# Package 'OhdsiRTools'

January 18, 2017

Type Package
Title Tools for Maintaining OHDSI R Packages
Version 1.2.0
<b>Date</b> 2017-1-18
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<b>Description</b> Format and check syntax of R code and packages following the OHDSI R style guidelines. Support for parallel computation.
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<b>Depends</b> R (>= $3.1.0$ )
Imports devtools,     codetools,     formatR,     snow,     RJSONIO,     RCurl,     XML,     jsonlite,     methods,     utils
Suggests testthat
NeedsCompilation no
RoxygenNote 5.0.1
R topics documented:  checkUsagePackage 2 clusterApply 3 clusterRequire 4 convertArgsToList 4 createArgFunction 5 excludeFromList 5
formatRFile       6         formatRFolder       6         formatRText       7

2 checkUsagePackage

	insertCirceDefinitionInPackage	7
	loadSettingsFromJson	
	makeCluster	8
	matchInList	9
	OhdsiRTools	9
	prettyPrint	10
	restoreEnvironment	10
	saveSettingsToJson	1
	selectFromList	1
	stopCluster	12
	takeEnvironmentSnapshot	12
	updateCopyrightYearFile	13
	updateCopyrightYearFolder	13
Index		14

checkUsagePackage

Check all code in a package

## **Description**

Check all code in a package

## Usage

```
checkUsagePackage(package, ignoreHiddenFunctions = TRUE,
    suppressBindingKeywords = c("ggplot2", "ffwhich", "subset.ffdf", "glm"))
```

## **Arguments**

package The name of the package to check.

ignoreHiddenFunctions

Ignore functions for which the definition cannot be retrieved?

suppressBindingKeywords

A set of keywords that are indicative of non-standard evaluation.

## **Details**

This function uses the codetools package to check the code from problems. Heuristics are used to elimite false positives due to non-standard evaluation.

clusterApply 3

clusterApply	Apply a function to a list using the cluster
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## Description

Apply a function to a list using the cluster

#### Usage

```
clusterApply(cluster, x, fun, ..., stopOnError = FALSE, progressBar = TRUE,
    divideFfMemory = TRUE, setFfTempDir = TRUE)
```

## **Arguments**

cluster	ter The cluster of threads to run the function.	
x	The list on which the function will be applied.	
fun	The function to apply. Note that the context in which the function is specifies matters (see details).	
•••	Additional parameters for the function.	
stopOnError	Stop when one of the threads reports an error? If FALSE, all errors will be reported at the end.	
progressBar	Show a progress bar?	
divideFfMemory	When TRUE, the memory available for processing ff and ffdf objects will be equally divided over the threads.	
setFfTempDir	When TRUE, the ffTempDir option will be copied to each thread.	

## **Details**

The function will be executed on each element of x in the threads of the cluster. If there are more elements than threads, the elements will be queued. The progress bar will show the number of elements that have been completed. It can sometimes be important to realize that the context in which a function is created is also transmitted to the worker node. If a function is defined inside another function, and that outer function is called with a large argument, that argument will be transmitted to the worker node each time the function is executed. It can therefore make sense to define the function to be called at the package level rather than inside a function, to save overhead.

## Value

A list with the result of the function on each item in x.

4 convertArgsToList

clusterRequire

Require a package in the cluster

## **Description**

Require a package in the cluster

#### Usage

```
clusterRequire(cluster, package)
```

## **Arguments**

cluster The cluster object.

package The name of the package to load in all nodes.

 ${\tt convertArgsToList}$ 

Deprecated: Convert arguments used in call to a list

## Description

Deprecated: Convert arguments used in call to a list

## Usage

```
convertArgsToList(matchCall, resultClass = "list")
```

## **Arguments**

```
matchCall The result of match.call().
resultClass The class of the resulting object.
```

#### **Details**

Takes the argument values (both default and user-specified) and store them in a list. This function is deprecated because it fails when used in a function that is called using ::.

