# Package 'Strategus'

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```
Type Package
Title Coordinating and Executing Analytics Using HADES Modules
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Maintainer Anthony Sena <sena@ohdsi.org>
Description An R package for coordinating and executing analytics using HADES modules.
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BugReports https://github.com/OHDSI/Strategus/issues
Depends R (>= 4.2.0),
     CohortGenerator (>= 0.8.0),
     DatabaseConnector (>= 6.2.3)
Imports targets,
     renv (>= 1.0.0),
     ParallelLogger (>= 3.1.0),
     dplyr,
     checkmate,
     keyring,
     rlang,
     utils,
     R.utils,
     digest,
     methods,
     tibble,
     ResultModelManager (>= 0.3.0),
     SqlRender (>= 1.11.0),
     semver
Suggests testthat (>= 3.0.0),
     fs,
     knitr,
     rmarkdown,
     Eunomia,
     withr
Remotes ohdsi/CohortGenerator,
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```

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 ${\it add} {\it Module Specifications}$ 

Add module specifications to analysis specifications

### Description

Add module specifications to analysis specifications

### Usage

 ${\tt addModuleSpecifications(analysisSpecifications,\ moduleSpecifications)}$ 

### Arguments

analysisSpecifications

 $An \ object \ of \ type \ Analysis Specifications \ as \ created \ by \ create Empty Analysis Specifications \ module Specifications$ 

An object of type Module Specifications.

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#### Value

Returns the analysisSpecifications object with the module specifications added.

addSharedResources

Add shared resources to analysis specifications

#### **Description**

Add shared resources to analysis specifications

#### Usage

add Shared Resources (analysis Specifications, shared Resources)

### **Arguments**

analysisSpecifications

sharedResources

 $An \ object \ of \ type \ Analysis Specifications \ as \ created \ by \ create Empty Analysis Specificiations \ and \ created \ by \ create Empty Analysis Specific \ at ions \ and \ created \ by \ create Empty Analysis Specific \ at ions \ and \ created \ by \ create Empty Analysis Specific \ at ions \ and \ created \ by \ create Empty Analysis Specific \ at ions \ and \ created \ by \ create Empty Analysis Specific \ at ions \ and \ created \ by \ create Empty Analysis Specific \ at ions \ and \ created \ by \ create Empty Analysis Specific \ at ions \ and \ created \ by \ create Empty Analysis Specific \ at ions \ and \ created \ by \ create Empty Analysis Specific \ at ions \ and \ created \ by \ create Empty Analysis Specific \ at ions \ and \ created \ by \ create Empty Analysis Specific \ at ions \ and \ created \ by \ create Empty Analysis Specific \ at ions \ and \ created \ by \ create Empty Analysis Specific \ at ions \ and \ created \ by \ create Empty Analysis Specific \ at ions \ and \ created \ at ions \ at ions \ at ions \ at ions \ and \ at ions \ at i$ 

An object of type SharedResources.

#### Value

Returns the analysisSpecifications object with the module specifications added.

compareLockFiles

Compare two renv.lock files

#### **Description**

Used to compare renv.lock files and return the results in a data.frame. The return value will include a "full join" representation of the packages across the two lock files.

#### Usage

```
compareLockFiles(filename1, filename2)
```

### **Arguments**

filename1 The first renv.lock file name
filename2 The second renv.lock file name

### Value

A data.frame with the comparison of the rev.lock files

createCdmExecutionSettings

Create CDM execution settings

#### **Description**

Create CDM execution settings

### Usage

```
createCdmExecutionSettings(
  connectionDetailsReference,
  workDatabaseSchema,
  cdmDatabaseSchema,
  cohortTableNames = CohortGenerator::getCohortTableNames(cohortTable = "cohort"),
  tempEmulationSchema = getOption("sqlRenderTempEmulationSchema"),
  workFolder,
  resultsFolder,
  minCellCount = 5,
  integerAsNumeric = getOption("databaseConnectorIntegerAsNumeric", default = TRUE),
  integer64AsNumeric = getOption("databaseConnectorInteger64AsNumeric", default = TRUE),
  resultsConnectionDetailsReference = NULL,
  resultsDatabaseSchema = NULL
)
```

#### **Arguments**

connectionDetailsReference

A string that can be used to retrieve database connection details from a secure local store.

workDatabaseSchema

A database schema where intermediate data can be stored. The user (as identified in the connection details) will need to have write access to this database schema.

cdmDatabaseSchema

The database schema containing the data in CDM format. The user (as identified in the connection details) will need to have read access to this database schema.

