# Package 'OhdsiShinyModules'

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```
Type Package
Title Repository of Shiny Modules for OHDSI Result Viewers
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Description Install this package to access useful shiny modules for building shiny apps to ex-
     plore results using the OHDSI tools .
License Apache License 2.0
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LazyData true
Depends R (>=3.3.0)
VignetteBuilder knitr
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Imports checkmate,
     CirceR,
     cowplot,
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     gridExtra,
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     lubridate,
     methods,
     ParallelLogger,
     plotly,
     purrr,
     reactable,
     readr,
     RJSONIO,
     rlang,
     rmarkdown,
     scales,
     shiny,
     shinycssloaders,
     shinydashboard,
```

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$\operatorname{SqlRender},$
m stringi,
m stringr,
tibble,
tidyr,
${ m tidyselect},$
$\mathrm{tippy},$
RColorBrewer,
$\operatorname{markdown}$
Suggests kableExtra,
knitr,
$\operatorname{ResultModelManager},$
$\operatorname{RSQLite},$
${ m testthat},$
$\operatorname{withr}$
Remotes ohdsi/CirceR, ohdsi/ResultModelManager
RoxygenNote 7.2.3

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sccsServer
sccsView
timeDistributionsView
visitContextView

 $about {\tt HelperFile}$ 

The location of the about module helper file

# Description

The location of the about module helper file

# ${\bf Usage}$

aboutHelperFile()

# Details

Returns the location of the about helper file

# Value

string location of the about helper file

aboutServer 5

aboutServer

The module server for the shiny app home

# Description

The module server for the shiny app home

### Usage

```
aboutServer(
  id = "homepage",
  connectionHandler = NULL,
  resultDatabaseSettings = NULL
)
```

# Arguments

 $\begin{tabular}{ll} id & the unique reference id for the module \\ connection Handler \\ \end{tabular}$ 

a connection to the database with the results

resultDatabaseSettings

a list containing the characterization result schema, dbms, table Prefix, database Table and  ${\it cgTablePrefix}$ 

#### **Details**

The user specifies the id for the module

# Value

The server for the shiny app home

aboutViewer

The module viewer for the shiny app home

### Description

The module viewer for the shiny app home

# Usage

```
aboutViewer(id = "homepage")
```

#### Arguments

id

the unique reference id for the module

#### **Details**

The user specifies the id for the module

#### Value

The user interface to the home page module

characterization Aggregate Features Server

The module server for exploring aggregate features results

# Description

The module server for exploring aggregate features results

### Usage

```
characterizationAggregateFeaturesServer(
  id,
  connectionHandler,
  resultDatabaseSettings
)
```

### Arguments

```
\begin{tabular}{ll} id & the unique reference id for the module \\ connectionHandler & the connection to the prediction result database \\ resultDatabaseSettings & a list containing the characterization result schema, dbms, tablePrefix, databaseTable and cgTablePrefix \\ \end{tabular}
```

# **Details**

The user specifies the id for the module

### Value

The server to the description aggregate features module

```
characterization Aggregate Features Viewer\\
```

The module viewer for exploring aggregate feature results

### Description

The module viewer for exploring aggregate feature results

#### Usage

characterizationAggregateFeaturesViewer(id)

# Arguments

id the unique reference id for the module

#### **Details**

The user specifies the id for the module

### Value

The user interface to the description aggregate feature module

```
characterization Dechallenge Rechallenge Server\\
```

The module server for exploring Dechallenge Rechallenge results

# Description

The module server for exploring Dechallenge Rechallenge results

# Usage

```
characterizationDechallengeRechallengeServer(
  id,
  connectionHandler,
  resultDatabaseSettings
)
```

# Arguments

```
\begin{tabular}{ll} id & the unique reference id for the module \\ connectionHandler & the connection to the prediction result database \\ resultDatabaseSettings & a list containing the characterization result schema, dbms, tablePrefix, databaseTable and cgTablePrefix \\ \end{tabular}
```

#### **Details**

The user specifies the id for the module

### Value

The server to the Dechallenge Rechallenge module

#### characterizationDechallengeRechallengeViewer

The module viewer for exploring Dechallenge Rechallenge results

# Description

The module viewer for exploring Dechallenge Rechallenge results

### Usage

characterizationDechallengeRechallengeViewer(id)

# Arguments

id

the unique reference id for the module

#### Details

The user specifies the id for the module

### Value

The user interface to the description Dechallenge Rechallenge module

# $characterization {\tt HelperFile}$

The location of the characterization module helper file

# Description

The location of the characterization module helper file

# Usage

characterizationHelperFile()

# Details

Returns the location of the characterization helper file

# Value

string location of the characterization helper file

#### characterization Incidence Server

The module server for exploring incidence results

### Description

The module server for exploring incidence results

# Usage

characterizationIncidenceServer(id, connectionHandler, resultDatabaseSettings)

### Arguments

id the unique reference id for the module

connectionHandler

the connection to the prediction result database

resultDatabaseSettings

a list containing the characterization result schema, dbms, table Prefix, database Table and  ${\it cgTablePrefix}$ 

#### **Details**

The user specifies the id for the module

#### Value

The server to the prediction incidence module

# characterizationIncidenceViewer

The module viewer for exploring incidence results

### Description

The module viewer for exploring incidence results

# Usage

characterizationIncidenceViewer(id)

#### Arguments

id the unique reference id for the module

#### **Details**

The user specifies the id for the module

#### Value

The user interface to the description incidence module

characterizationServer

 $The \ module \ server \ for \ exploring \ characterization \ studies$ 

# Description

The module server for exploring characterization studies

# Usage

```
characterizationServer(
  id,
  connectionHandler,
  resultDatabaseSettings = list(port = 1)
)
```

# Arguments

### **Details**

The user specifies the id for the module

#### Value

The server for the characterization module

 ${\tt characterization Table Server}$ 

The module server for exploring 1 or more cohorts features

# Description

The module server for exploring 1 or more cohorts features

# Usage

characterizationTableServer(id, connectionHandler, resultDatabaseSettings)

### Arguments

id the unique reference id for the module

connectionHandler

the connection to the prediction result database

resultDatabaseSettings

a list containing the characterization result schema, dbms, table Prefix, database Table and  ${\it cgTablePrefix}$ 

### **Details**

The user specifies the id for the module

### Value

The server to the cohorts features server

characterizationTableViewer

The module viewer for exploring 1 or more cohorts features

### Description

The module viewer for exploring 1 or more cohorts features

# Usage

characterizationTableViewer(id)

### Arguments

id the unique reference id for the module

### **Details**

The user specifies the id for the module

### Value

The user interface to the description cohorts features

#### $characterization {\tt TimeToEventServer}$

The module server for exploring time to event results

### Description

The module server for exploring time to event results

# Usage