### Value

An object of the class specified in resultClass.

```
myFun <- function(x = 1, y = 2) {
  return(convertArgsToList(match.call()))
}</pre>
```

createArgFunction 5

|--|

## **Description**

Create an argument function

## Usage

```
createArgFunction(functionName, excludeArgs = c(), includeArgs = NULL,
  addArgs = list(), rCode = c(), newName)
```

## Arguments

functionName	The name of the function for which we want to create an args function.
excludeArgs	Exclude these arguments from appearing in the args function.
includeArgs	Include these arguments in the args function.
addArgs	Add these arguments to the args functions. Defined as a list with format name = default.
rCode	A character vector representing the R code where the new function should be appended to.
newName	The name of the new function. If not specified, the new name will be automati-

## **Details**

This function can be used to create a function that has (almost) the same interface as the specified function, and the output of this function will be a list of argument values.

#### Value

A character vector with the R code including the new function.

cally derived from the old name.

## **Examples**

```
createArgFunction("read.csv", addArgs = list(exposureId = "exposureId"))
```

excludeFromList Exclude variables from a list of objects of the same type

## Description

Exclude variables from a list of objects of the same type

## Usage

```
excludeFromList(x, exclude)
```

6 formatRFolder

## **Arguments**

x A list of objects of the same type.

exclude A character vector of names of variables to exclude.

 ${\tt formatRFile}$ 

Format an R file

## **Description**

Format an R file

## Usage

```
formatRFile(file, width.cutoff = 100)
```

## **Arguments**

file The path to the file.

width.cutoff Number of characters that each line should be limited to.

formatRFolder

Format all R files in a folder

## Description

Format all R files in a folder

## Usage

```
formatRFolder(path = ".", recursive = TRUE, skipAutogenerated = TRUE, ...)
```

## **Arguments**

Path to the folder containing the files to format. Only files with the .R extension

will be formatted.

recursive Include all subfolders?

skipAutogenerated

Skip autogenerated files such as RcppExports.R?

... Parameters to be passed on the the formatRFile function

```
## Not run:
formatRFolder()
## End(Not run)
```

formatRText 7

formatRText	Format R code
formatRText	Format R code

## **Description**

Format R code

#### Usage

```
formatRText(text, width.cutoff = 100)
```

## Arguments

text A character vector with the R code to be formatted.

width.cutoff Number of characters that each line should be limited to.

#### Value

A character vector with formatted R code.

insertCirceDefinitionInPackage

Load a Circe definition and insert it into this package

## **Description**

Load a Circe definition and insert it into this package

## Usage

```
insertCirceDefinitionInPackage(definitionId, name = NULL,
  baseUrl = "http://hix.jnj.com:8080/WebAPI")
```

## Arguments

definitionId The number indicating which Circe definition to fetch.

name The name that will be used for the json and SQL files. If not provided, the name

in Circe will be used, but this may not lead to valid file names.

baseUrl The base URL for the WebApi instance.

#### **Details**

Load a Circe definition from a WebApi instance and insert it into this package. This will fetch the json object and store it in the 'inst/circe' folder, and fetch the template SQL and store it in the 'inst/sql/sql\_server' folder. Both folders will be created if they don't exist.

8 makeCluster

## **Examples**

```
## Not run:
# This will create 'inst/circe/Angioedema.json' and 'inst/sql/sql_server/Angioedema.sql':
insertCirceDefinitionInPackage(2132, "Angioedema")
## End(Not run)
```

loadSettingsFromJson Load a settings object from a JSON file

## Description

Load a settings object from a JSON file

## Usage

loadSettingsFromJson(fileName)

## **Arguments**

fileName

Name of the JSON file to load.

## **Details**

Load a settings object from a JSON file, restoring object classes and attributes.

## Value

An R object as specified by the JSON.

makeCluster

Create a cluster of nodes for parallel computation

## Description

Create a cluster of nodes for parallel computation

## Usage

```
makeCluster(numberOfThreads, singleThreadToMain = TRUE)
```

## **Arguments**

numberOfThreads

Number of parallel threads.

singleThreadToMain

If numberOfThreads is 1, should we fall back to running the process in the main thread?

matchInList 9

## Value

An object representing the cluster.

matchInList

In a list of object of the same type, find those that match the input

## Description

In a list of object of the same type, find those that match the input

## Usage

```
matchInList(x, toMatch)
```

## **Arguments**

x A list of objects of the same type.

toMatch The object to match.

