on to shiny::runApp

... Extra parameters for shiny::runApp() like "port" or "host"

writeJsonResultsToCsv Write JSON Results to CSV file

## **Description**

Write JSON Results to CSV file

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#### Usage

```
writeJsonResultsToCsv(
    jsonPath,
    csvPath,
    columns = c("checkId", "failed", "passed", "isError", "notApplicable", "checkName",
        "checkDescription", "thresholdValue", "notesValue", "checkLevel", "category",
        "subcategory", "context", "checkLevel", "cdmTableName", "cdmFieldName", "conceptId",
        "unitConceptId", "numViolatedRows", "pctViolatedRows", "numDenominatorRows",
        "executionTime", "notApplicableReason", "error", "queryText"),
    delimiter = ","
)
```

#### **Arguments**

jsonPath Path to the JSON results file generated using the execute function

csvPath Path to the CSV output file

columns (OPTIONAL) List of desired columns

delimiter (OPTIONAL) CSV delimiter

writeJsonResultsToTable

Write JSON Results to SQL Table

## **Description**

Write JSON Results to SQL Table

## Usage

```
writeJsonResultsToTable(
  connectionDetails,
  resultsDatabaseSchema,
  jsonFilePath,
  writeTableName = "dqdashboard_results",
  cohortDefinitionId = c()
)
```

#### **Arguments**

connectionDetails

A connectionDetails object for connecting to the CDM database

resultsDatabaseSchema

The fully qualified database name of the results schema

jsonFilePath Path to the JSON results file generated using the execute function

 $\label{lem:writeTableName} \begin{tabular}{ll} Name of table in the database to write results to cohortDefinitionId \end{tabular}$ 

If writing results for a single cohort this is the ID that will be appended to the table name