```
characterizationTimeToEventServer(
  id,
  connectionHandler,
  resultDatabaseSettings
)
```

# Arguments

```
\begin{tabular}{ll} id & the unique reference id for the module \\ connectionHandler & the connection to the prediction result database \\ \end{tabular}
```

 $\label{eq:containing} a \ \ \mbox{list containing the characterization result schema, dbms, table Prefix,} \\ database Table \ \mbox{and} \ \mbox{cgTable Prefix}$ 

# Details

The user specifies the id for the module

### Value

The server to the prediction time to event module

characterizationTimeToEventViewer

The module viewer for exploring time to event results

# Description

The module viewer for exploring time to event results

### Usage

```
characterizationTimeToEventViewer(id)
```

#### Arguments

id the unique reference id for the module

characterizationViewer 13

#### **Details**

The user specifies the id for the module

#### Value

The user interface to the characterization time to event module

characterizationViewer

The module viewer for exploring characterization studies

# Description

The module viewer for exploring characterization studies

# Usage

```
characterizationViewer(id = 1)
```

# Arguments

id

the unique reference id for the module

### **Details**

The user specifies the id for the module

### Value

The user interface to the characterization viewer module

cohortCountsModule

Shiny module for cohort counts

# Description

Shiny module for cohort counts. Displays reactable table of cohort counts

### Usage

```
cohortCountsModule(
   id,
   dataSource,
   cohortTable = dataSource$cohortTable,
   databaseTable = dataSource$dbTable,
   selectedCohorts,
   selectedDatabaseIds,
   cohortIds
)
```

14 cohort Definitions Module

# Arguments

id namespace id

dataSource Backend Data source (DatabaseConnection)

cohortTable data.frame of all cohorts
databaseTable data.frame of all databases

selectedCohorts

shiny::reactive - should return cohorts selected or NULL

selectedDatabaseIds

shiny::reactive - should return cohorts selected or NULL

cohortIds shiny::reactive - should return cohorts selected integers or NULL

cohortCountsView

Cohort Counts View

# Description

Shiny view for cohort counts module

### Usage

```
cohortCountsView(id)
```

# Arguments

id

Namespace id

cohortDefinitionsModule

Cohort Definition module

# Description

cohort defintion conceptsets, json etc

### Usage

```
cohortDefinitionsModule(
   id,
   dataSource,
   cohortDefinitions,
   cohortTable = dataSource$cohortTable,
   cohortCountTable = dataSource$cohortCountTable,
   databaseTable = dataSource$dbTable
)
```

cohort Definitions View 15

# Arguments

id Namespace id

dataSource DatabaseConnection

cohortDefinitions

reactive of cohort definitions to display

cohortTable data.frame of cohorts, cohortId, cohortName

cohortCountTable

data.frame of cohortCounts, cohortId, subjects records

databaseTable data.frame of databasese, databaseId, name

 ${\tt cohortDefinitionsView}$   ${\tt Cohort\ Definitions\ View}$ 

# Description

Outputs cohort definitions

### Usage

cohortDefinitionsView(id)

### Arguments

id Namespace id for module

 ${\tt cohortDiagCharacterizationView}$ 

characterization

# Description

Use for customizing UI

# Usage

cohortDiagCharacterizationView(id)

# Arguments

id Namespace Id - use namespaced id ns("characterization") inside diagnos-

ticsExplorer module

 ${\tt cohortDiagnosticsHelperFile}$ 

The location of the description module helper file

# Description

The location of the description module helper file

# Usage

```
cohortDiagnosticsHelperFile()
```

#### **Details**

Returns the location of the description helper file

#### Value

string location of the description helper file

```
{\tt cohortDiagnosticsServer}
```

Cohort Diagnostics Explorer main module

# Description

Cohort Diagnostics Explorer main module

# Usage

```
cohortDiagnosticsServer(
   id,
   connectionHandler,
   resultDatabaseSettings,
   dataSource = NULL
)
```

# Arguments

id module Id

connectionHandler

 $Result Model Manager\ Connection Hander\ instance$ 

resultDatabaseSettings

results database settings

 ${\tt dataSource} \qquad {\tt dataSource} \quad {\tt d$ 

 ${\tt cohortDiagnosticsView} \quad \textit{View for cohort diagnostics module}$ 

# Description

View for cohort diagnostics module

### Usage

```
cohortDiagnosticsView(id = "DiagnosticsExplorer")
```

#### Arguments

id

the unique reference id for the module

### **Details**

The user specifies the id for the module

### Value

The user interface to the cohort diagnostics viewer module

```
{\tt cohortGeneratorHelperFile}
```

The location of the cohort-generator module helper file

# Description

The location of the cohort-generator module helper file

# Usage

```
cohortGeneratorHelperFile()
```

# Details

Returns the location of the cohort-generator helper file

### Value

string location of the cohort-generator helper file

18 cohortGeneratorViewer

cohortGeneratorServer The module server for the main cohort generator module

# Description

The module server for the main cohort generator module

### Usage

cohortGeneratorServer(id, connectionHandler, resultDatabaseSettings)

# Arguments

a named list containing the cohort generator results database details (schema, table prefix)

### Value

the cohort generator results viewer main module server

 ${\tt cohortGenerator Viewer} \quad \textit{The viewer of the main cohort generator module}$ 

# Description

The viewer of the main cohort generator module

### Usage

cohortGeneratorViewer(id)

### Arguments

id the unique reference id for the module

#### Value

The user interface to the cohort generator results viewer

 ${\tt cohortMethodAttritionServer}$ 

 $The\ module\ server\ for\ rendering\ the\ PLE\ attrition\ results$ 

# Description

The module server for rendering the PLE attrition results

#### Usage

```
cohortMethodAttritionServer(
  id,
  selectedRow,
  connectionHandler,
  resultDatabaseSettings
)
```

# Arguments