cohortTableNames

An object identifying the various cohort table names that will be created in the workDatabaseSchema. This object can be created using the CohortGenerator::getCohortTableNafunction.

tempEmulationSchema

Some database platforms like Oracle and Impala do not truly support temp tables. To emulate temp tables, provide a schema with write privileges where temp tables can be created.

workFolder A folder in the local file system where intermediate results can be written.

resultsFolder A folder in the local file system where the module output will be written.

minCellCount 
The minimum number of subjects contributing to a count before it can be in-

cluded in results.

integerAsNumeric

Logical: should 32-bit integers be converted to numeric (double) values? If FALSE 32-bit integers will be represented using R's native Integer class. Default is TRUE

integer64AsNumeric

Logical: should 64-bit integers be converted to numeric (double) values? If FALSE 64-bit integers will be represented using bit64::integer64. Default is TRUE

 $results {\tt ConnectionDetails} Reference$ 

A string that can be used to retrieve the results database connection details from a secure local store.

 ${\tt resultsDatabaseSchema}$ 

A schema where the results tables are stored

#### Value

An object of type ExecutionSettings.

 $create {\tt EmptyAnalysisSpecificiations}$ 

Create an empty analysis specifications object.

#### **Description**

Create an empty analysis specifications object.

#### Usage

```
createEmptyAnalysisSpecificiations()
```

### Value

An object of type AnalysisSpecifications.

createResultDataModels

Create Result Data Models

### Description

Use this at the study design stage to create data models for modules This functions loads modules and executes any custom code to create schemas in a results database If recreate is set to TRUE all existing data will be removed, otherwise

### Usage

```
createResultDataModels(
  analysisSpecifications,
  executionSettings,
  executionScriptFolder = NULL,
  keyringName = NULL,
  restart = FALSE
)
```

#### **Arguments**

analysisSpecifications

 $An object of type \verb| AnalysisSpecifications| as created by \verb| createEmptyAnalysisSpecificiations| and the substitution of type and the substitution of type analysisSpecifications| as created by \verb| createEmptyAnalysisSpecifications| and the substitution of type analysisSpecifications| and the substitution of type analysisSpecification of$ 

executionSettings

 $An \,object\,of\,type\,{\tt ExecutionSettings}\,as\,created\,by\,{\tt createCdmExecutionSettings()}$ 

 $or\ create {\tt ResultsExecutionSettings()}.$ 

executionScriptFolder

Optional: the path to use for storing the execution script. when NULL, this

function will use a temporary file location to create the script to execute.

keyringName The name of the keyring to operate on. This function assumes you have created

the keyring before calling this function. It defaults to NULL to select the default keyring. If the keyring is password protected, the password must be stored in the environment variable STRATEGUS\_KEYRING\_PASSWORD so it is retrieved using the command Sys.getenv("STRATEGUS\_KEYRING\_PASSWORD")

restart Restart run? Requires executionScriptFolder to be specified, and be the

same as the executionScriptFolder used in the run to restart.

createResultsExecutionSettings

Create Results execution settings

### **Description**

Create Results execution settings

### Usage

```
createResultsExecutionSettings(
  resultsConnectionDetailsReference,
  resultsDatabaseSchema,
  workFolder,
  resultsFolder,
  minCellCount = 5,
  integerAsNumeric = getOption("databaseConnectorIntegerAsNumeric", default = TRUE),
  integer64AsNumeric = getOption("databaseConnectorInteger64AsNumeric", default = TRUE))
```

#### **Arguments**

resultsConnectionDetailsReference

A string that can be used to retrieve the results database connection details from

a secure local store.

resultsDatabaseSchema

A schema where the results tables are stored

workFolder A folder in the local file system where intermediate results can be written.

resultsFolder A folder in the local file system where the module output will be written.

minCellCount The minimum number of subjects contributing to a count before it can be in-

cluded in results.

integerAsNumeric

Logical: should 32-bit integers be converted to numeric (double) values? If FALSE 32-bit integers will be represented using R's native Integer class. Default is TRUE

integer64AsNumeric

Logical: should 64-bit integers be converted to numeric (double) values? If FALSE 64-bit integers will be represented using bit64::integer64. Default is TRUE

#### Value

An object of type ExecutionSettings.

ensureAllModulesInstantiated

Ensure all modules are instantiated

### **Description**

Ensure that all modules referenced in the analysis specifications are instantiated locally in the folder specified in the INSTANTIATED\_MODULES\_FOLDER environmental variable.

Missing modules will be fetched from remote repositories.

This function will also check whether there are different versions of the same module specified, which is not allowed, and whether all modules required by the specified modules are also instantiated.