## **Details**

Typically, toMatch will contain a subset of the variables that are in the objects in the list. Any object matching all variables in toMatch will be included in the result.

## Value

A list of objects that match the toMatch object.

OhdsiRTools

OhdsiRTools

## Description

OhdsiRTools

10 restoreEnvironment

prettyPrint

Print a list of objects

## Description

Print a list of objects

## Usage

prettyPrint(object)

## **Arguments**

object

The list to print.

#### **Details**

Will print nested lists using indentation.

restoreEnvironment

Restore the R environment to a snapshot

## Description

Restore the R environment to a snapshot

## Usage

restoreEnvironment(snapshot, stopOnWrongRVersion = FALSE)

## **Arguments**

snapshot

The snapshot data frame as generated using the takeEnvironmentSnapshot

function.

stopOnWrongRVersion

Should the function stop when the wrong version of R is installed? Else just a warning will be thrown when the version doesn't match.

#### **Details**

This function restores the R environment to a previous snapshot, meaning all the packages will be restored to the versions they were at at the time of the snapshot. Note: on Windows you will very likely need to have RTools installed to build the various packages.

saveSettingsToJson 11

## **Examples**

```
## Not run:
snapshot <- takeEnvironmentSnapshot("OhdsiRTools")
write.csv(snapshot, "snapshot.csv")
# 5 years later
snapshot <- read.csv("snapshot.csv")
restoreEnvironment(snapshot)
## End(Not run)</pre>
```

 ${\tt save Settings To Json}$ 

Save a settings object as JSON file

## **Description**

Save a settings object as JSON file

## Usage

```
saveSettingsToJson(object, fileName)
```

## **Arguments**

object R object to be saved.

fileName File name where the object should be saved.

## **Details**

Save a setting object as a JSON file, using pretty formatting and preserving object classes and attributes.

selectFromList

Select variables from a list of objects of the same type

## **Description**

Select variables from a list of objects of the same type

## Usage

```
selectFromList(x, select)
```

## Arguments

x A list of objects of the same type.

select A character vector of names of variables to select.

stopCluster

Stop the cluster

## **Description**

Stop the cluster

## Usage

```
stopCluster(cluster)
```

## **Arguments**

cluster

The cluster to stop

take Environment Snapshot

Take a snapshot of the R environment

## Description

Take a snapshot of the R environment

## Usage

takeEnvironmentSnapshot(rootPackage)

#### **Arguments**

rootPackage

The name of the root package

## **Details**

This function records all versions used in the R environment that are used by one root package. This can be used for example to restore the environment to the state it was when a particular study package was run using the restoreEnvironment function.

#### Value

A data frame listing all the dependencies of the root package and their version numbers, in the order in which they should be installed.

```
snapshot <- takeEnvironmentSnapshot("OhdsiRTools")
snapshot</pre>
```

updateCopyrightYearFile

Update the copyright year in a R or SQL file

## Description

Update the copyright year in a R or SQL file

## Usage

```
updateCopyrightYearFile(file)
```

## Arguments

file

The path to the file.

updateCopyrightYearFolder

Update the copyright year in all R and SQL files in a folder

## Description

Update the copyright year in all R and SQL files in a folder

## Usage

```
updateCopyrightYearFolder(path = ".", recursive = TRUE)
```

## **Arguments**

path

Path to the folder containing the files to update. Only files with the .R and .SQL

extension will be updated.

recursive

Include all subfolders?

```
## Not run:
updateCopyrightYearFolder()
## End(Not run)
```

## **Index**

```
check Usage Package, \\ \\ 2
clusterApply, 3
clusterRequire, 4
convertArgsToList, 4
createArgFunction, 5
excludeFromList, 5
formatRFile, 6
formatRFolder, 6
formatRText, 7
insertCirceDefinitionInPackage, 7
loadSettingsFromJson, 8
makeCluster, 8
matchInList, 9
OhdsiRTools, 9
OhdsiRTools-package (OhdsiRTools), 9
prettyPrint, 10
restoreEnvironment, 10, 12
{\tt save Settings To Json, 11}
selectFromList, 11
stopCluster, 12
takeEnvironmentSnapshot, 10, 12
updateCopyrightYearFile, 13
updateCopyrightYearFolder, 13
```