```
\begin{tabular}{lll} id & the unique reference id for the module \\ selectedRow & the selected row from the main results table \\ connectionHandler & the connection to the PLE results database \\ resultDatabaseSettings & a list containing the result schema and prefixes \\ \end{tabular}
```

#### Value

the PLE attrition results content server

 ${\tt cohortMethodAttritionViewer}$ 

The module viewer for rendering the PLE attrition results

# Description

The module viewer for rendering the PLE attrition results

#### Usage

```
cohortMethodAttritionViewer(id)
```

# Arguments

id the unique reference id for the module

# Value

The user interface to the cohort method attrition

 $cohort {\tt MethodCovariateBalanceServer}$ 

The module server for rendering the covariate balance plot

# Description

The module server for rendering the covariate balance plot

# Usage

```
cohortMethodCovariateBalanceServer(
  id,
  selectedRow,
  connectionHandler,
  resultDatabaseSettings,
  metaAnalysisDbIds = NULL
)
```

#### Arguments

```
\begin{tabular}{lll} id & the unique reference id for the module \\ selectedRow & the selected row from the main results table \\ connectionHandler & the connection to the PLE results database \\ resultDatabaseSettings & a list containing the result schema and prefixes \\ metaAnalysisDbIds & metaAnalysisDbIds \\ \hline \end{tabular}
```

### Value

the PLE covariate balance content server

cohortMethodCovariateBalanceViewer

 $\label{eq:covariate} \textit{The module viewer for rendering the PLE covariate balance analysis}$ 

# Description

The module viewer for rendering the PLE covariate balance analysis

#### Usage

```
cohortMethodCovariateBalanceViewer(id)
```

### Arguments

id the unique reference id for the module

#### Value

The user interface to the cohort method covariate balance results

 $cohort {\tt MethodDiagnosticsSummaryServer}$ 

The module server for rendering the PLE diagnostics summary

# Description

The module server for rendering the PLE diagnostics summary

### Usage

```
cohortMethodDiagnosticsSummaryServer(
  id,
  connectionHandler,
  resultDatabaseSettings,
  inputSelected
)
```

#### Arguments

id the unique reference id for the module

connectionHandler

the connection to the PLE results database

resultDatabaseSettings

a list containing the result schema and prefixes

inputSelected The target id, comparator id, outcome id and analysis id selected by the

user

#### Value

the PLE diagnostics summary results

cohortMethodDiagnosticsSummaryViewer

The module viewer for rendering the PLE diagnostics results

### Description

The module viewer for rendering the PLE diagnostics results

# Usage

cohortMethodDiagnosticsSummaryViewer(id)

#### Arguments

id

the unique reference id for the module

#### Value

The user interface to the cohort method diagnostics viewer

```
cohortMethodHelperFile
```

The location of the cohort method module helper file

# Description

The location of the cohort method module helper file

#### Usage

```
cohortMethodHelperFile()
```

#### **Details**

Returns the location of the cohort method helper file

#### Value

string location of the cohort method helper file

```
cohort Method Kaplan Meier Server
```

The module server for rendering the Kaplan Meier curve

### Description

The module server for rendering the Kaplan Meier curve

# ${\bf Usage}$

```
cohortMethodKaplanMeierServer(
  id,
  selectedRow,
  connectionHandler,
  resultDatabaseSettings
)
```

# Arguments

```
\begin{tabular}{lll} id & the unique reference id for the module \\ selectedRow & the selected row from the main results table \\ connectionHandler & the connection to the PLE results database \\ resultDatabaseSettings & a list containing the result schema and prefixes \\ \end{tabular}
```

#### Value

the PLE Kaplain Meier content server

cohortMethodKaplanMeierViewer

The module viewer for rendering the PLE Kaplan Meier curve

### Description

The module viewer for rendering the PLE Kaplan Meier curve

### Usage

```
cohortMethodKaplanMeierViewer(id)
```

# Arguments

id

the unique reference id for the module

#### Value

The module viewer for Kaplan Meier objects

 $cohort {\tt MethodPopulationCharacteristicsServer}$ 

The module server for rendering the population characteristics

### Description

The module server for rendering the population characteristics

# Usage

```
cohortMethodPopulationCharacteristicsServer(
  id,
  selectedRow,
  connectionHandler,
  resultDatabaseSettings
)
```

### Arguments

```
id the unique reference id for the module selectedRow the selected row from the main results table connectionHandler the connection to the PLE results database resultDatabaseSettings a list containing the result schema and prefixes
```

# Value

the PLE population characteristics content server

### $cohort {\tt MethodPopulationCharacteristicsViewer}$

The module viewer for rendering the PLE population characteristics

### Description

The module viewer for rendering the PLE population characteristics

#### Usage

cohortMethodPopulationCharacteristicsViewer(id)

### Arguments

id

the unique reference id for the module

#### Value

The user interface to the cohort method population characteristics objects

cohortMethodPowerServer

The module server for rendering the PLE power analysis results

### Description

The module server for rendering the PLE power analysis results

### Usage

```
cohortMethodPowerServer(
   id,
   selectedRow,
   connectionHandler,
   resultDatabaseSettings)
```

### Arguments

# Value

the PLE systematic error power server

cohortMethodPowerViewer

The module viewer for rendering the PLE power analysis

### Description

The module viewer for rendering the PLE power analysis

# Usage

```
cohortMethodPowerViewer(id)
```

# Arguments

id

the unique reference id for the module

#### Value

The user interface to the cohort method power calculation results

```
cohort Method Propensity Model Server
```

The module server for rendering the propensity score model

### Description

The module server for rendering the propensity score model

# Usage

```
cohortMethodPropensityModelServer(
  id,
  selectedRow,
  connectionHandler,
  resultDatabaseSettings
)
```

### Arguments

```
id the unique reference id for the module selectedRow the selected row from the main results table connectionHandler the connection to the PLE results database resultDatabaseSettings a list containing the result schema and prefixes
```

### Value

the PLE propensity score model

 ${\tt cohortMethodPropensityModelViewer}$ 

 $\label{lem:covariates} The \ module \ viewer \ for \ rendering \ the \ PLE \ propensity \ score \ model \\ covariates/coefficients$ 

#### Description

The module viewer for rendering the PLE propensity score model covariates/coefficients

#### Usage

cohortMethodPropensityModelViewer(id)

### Arguments

id

the unique reference id for the module

#### Value

The user interface to the cohort method propensity score model covariates/coefficients

 ${\tt cohortMethodPropensityScoreDistServer}$ 

The module server for rendering a PLE propensity score distribution

# Description

The module server for rendering a PLE propensity score distribution

### Usage