### Usage

ensureAllModulesInstantiated(analysisSpecifications, forceVerification = FALSE)

#### Arguments

analysisSpecifications

An object of type AnalysisSpecifications as created by createEmptyAnalysisSpecificiations

forceVerification

When set to TRUE, the verification process is forced to re-evaluate if a module is properly installed. The default is FALSE since if a module is successfully validated, the module will contain the hash value of the module's renv.lock file in the file system so it can by-pass running this check every time.

#### Value

A list containing the install status of all modules (TRUE if all are installed properly) and a tibble listing the instantiated modules.

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execute

Execute analysis specifications.

#### **Description**

Execute analysis specifications.

#### Usage

```
execute(
  analysisSpecifications,
  executionSettings,
  executionScriptFolder = NULL,
  keyringName = NULL,
  restart = FALSE
)
```

#### **Arguments**

analysisSpecifications

An object of type AnalysisSpecifications as created by createEmptyAnalysisSpecificiations

executionSettings

An object of type ExecutionSettings as created by createCdmExecutionSettings() or createResultsExecutionSettings().

executionScriptFolder

Optional: the path to use for storing the execution script. when NULL, this function will use a temporary file location to create the script to execute.

keyringName

The name of the keyring to operate on. This function assumes you have created the keyring before calling this function. It defaults to NULL to select the default keyring. If the keyring is password protected, the password must be stored in the environment variable STRATEGUS\_KEYRING\_PASSWORD so it is retrieved using the command Sys.getenv("STRATEGUS\_KEYRING\_PASSWORD")

restart

Restart run? Requires executionScriptFolder to be specified, and be the same as the executionScriptFolder used in the run to restart.

#### Value

Does not return anything. Is called for the side-effect of executing the specified analyses.

getModuleList

Provides a list of HADES modules to run through Strategus

#### **Description**

This function provides a list of modules and their locations that may be used with Strategus.

#### Usage

```
getModuleList()
```

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#### Value

A data.frame() of modules that work with Strategus. This will contain: module = The name of the module version = The version of the module remote\_repo = The remote location of the module (i.e. github.com) remote\_username = The organization of the module (i.e. OHDSI) module\_type = 'cdm' or 'results'. 'cdm' refers to modules that are designed to work against patient level data in the OMOP CDM format. 'results' refers to modules that are designed to work against a results database containing output from a 'cdm' module.

retrieveConnectionDetails

Retrieve connection details from the secure location

#### **Description**

Retrieve connection details from the secure location

#### Usage

retrieveConnectionDetails(connectionDetailsReference, keyringName = NULL)

#### **Arguments**

connectionDetailsReference

A string that can be used to retrieve the settings from the secure store.

keyringName

The name of the keyring to operate on. This function assumes you have created the keyring before calling this function. It defaults to NULL to select the default keyring. If the keyring is password protected, the password must be stored in the environment variable STRATEGUS\_KEYRING\_PASSWORD so it is retrieved using the command Sys.getenv("STRATEGUS\_KEYRING\_PASSWORD")

### Value

Returns an object of type connectionDetails.

### See Also

storeConnectionDetails()

runSchemaCreation

Create module(s) result data model

#### Description

This function will create the results data model for the modules in the analysisSpecifications. A module can implement its own results data model creation function by implementing the function createDataModelSchema in its Main.R. The default behavior is to use the ResultsModelManager to create the results data model based on the resultsDataModelSpecification.csv in the module's results folder.

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#### **Usage**

```
runSchemaCreation(
  analysisSpecifications,
  keyringSettings,
 moduleIndex,
 executionSettings,
```

#### **Arguments**

analysisSpecifications

An object of type AnalysisSpecifications as created by createEmptyAnalysisSpecificiations

keyringSettings

The keyringSettings from the executionSettings context

moduleIndex

The index of the module in the analysis specification

executionSettings

An object of type ExecutionSettings as created by createCdmExecutionSettings()

or createResultsExecutionSettings().

For future expansion

storeConnectionDetails

Store connection details in a secure location

### **Description**

Store connection details in a secure location

### Usage

```
storeConnectionDetails(
  connectionDetails,
  connectionDetailsReference,
  keyringName = NULL
)
```

### Arguments

connectionDetails

An object of type connectionDetails as created by the DatabaseConnector::createConnectionI function.

 ${\tt connectionDetailsReference}$ 

A string that can be used to retrieve the settings from the secure store.

keyringName

The name of the keyring to operate on. This function assumes you have created the keyring before calling this function. It defaults to NULL to select the default keyring. If the keyring is password protected, the password must be stored in the environment variable STRATEGUS\_KEYRING\_PASSWORD so it is retrieved using the command Sys.getenv("STRATEGUS\_KEYRING\_PASSWORD")

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### Value

Does not return anything. Is called for the side effect of having the connection details stored.

