Package 'DataQualityDashboard'

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Type Package

Title Execute and View Data Quality Checks on OMOP CDM Database

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Description Assesses data quality in Observational Medical Outcomes Partnership Common Data Model (OMOP CDM) databases. Executes data quality checks and provides an R 'shiny' application to view the results.

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

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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)

`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. By default, results will be written to a json file as well as a database table.

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(),
  checkSeverity = c("fatal", "convention", "characterization"),
  cohortDefinitionId = c(),
  cohortDatabaseSchema = resultsDatabaseSchema,
  cohortTableName = "cohort",
  tablesToExclude = c("CONCEPT", "VOCABULARY", "CONCEPT_ANCESTOR",
   "CONCEPT_RELATIONSHIP", "CONCEPT_CLASS", "CONCEPT_SYNONYM", "RELATIONSHIP", "DOMAIN",
    "PACK_CONTENT", "CONCEPT_METADATA", "CONCEPT_RELATIONSHIP_METADATA"),
  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)
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)?
sqlOnlyUnionCount	(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.
checkSeverity	Choose which DQ check severity levels to execute. Default is all 3 (fatal, convention, characterization)
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. By default the vocabulary tables are excluded, including the CONCEPT_METADATA, CONCEPT_RELATIONSHIP_METADATA, and PACK_CONTENT tables added in CDM v5.5.

`cdmVersion`
: The CDM version to target for the data source. Options are "5.2", "5.3", "5.4", or "5.5". 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.

Value

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.

Value

A list containing check descriptions, table checks, field checks, and concept checks

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`reEvaluateThresholds` *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"
)
```

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 specified the default thresholds will be applied.

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

Value

A list containing the re-evaluated DQD results

`viewDqDashboard` *View DQ Dashboard*

Description

View DQ Dashboard

Usage

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

Arguments

`jsonPath`	The fully-qualified 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"

Value

NULL (launches Shiny application)

`writeDBResultsToJson` *Write DQD results database table to json*

Description

Write DQD results database table to json

Usage

```
writeDBResultsToJson(
  connection,
  resultsDatabaseSchema,
  cdmDatabaseSchema,
  writeTableName,
  outputFolder,
  outputFile
)
```

Arguments

connection	A connection object
resultsDatabaseSchema	The fully qualified database name of the results schema
cdmDatabaseSchema	The fully qualified database name of the CDM schema
writeTableName	Name of DQD results table in the database to read from
outputFolder	The folder to output the json results file to
outputFile	The output filename of the json results file

Value

NULL (writes results to JSON file)

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writeJsonResultsToCsv *Write JSON Results to CSV file*

Description

Write JSON Results to CSV file

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

Value

NULL (writes results to CSV file)

`writeJsonResultsToTable` *Write JSON Results to SQL Table*

Description

Write JSON Results to SQL Table

Usage

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

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
`writeTableName`	Name of table in the database to write results to
`cohortDefinitionId`	If writing results for a single cohort this is the ID that will be appended to the table name
`singleTable`	If TRUE, writes all results to a single table. If FALSE (default), writes to 3 separate tables by check level (table, field, concept) (NOTE this default behavior will be deprecated in the future)

Value

NULL (writes results to database table)

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