```
cohortMethodPropensityScoreDistServer(
  id,
  selectedRow,
  connectionHandler,
  resultDatabaseSettings,
  metaAnalysisDbIds = F
)
```

# Arguments

```
\begin{tabular}{lll} id & the unique reference id for the module \\ selectedRow & the selected row from the main results table \\ connectionHandler & the connection to the PLE results database \\ resultDatabaseSettings & a list containing the result schema and prefixes \\ metaAnalysisDbIds & metaAnalysisDbIds \\ \end{tabular}
```

#### Value

the PLE propensity score distribution content server

 $cohort {\tt MethodPropensityScoreDistViewer}$ 

The module viewer for rendering the propensity score distribution

### Description

The module viewer for rendering the propensity score distribution

### Usage

```
cohortMethodPropensityScoreDistViewer(id)
```

### Arguments

id

the unique reference id for the module

#### Value

The user interface to the cohort method propensity score distribution

 ${\tt cohortMethodResultSummaryServer}$ 

The module server for rendering the PLE diagnostics summary

# Description

The module server for rendering the PLE diagnostics summary

### Usage

```
cohortMethodResultSummaryServer(
  id,
  connectionHandler,
  resultDatabaseSettings,
  inputSelected
)
```

### Arguments

```
id the unique reference id for the module
```

connectionHandler

the connection to the PLE results database

resultDatabaseSettings

a list containing the result schema and prefixes

inputSelected The target id, comparator id, outcome id and analysis id selected by the

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#### Value

the PLE diagnostics summary results

 $cohort {\tt MethodResultSummaryViewer}$ 

The module viewer for rendering the cohort method results

# Description

The module viewer for rendering the cohort method results

#### Usage

cohortMethodResultSummaryViewer(id)

#### Arguments

id the unique reference id for the module

### Value

The user interface to the cohort method diagnostics viewer

 ${\tt cohortMethodServer}$ 

The module server for the main cohort method module

### Description

The module server for the main cohort method module

# Usage

cohortMethodServer(id, connectionHandler, resultDatabaseSettings)

# Arguments

 $\mbox{id} \qquad \qquad \mbox{the unique reference id for the module}$ 

connectionHandler

a connection to the database with the results

 ${\tt resultDatabaseSettings}$ 

a named list containing the PLE results database connection details

### Value

the PLE results viewer main module server

 ${\tt cohortMethodSystematicErrorServer}$ 

The module server for rendering the systematic error objects

# Description

The module server for rendering the systematic error objects

# Usage

```
cohortMethodSystematicErrorServer(
  id,
  selectedRow,
  connectionHandler,
  resultDatabaseSettings
)
```

#### Arguments

```
\begin{tabular}{lll} id & the unique reference id for the module \\ selectedRow & the selected row from the main results table \\ connectionHandler & the connection handler to the result databases \\ resultDatabaseSettings & a list containing the result schema and prefixes \\ \end{tabular}
```

#### Value

the PLE systematic error content server

```
cohortMethodSystematicErrorViewer
```

The module viewer for rendering the PLE systematic error objects

# Description

The module viewer for rendering the PLE systematic error objects

#### Usage

```
cohortMethodSystematicErrorViewer(id)
```

# Arguments

id the unique reference id for the module

# Value

The user interface to the cohort method systematic error module

cohortMethodViewer

The viewer of the main cohort method module

# Description

The viewer of the main cohort method module

#### Usage

```
cohortMethodViewer(id)
```

#### Arguments

id

the unique reference id for the module

### Value

The user interface to the cohort method results viewer

cohortOverlapView

Cohort Overlap View

# Description

Use for customizing UI

# Usage

```
cohortOverlapView(id)
```

# Arguments

id

Name space  $\operatorname{Id}$  - use name spaced id  $\operatorname{ns}("\operatorname{cohortOverlap"})$  inside diagnostics Explorer module

 ${\tt compare Cohort Characterization View}$ 

 $compare\ characterization\ view$ 

# Description

Use for customizing UI

# Usage

```
compareCohortCharacterizationView(
  id,
  title = "Compare cohort characterization"
)
```

#### Arguments

id Namespace Id - use namespaced id ns("compareCohortCharacterization")

 $inside\ diagnostics Explorer\ module$ 

title Optional string title field

#### conceptsInDataSourceView

concepts In DataSource View

# Description

Use for customizing UI

#### Usage

```
conceptsInDataSourceView(id)
```

#### Arguments

id

Namespace Id - use namespaced id ns ("conceptsInDataSource") inside diagnostics Explorer module

createCdDatabaseDataSource

Create a CD data source from a database

#### Description

use this to create an interface to cohort diagnostics results data NOTE: I think this would make a good R6 class for other objects in this package so you could query them outside of a shiny app. E.g. if you wanted to make a custom R markdown template

# Usage

```
createCdDatabaseDataSource(
  connectionHandler,
  resultDatabaseSettings,
  dataModelSpecificationsPath = system.file("cohort-diagnostics-ref",
        "resultsDataModelSpecification.csv", package = utils::packageName()),
  dataMigrationsRef = system.file("cohort-diagnostics-ref", "migrations.csv", package =
        utils::packageName()),
  displayProgress = FALSE
)
```

32 createCustomColDefList

### Arguments

# connectionHandler

An instance of a ResultModelManager::connectionHander - manages a connection to a database.

resultDatabaseSettings

a list containing the result schema and prefixes

dataModelSpecificationsPath

The path to a file containing specifications for the data model used by the database.

data Migrations Ref

The path to a file listing all migrations for the data model that should have been applied

displayProgress

display a progress messaage (can only be used inside a shiny reactive context)

#### Value

An object of class 'CdDataSource'.

createCustomColDefList

Creating a list of custom column definitions for use in reactables

#### Description

Creating a list of custom column definitions for use in reactables

### Usage

```
createCustomColDefList(
  rawColNames,
  niceColNames = NULL,
  tooltipText = NULL,
  case = NULL,
  customColDefOptions = NULL)
```

#### Arguments

 ${\tt rawColNames} \qquad \text{The raw column names taken directly from the source data table that are} \\$ 

to be overwritten in the reactable

tooltipText The text to be displayed in a toolTip when hovering over the column in

the reactable

case Optional argument to convert raw column names to snake or camel case.

Defaults to NULL and preserves whatever raw column names are passed

 $_{
m in}$ 

customColDefOptions

A list of lists, where the inner lists are any custom options from reactable::colDef for each column

#### Value

A named list of reactable::colDef objects