#### See Also

retrieveConnectionDetails()

syncLockFile	Synchronize renv.lock files and overwrite the target file (read the description)

### Description

Used to synchronize the values from the "source of truth" renv.lock file to the target renv.lock file. Packages are compared (by name) and if the version of the package in the "source of truth" is greater the one found in the target, the target renv.lock file will be updated. This function will automatically update the target file.

Version comparison is handled by the semver package and since most packages use semantic versioning. When a package does not use semantic versioning, a warning is provided so the user can review

#### Usage

syncLockFile(sourceOfTruthLockFileName, targetLockFileName)

#### **Arguments**

sourceOfTruthLockFileName

 $\label{thm:continuity} The \ renv.lock \ file \ to \ use \ as \ the \ source \ of \ truth \\ \texttt{targetLockFileName}$ 

The target renv.lock file that will be synced with the source of truth

### Value

A data frame containing the different packages and their version that were involved in the synchronization process

unlockKeyring	Helper function to unlock a keyring	
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### Description

This helper function is used to unlock a keyring by using the password stored in Sys.getenv("STRATEGUS\_KEYRING\_I It will alert the user if the environment variable with the password is not set.

#### Usage

unlockKeyring(keyringName)

#### **Arguments**

keyringName

The name of the keyring to operate on. This function assumes you have created the keyring before calling this function. It defaults to NULL to select the default keyring. If the keyring is password protected, the password must be stored in the environment variable STRATEGUS\_KEYRING\_PASSWORD so it is retrieved using the command Sys.getenv("STRATEGUS\_KEYRING\_PASSWORD")

#### Value

Returns TRUE if the keyring was unlocked using the password otherwise it returns FALSE

validateLockFile

Validate an renv.lock file to ensure it is ready for use by Strategus

#### **Description**

Will check an renv.lock file for a module to verify that it only references tagged packages and includes the packages required by Strategus. It will also check for suggested packages that are useful for testing, such as RSQLite.

#### Usage

validateLockFile(filename)

#### **Arguments**

filename

The renv.lock file to validate

verifyModuleInstallation

Verify a module is properly installed

### **Description**

In some instances a module may fail to instantiate and install due to problems when calling renv::restore for the module's renv.lock file. This function will allow you to surface inconsistencies between the module renv.lock file and the module's renv project library. This function will check to that a module has been properly installed using internal functions of the renv package. If a module is verified to work via this function, the hash of the module's renv.lock file will be written to a text file in the module directory to indicate that it is ready for use. This will allow subsequent calls to work faster since the initial verification process can take some time.It is possible to re-run the verification of a module by using the forceVerification parameter.

To fix issues with a module, you will need to open the module's .Rproj in RStudio instance and debug the issues when calling renv::restore().

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#### **Usage**

```
verifyModuleInstallation(
  module,
  version,
  silent = FALSE,
  forceVerification = FALSE)
```

### **Arguments**

module The name of the module to verify (i.e. "CohortGeneratorModule")

version The version of the module to verify (i.e. "0.2.1")

silent When TRUE output of this verification process is suppressed

forceVerification

When set to TRUE, the verification process is forced to re-evaluate if a module is properly installed. The default is FALSE since if a module is successfully validated, the module will contain the hash value of the module's renv.lock file in the file system so it can by-pass running this check every time.

#### Value

A list with the output of the consistency check

withModuleRenv

Load module execution space inside and renv inspired by targets::tar\_script but allowing custom variable execution

### **Description**

Designed to allow more human readable code that is executed inside a module as well as simple variable substitution for injecting constants (e.g. simple parameters or file paths used inside and outside of modules)

#### Usage

```
withModuleRenv(
  code,
  moduleFolder,
  injectVars = list(),
  tempScriptFile = tempfile(fileext = ".R"),
  job = FALSE,
  processName = paste(moduleFolder, "_renv_run")
```

#### Arguments

code code block to execute

moduleFolder Instantiated Strategus module folder

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injectVars list of var names list(name=value) to replace (e.g. replace list(foo = "some

string") will find the pattern foo and replace it with the string some string -

be careful!

tempScriptFile tempFile to write script to

job run as rstudio job

processName String name for process

#### **Details**

This pattern also allows dependency injection which could be used if you don't want to use and renv and (instead) would like to use docker images or just execution in the base environment for testing/debugging

### Value

NULL invisibly

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