```
{\tt createLargeSqlQueryDt} \quad \textit{Create Large Sql Query Data Table}
```

# Description

Construct an instance of a LargeDataTable R6 instance for use inside largeTableServer This should pass a parameterized sql query that can be used to iteratively return data from a table rather than returning the entire object.

# Usage

```
createLargeSqlQueryDt(
  connectionHandler = NULL,
  connectionDetails = NULL,
  baseQuery,
  countQuery = NULL
)
```

#### Arguments

connectionHandler

 $Result Model Manager\ connection Handler\ instance$ 

connectionDetails

DatabaseConnector connectionDetails instance

baseQuery

base sql query

countQuery

count query string (should match query). Can be auto generated with

sub query (default) but this will likely result in slow results

#### databaseInformationView

database Information View

# Description

Use for customizing UI

### Usage

databaseInformationView(id)

# Arguments

id

Name space  $\operatorname{Id}$  - use name spaced id  $\operatorname{ns}(\operatorname{"databaseInformation"})$  inside diagnostics Explorer module dataDiagnosticDrillServer

The module server for exploring prediction summary results

### Description

The module server for exploring prediction summary results

# Usage

dataDiagnosticDrillServer(id, connectionHandler, resultDatabaseSettings)

# Arguments

#### **Details**

The user specifies the id for the module

#### Value

The server to the summary module

```
dataDiagnosticDrillViewer
```

 $\label{lem:condition} The\ module\ viewer\ for\ exploring\ data-diagnostic\ results\ in\ more\ detail$ 

# Description

The module viewer for exploring data-diagnostic results in more detail

# Usage

```
dataDiagnosticDrillViewer(id)
```

### Arguments

id the unique reference id for the module

### Details

The user specifies the id for the module

#### Value

The user interface to the summary module

 ${\tt dataDiagnosticHelperFile}$ 

The location of the data-diagnostic module helper file

### Description

The location of the data-diagnostic module helper file

# Usage

```
dataDiagnosticHelperFile()
```

#### Details

Returns the location of the data-diagnostic helper file

#### Value

string location of the data-diagnostic helper file

 ${\tt dataDiagnosticServer}$  The module server for exploring data-diagnostic

### Description

The module server for exploring data-diagnostic

# Usage

```
dataDiagnosticServer(
  id = "dataDiag",
  connectionHandler,
  resultDatabaseSettings = list(port = 1)
)
```

# Arguments

```
\begin{tabular}{ll} id & the unique reference id for the module \\ connectionHandler & a connection to the database with the results \\ resultDatabaseSettings & a list containing the data-diagnostic result schema \\ \end{tabular}
```

### **Details**

The user specifies the id for the module

# Value

The server for the data-diagnostic module

dataDiagnosticSummaryServer

The module server for exploring prediction summary results

### Description

The module server for exploring prediction summary results

# Usage

 $\tt dataDiagnosticSummaryServer(id, connectionHandler, resultDatabaseSettings)$ 

# Arguments

#### **Details**

The user specifies the id for the module

#### Value

The server to the summary module

```
dataDiagnosticSummaryViewer
```

 $The\ module\ viewer\ for\ exploring\ data\mbox{-} diagnostic\ summary\ results$ 

# Description

The module viewer for exploring data-diagnostic summary results

# Usage

```
dataDiagnosticSummaryViewer(id)
```

### Arguments

id the unique reference id for the module

#### **Details**

The user specifies the id for the module

#### Value

The user interface to the summary module

data Diagnostic Viewer

# Description

The module viewer for exploring data-diagnostic

# Usage

```
dataDiagnosticViewer(id = "dataDiag")
```

# Arguments

 $\operatorname{id}$ 

the unique reference id for the module

# Details

The user specifies the id for the module

# Value

The user interface to the data-diagnostic viewer module

 ${\tt datasources Helper File} \ \ \textit{Define the helper file for the module}$ 

# Description

Define the helper file for the module

# Usage

```
datasourcesHelperFile()
```

#### Value

The helper html file for the datasources module

38 datasourcesViewer

datasourcesServer

The server function for the datasources module

#### Description

The server function for the datasources module

#### Usage

```
datasourcesServer(id, connectionHandler, resultDatabaseSettings)
```

#### Arguments

 $\begin{tabular}{ll} \begin{tabular}{ll} \beg$ 

A connection to the database with the results

result Database Settings

A named list containing the cohort generator results database details (schema, table prefix)

# Value

The server for the datasources module

datasourcesViewer

The viewer function for hte datasources module

#### Description

The viewer function for hte datasources module

#### Usage

```
datasourcesViewer(id)
```

# Arguments

id

The unique id for the datasources viewer namespace

# Value

The UI for the datasources module

evidenceSynthesisHelperFile

The location of the evidence synthesis module helper file

#### Description

The location of the evidence synthesis module helper file

#### Usage

```
evidenceSynthesisHelperFile()
```

#### Details

Returns the location of the evidence synthesis helper file

#### Value

string location of the evidence synthesis helper file

evidenceSynthesisServer

 $The \ module \ server \ for \ exploring \ Patient Level Prediction$ 

# Description

The module server for exploring PatientLevelPrediction

#### Usage

```
evidenceSynthesisServer(
  id,
  connectionHandler,
  resultDatabaseSettings = list(port = 1)
)
```

# Arguments

#### **Details**

The user specifies the id for the module

#### Value

The server for the PatientLevelPrediction module

evidenceSynthesisViewer

The module viewer for exploring evidence-synthesis

#### Description

The module viewer for exploring evidence-synthesis

#### Usage

```
evidenceSynthesisViewer(id = 1)
```

#### Arguments

id

the unique reference id for the module

#### Details

The user specifies the id for the module

#### Value

The user interface to the evidence-synthesis viewer module

```
getCirceRenderedExpression
```

Returns list with circe generated documentation

#### Description

Returns list with circe generated documentation

# Usage

```
getCirceRenderedExpression(
  cohortDefinition,
  cohortName = "Cohort Definition",
  includeConceptSets = FALSE
)
```

#### Arguments

cohortDefinition

An R object (list) with a list representation of the cohort definition expression, that may be converted to a cohort expression JSON using RJ-SONIO::toJSON(x = cohortDefinition, digits = 23, pretty = TRUE)

 $\begin{array}{ll} \textbf{CohortName} & \textbf{Name for the cohort definition} \\ \textbf{includeConceptSets} \end{array}$ 

Do you want to inclued concept set in the documentation

#### Value

list object

 ${\tt getEnabledCdReports}$ 

Get enable cd reports from available data

# Description

Get enable cd reports from available data

# Usage

getEnabledCdReports(dataSource)

#### Arguments

dataSource

 $\mathbf{C}$ 

getLogoImage

The location of the OHDSI logo

# Description

The location of the OHDSI logo

#### Usage

getLogoImage()

#### **Details**

Returns the location of the OHDSI logo

# Value

string location of the OHDSI logo

incidenceRatesView

incidence Rates View

# Description

Use for customizing UI

#### Usage

incidenceRatesView(id)

#### Arguments

id

Name space  $\operatorname{Id}$  - use name spaced id  $\operatorname{ns}(\operatorname{"incidenceRates"})$  inside diagnostics Explorer module 42 LargeDataTable

inclusionRulesView

inclusion Rules View

# Description

Use for customizing UI

#### Usage

inclusionRulesView(id)

#### Arguments

id

Name space  $\operatorname{Id}$  - use name spaced id  $\operatorname{ns}("\operatorname{inclusionRules"})$  inside diagnostics Explorer module

indexEventBreakdownView

Index event breakdown view

#### Description

Use for customizing UI

# Usage

indexEventBreakdownView(id)

# Arguments

id

Name space  $\operatorname{Id}$  - use name spaced id  $\operatorname{ns}(\operatorname{"indexEvents"})$  inside diagnostics Explorer module

LargeDataTable

Large Data Table

#### Description

Large data table R6 class.

Uses ResultModelManager::ConnectionHandler class to create paginating tables

NOTE Only currently works with sqlite and postgresql database backends (probably redshift too) as this method uses limit and offset for the queries

Alternatively, you might want to subclass this class. For example, if your backend query is against an API such as and ATLAS instance or ATHENA

If subclassing use inheritance and treat this class as an interface to implement - implementing the methods:

get

LargeDataTable 43

#### Public fields

```
baseQuery query string sql
countQuery count query string (should match query). Can be auto generated with sub
    query (default) but this will likely result in slow results
connectionHandler ResultModelManager connection handler to execute query inside ini-
    tialize
```

#### Methods

```
Public methods:
 • LargeDataTable$new()
 • LargeDataTable$getCount()
 • LargeDataTable$getPage()
 • LargeDataTable$getAllResults()
 • LargeDataTable$clone()
Method new():
 Usage:
 LargeDataTable$new(connectionHandler, baseQuery, countQuery = NULL)
 Arguments:
 connectionHandler ResultModelManager connectionHandler instance
 baseQuery base sql query
 countQuery count query string (should match query). Can be auto generated with sub
    query (default) but this will likely result in slow results
 Returns: self get count
Method getCount(): execute count query with specified parameters
 Usage:
 LargeDataTable$getCount(...)
 Arguments:
 Returns: count Get Page
```

#### Method getPage():

```
Usage:
LargeDataTable$getPage(pageNum, pageSize = self$pageSize, ...)
Arguments:
pageNum page number
pageSize page size
Returns: data.frame of query result get all results
```

## Method getAllResults():

```
Usage:
LargeDataTable$getAllResults(...)
Arguments:
```

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. . .

Returns: data.frame of all results. Used for large file downloads

Method clone(): The objects of this class are cloneable with this method.

Usage:

LargeDataTable\$clone(deep = FALSE)

Arguments:

deep Whether to make a deep clone.

largeTableServer

Large Table Component Server

## Description

Display large data tables in a consistent way - server side pagination for reactable objects

#### Usage

```
largeTableServer(
   id,
   ldt,
   inputParams,
   modifyData = NULL,
   columns = shiny::reactive(list()),
   ...
)
```

## Arguments

id Shiny module id. Must match Large Table Viewer

ldt LargeDataTable instance

inputParams reactive that returns list of parameters to be passed to ldt

modifyData optional callback function that takes the data page, page number, page

size as parameters must return data.frame compatable instance

 ${\bf columns} \qquad \qquad {\bf List \ or \ reactable \ returning \ list \ of \ reactable :: columnDef \ objects}$ 

... Additional reactable options (searchable, sortable

largeTableView 45

largeTableView

Large Table Component Viewer

#### Description

Components for results sets with many thousands of rows More limited than other table components in terms of automatic handling of search and filtering but will allow responsive apps

# Usage

```
largeTableView(
  id,
  pageSizeChoices = c(10, 25, 50, 100),
  selectedPageSize = 10,
  fullDownloads = TRUE
)
```

# Arguments

makeButtonLabel

Make a label for an html button

# Description

Make a label for an html button

#### Usage

```
makeButtonLabel(label)
```

# Arguments

label

The desired label for hte button

# Value

html code to make a button label

OhdsiShinyModules

Ohds i Shiny Modules

### Description

A selection of shiny modules for exploring standardized OHDSI results

orpahanConceptsView

Orphan Concepts View

## Description

Use for customizing UI

#### Usage

```
orpahanConceptsView(id)
```

## Arguments

id

Name space  $\operatorname{Id}$  - use name spaced id  $\operatorname{ns}(\operatorname{"orphanConcepts"})$  inside diagnostics Explorer module

patientLevelPredictionCalibrationServer

The module server for exploring prediction validation results

## Description

The module server for exploring prediction validation results

## Usage

```
patientLevelPredictionCalibrationServer(
   id,
   performanceId,
   connectionHandler,
   inputSingleView,
   resultDatabaseSettings
)
```

# Arguments

#### **Details**

The user specifies the id for the module

#### Value

The server to the prediction calibration module

```
patient Level Prediction Calibration Viewer\\
```

 $\label{lem:condition} \textit{The module viewer for exploring prediction model calibration } \textit{results}$ 

#### Description

The module viewer for exploring prediction model calibration results

### Usage

```
patientLevelPredictionCalibrationViewer(id)
```

#### Arguments

id

the unique reference id for the module

#### **Details**

The user specifies the id for the module

## Value

The user interface to the prediction model calibration module

```
patient Level Prediction Covariate Summary Server\\
```

 $\label{lem:covariate} The \ module \ server \ for \ exploring \ prediction \ covariate \ summary \ results$ 

# Description

The module server for exploring prediction covariate summary results

#### Usage

```
patientLevelPredictionCovariateSummaryServer(
   id,
   modelDesignId,
   developmentDatabaseId,
   performanceId,
   connectionHandler,
   inputSingleView,
   resultDatabaseSettings
)
```

#### Arguments

a list containing the result schema and prefixes

#### **Details**

The user specifies the id for the module

#### Value

The server to the covariate summary module

#### patientLevelPredictionCovariateSummaryViewer

The module viewer for exploring prediction covariate summary results

# Description

The module viewer for exploring prediction covariate summary results

## Usage

patientLevelPredictionCovariateSummaryViewer(id)

#### Arguments

id the unique reference id for the module

#### **Details**

The user specifies the id for the module

# Value

The user interface to the covariate summary module

```
patientLevelPredictionCutoffServer
```

The module server for exploring prediction cut-off results

#### Description

The module server for exploring prediction cut-off results

#### Usage

```
patientLevelPredictionCutoffServer(
   id,
   performanceId,
   connectionHandler,
   inputSingleView,
   resultDatabaseSettings
)
```

## Arguments

#### **Details**

The user specifies the id for the module

#### Value

The server to the prediction cut-off module

```
\verb|patientLevelPredictionCutoffViewer|
```

The module viewer for exploring prediction cut-off results

# Description

The module viewer for exploring prediction cut-off results

#### Usage

```
patientLevelPredictionCutoffViewer(id)
```

#### Arguments

id the unique reference id for the module

#### **Details**

The user specifies the id for the module

#### Value

The user interface to the prediction cut-off module

```
patient Level Prediction Design Summary Server\\
```

The module server for exploring prediction designs in the results

# Description

The module server for exploring prediction designs in the results

## Usage

```
patientLevelPredictionDesignSummaryServer(
  id,
  connectionHandler,
  resultDatabaseSettings
)
```

#### Arguments

```
\begin{tabular}{ll} \begin{tabular}{ll} the unique reference id for the module \\ \hline connectionHandler \\ \hline the connection to the prediction result database \\ \hline resultDatabaseSettings \\ & a list containing the result schema and prefixes \\ \hline \end{tabular}
```

#### **Details**

The user specifies the id for the module

# Value

The server to the prediction design module

```
patient Level Prediction Design Summary Viewer\\
```

 $\label{thm:condition} The \ module \ viewer for \ exploring \ prediction \ designs \ that \ have \ been \\ run$ 

#### Description

The module viewer for exploring prediction designs that have been run

#### Usage

```
patientLevelPredictionDesignSummaryViewer(id)
```

#### Arguments

id the unique reference id for the module

#### **Details**

The user specifies the id for the module

#### Value

The user interface to the prediction design module

```
patient Level Prediction Diagnostics Server\\
```

The module server for exploring prediction diagnostic results

#### Description

The module server for exploring prediction diagnostic results

## Usage

```
patientLevelPredictionDiagnosticsServer(
  id,
  modelDesignId,
  connectionHandler,
  resultDatabaseSettings
)
```

#### Arguments

#### **Details**

The user specifies the id for the module

#### Value

The server to the prediction diagnostic module

```
patientLevelPredictionDiagnosticsViewer
```

The module viewer for exploring prediction diagnostic results

#### Description

The module viewer for exploring prediction diagnostic results

#### Usage

```
patientLevelPredictionDiagnosticsViewer(id)
```

# Arguments

id

the unique reference id for the module

#### **Details**

The user specifies the id for the module

# Value

The user interface to the prediction diagnostic module

```
patient Level Prediction Discrimination Server \\
```

 $\label{lem:condition} \textit{The module server for exploring prediction model discrimination} \\ \textit{results}$ 

# Description

The module server for exploring prediction model discrimination results

#### Usage

```
patientLevelPredictionDiscriminationServer(
   id,
   performanceId,
   connectionHandler,
   inputSingleView,
   resultDatabaseSettings
)
```

#### Arguments

#### **Details**

The user specifies the id for the module

#### Value

The server to the model discrimination module

```
patient Level Prediction Discrimination Viewer\\
```

 $\label{lem:condition} The \ module \ viewer \ for \ exploring \ prediction \ model \ discrimination \ results$ 

# Description

The module viewer for exploring prediction model discrimination results

# Usage

```
patientLevelPredictionDiscriminationViewer(id)
```

# Arguments

id the unique reference id for the module

#### **Details**

The user specifies the id for the module

## Value

The user interface to the model discrimination results module

```
patientLevelPredictionHelperFile
```

The location of the prediction module helper file

# Description

The location of the prediction module helper file

#### Usage

```
patientLevelPredictionHelperFile()
```

#### **Details**

Returns the location of the prediction helper file

#### Value

string location of the prediction helper file

```
patient Level Prediction Model Summary Server\\
```

The module server for exploring prediction summary results

# Description

The module server for exploring prediction summary results

#### Usage

```
patientLevelPredictionModelSummaryServer(
   id,
   connectionHandler,
   resultDatabaseSettings,
   modelDesignId
)
```

# Arguments

#### **Details**

The user specifies the id for the module

# Value

The server to the summary module

```
patientLevelPredictionModelSummaryViewer
```

The module viewer for exploring prediction summary results

# Description

The module viewer for exploring prediction summary results

# Usage

```
patientLevelPredictionModelSummaryViewer(id)
```

#### Arguments

id

the unique reference id for the module

#### **Details**

The user specifies the id for the module

## Value

The user interface to the summary module

```
patientLevelPredictionNbServer
```

 $The \ module \ server \ for \ exploring \ prediction \ net-benefit \ results$ 

# Description

The module server for exploring prediction net-benefit results

# Usage

```
patientLevelPredictionNbServer(
  id,
  performanceId,
  connectionHandler,
  inputSingleView,
  resultDatabaseSettings
```

#### Arguments

#### **Details**

The user specifies the id for the module

#### Value

The server to the net-benefit module

#### patientLevelPredictionNbViewer

The module viewer for exploring prediction net-benefit results

# Description

The module viewer for exploring prediction net-benefit results

# Usage

```
patientLevelPredictionNbViewer(id)
```

# Arguments

id the unique reference id for the module

#### **Details**

The user specifies the id for the module

#### Value

The user interface to the net-benefit module

```
patientLevelPredictionServer
```

The module server for exploring PatientLevelPrediction

#### Description

The module server for exploring PatientLevelPrediction

#### Usage

```
patientLevelPredictionServer(
   id,
   connectionHandler,
   resultDatabaseSettings = list(port = 1)
)
```

# Arguments

# Details

The user specifies the id for the module

#### Value

The server for the PatientLevelPrediction module

```
patient Level Prediction Settings Server\\
```

The module server for exploring prediction settings

# Description

The module server for exploring prediction settings

#### Usage

```
patientLevelPredictionSettingsServer(
  id,
  modelDesignId,
  developmentDatabaseId,
  performanceId,
  connectionHandler,
  inputSingleView,
  resultDatabaseSettings
)
```

#### Arguments

#### **Details**

The user specifies the id for the module

#### Value

The server to the settings module

## patient Level Prediction Settings Viewer

 $The \ module \ viewer for \ exploring \ prediction \ settings$ 

# Description

The module viewer for exploring prediction settings

## Usage

```
patientLevelPredictionSettingsViewer(id)
```

## Arguments

id the unique reference id for the module

#### Details

The user specifies the id for the module

#### Value

The user interface to the settings module

```
patientLevelPredictionValidationServer
```

 $The\ module\ server\ for\ exploring\ prediction\ validation\ results$ 

# Description

The module server for exploring prediction validation results

# Usage

```
patientLevelPredictionValidationServer(
   id,
   modelDesignId,
   developmentDatabaseId,
   performanceId,
   connectionHandler,
   inputSingleView,
   resultDatabaseSettings
)
```

# Arguments

#### **Details**

The user specifies the id for the module

#### Value

The server to the validation module

#### patientLevelPredictionValidationViewer

The module viewer for exploring prediction validation results

# Description

The module viewer for exploring prediction validation results

#### Usage

```
patientLevelPredictionValidationViewer(id)
```

#### Arguments

id

the unique reference id for the module

#### **Details**

The user specifies the id for the module

#### Value

The user interface to the validation module

```
patientLevelPredictionViewer
```

 $The \ module \ viewer \ for \ exploring \ Patient Level Prediction$ 

# Description

The module viewer for exploring PatientLevelPrediction

# Usage

```
patientLevelPredictionViewer(id = 1)
```

# Arguments

id

the unique reference id for the module

#### **Details**

The user specifies the id for the module

#### Value

The user interface to the PatientLevelPrediction viewer module

phevaluatorHelperFile The location of the phevaluator module helper file

#### Description

The location of the phevaluator module helper file

#### Usage

phevaluatorHelperFile()

#### Details

Returns the location of the cohort-generator helper file

#### Value

String location of the phevaluator helper file

phevaluatorServer

The module server for the main phevaluator module

# Description

The module server for the main phevaluator module

## Usage

phevaluatorServer(id, connectionHandler, resultDatabaseSettings)

# Arguments

id The unique reference id for the module

connectionHandler

A connection to the database with the results

 ${\tt resultDatabaseSettings}$ 

A named list containing the cohort generator results database details (schema, table prefix)

#### Value

The phevaluator main module server

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phevaluatorViewer

The viewer of the phevaluator module

# Description

The viewer of the phevaluator module

## Usage

```
phevaluatorViewer(id)
```

#### Arguments

id

The unique reference id for the module

#### Value

The user interface to the phevaluator results viewer

resultTableServer

Result Table Server

# Description

Result Table Server

## Usage

```
resultTableServer(
   id,
   df,
   colDefsInput,
   selectedCols = NULL,
   sortedCols = NULL,
   elementId = NULL,
   addActions = NULL,
   downloadedFileName = NULL,
   groupBy = NULL
```

## Arguments

id string, table id must match resultsTableViewer function

df reactive that returns a data frame colDefsInput named list of reactable::colDefs

selectedCols string vector of columns the reactable should display to start by default.

Defaults to ALL if not specified.

sortedCols string vector of columns the reactable should sort by by default. Defaults

to no sort if not specified.

result Table Viewer 63

elementId optional string vector of element Id name for custom dropdown filtering

if present in the customColDef list. Defaults to NULL.

addActions add a button row selector column to the table to a column called 'actions'.

actions must be a column in df

downloadedFileName

string, desired name of downloaded data file. can use the name from the

module that is being used

groupBy The columns to group by

#### Value

shiny module server

resultTableViewer

Result Table Viewer

#### Description

Result Table Viewer

#### Usage

```
resultTableViewer(id = "result-table", downloadedFileName = NULL)
```

#### Arguments

id string
downloadedFileName

string, desired name of downloaded data file. can use the name from the

module that is being used

# Value

shiny module UI

sccsHelperFile

The location of the description module helper file

#### Description

The location of the description module helper file

#### Usage

sccsHelperFile()

#### **Details**

Returns the location of the description helper file

#### Value

string location of the description helper file

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sccsServer

 $The \ module \ server \ for \ exploring \ SCCS$ 

# Description

The module server for exploring SCCS

# Usage

```
sccsServer(id, connectionHandler, resultDatabaseSettings = list(port = 1))
```

#### Arguments

 ${\tt id} \hspace{1cm} {\tt the \ unique \ reference \ id \ for \ the \ module}$ 

connectionHandler

a connection to the database with the results

resultDatabaseSettings

a list containing the prediction result schema and connection details

#### **Details**

The user specifies the id for the module

# Value

The server for the PatientLevelPrediction module

sccsView

 $SCCS\ shiny\ module\ UI\ code$ 

# Description

Load the ui for the sccs module

#### Usage

```
sccsView(id = "sccs-module")
```

# Arguments

id

id for module

timeDistributionsView 65

 ${\tt timeDistributionsView} \quad timeDistributions \ view$ 

# Description

Use for customizing UI

# Usage

timeDistributionsView(id)

# Arguments

id

Name space  $\operatorname{Id}$  - use name spaced id ns("imeDistributions") inside diagnostics Explorer module

visitContextView

Visit context module view

# Description

Use for customizing UI

## Usage

visitContextView(id)

# Arguments

id

Name space  $\operatorname{Id}$  - use name spaced id  $\operatorname{ns}(\operatorname{"vistConext"})$  inside diagnostics Explorer module