

# Package ‘SelfControlledCaseSeries’

November 10, 2015

**Type** Package

**Title** Self-Controlled Case Series

**Version** 0.0.2

**Date** 2015-11-10

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**Description** SelfControlledCaseSeries is an R package for performing Self-Controlled Case Series (SCCS) analyses in an observational database in the OMOP Common Data Model

**Depends** R (>= 3.2.2),  
Cyclops (>= 1.2.0),  
DatabaseConnector (>= 1.3.0)

**Imports** RJDBC,  
SqlRender (>= 1.1.1),  
bit,  
ff,  
ffbase (>= 0.12.1),  
Rcpp (>= 0.11.2),  
OhdsiRTools,  
splines,  
ggplot2

**Suggests** testthat

**License** Apache License 2.0

**LinkingTo** Rcpp

**NeedsCompilation** yes

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createSccsEraData	Create SCCS era data
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## Description

Create SCCS era data

## Usage

```
createSccsEraData(sccsData, exposureId, outcomeId = NULL, naivePeriod = 0,
  firstOutcomeOnly = FALSE, includeExposureOfInterest = TRUE,
  exposureOfInterestSettings = createCovariateSettings(stratifyByID = TRUE,
  start = 0, end = 0, addExposedDaysToEnd = TRUE, splitPoints = c()),
  includePreExposureOfInterest = FALSE,
  preExposureOfInterestSetting = createCovariateSettings(stratifyByID = TRUE,
  mergeErasBeforeSplit = FALSE, start = -30, end = -1, addExposedDaysToEnd =
  FALSE, splitPoints = c()), covariateSettingsList = list(),
  includeAgeEffect = FALSE, ageKnots = 5, includeSeasonality = FALSE,
  seasonKnots = 5, eventDependentObservation = FALSE)
```

## Arguments

sccsData	An object of type sccsData as created using the <a href="#">getDbSccsData</a> function.
naivePeriod	The number of days at the start of a patient's observation period that should not be included in the risk calculations. Note that the naive period can be used to determine current covariate status right after the naive period, and whether an outcome is the first one.
firstOutcomeOnly	Whether only the first occurrence of an outcome should be considered.
covariateStart	Start day relative to the start of a covariate when the covariate should be considered in the risk profile.
covariatePersistencePeriod	Number of days after the end of the covariate when the risk is assumed to stop.
excludeConceptIds	Concept IDs that should be excluded from the list of covariates.

## Details

This function chops patient time into periods during which all covariates remain constant. The output details these periods, their durations, and a sparse representation of the covariate values.

## Value

An object of type sccsEraData.

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`getDbSccsData`*Todo: add title*

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**Description**

Todo: add description

**Usage**

```
getDbSccsData(connectionDetails, cdmDatabaseSchema,  
  oracleTempSchema = cdmDatabaseSchema,  
  outcomeDatabaseSchema = cdmDatabaseSchema,  
  outcomeTable = "condition_occurrence", outcomeIds,  
  outcomeConditionTypeConceptIds = c(),  
  exposureDatabaseSchema = cdmDatabaseSchema, exposureTable = "drug_era",  
  exposureIds = c(), excludeConceptIds = c(), drugEraCovariates = FALSE,  
  conditionEraCovariates = FALSE, procedureCovariates = FALSE,  
  visitCovariates = FALSE, observationCovariates = FALSE,  
  measurementCovariates = FALSE, deleteCovariatesSmallCount = 100,  
  cdmVersion = "4")
```

**Arguments**

`connectionDetails`

An R object of type `ConnectionDetails` created using the function `createConnectionDetails` in the `DatabaseConnector` package.

`cdmDatabaseSchema`

`oracleTempSchema`

`outcomeDatabaseSchema`

`outcomeTable`

`outcomeIds`

`outcomeConditionTypeConceptIds`

`exposureDatabaseSchema`

`exposureTable`

`exposureIds`

`excludeConceptIds`

`drugEraCovariates`

`conditionEraCovariates`

`procedureCovariates`

`visitCovariates`

observationCovariates

measurementCovariates

deleteCovariatesSmallCount

cdmVersion

## Details

Todo: add details

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loadSccsData	<i>Load the SCCS data from a folder</i>
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## Description

loadSccsData loads an object of type sccsData from a folder in the file system.

## Usage

```
loadSccsData(folder, readOnly = TRUE)
```

## Arguments

folder	The name of the folder containing the data.
readOnly	If true, the data is opened read only.

## Details

The data will be written to a set of files in the folder specified by the user.

## Value

An object of class cohortData.

## Examples

```
# todo
```

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loadSccsEraData	<i>Load the SCCS era data from a folder</i>
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**Description**

loadSccsEraData loads an object of type `sccsEraData` from a folder in the file system.

**Usage**

```
loadSccsEraData(folder, readOnly = FALSE)
```

**Arguments**

<code>readOnly</code>	If true, the data is opened read only.
<code>file</code>	The name of the folder containing the data.

**Details**

The data will be written to a set of files in the folder specified by the user.

**Value**

An object of class `sccsEraData`

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saveSccsData	<i>Save the SCCS data to folder</i>
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**Description**

`sccsData` saves an object of type `sccsData` to folder.

**Usage**

```
saveSccsData(sccsData, folder)
```

**Arguments**

<code>sccsData</code>	An object of type <code>sccsData</code> as generated using <a href="#">getDbSccsData</a> .
<code>folder</code>	The name of the folder where the data will be written. The folder should not yet exist.

**Details**

The data will be written to a set of files in the specified folder.

**Examples**

```
# todo
```

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saveSccsEraData	<i>Save the SCCS era data to folder</i>
-----------------	---

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**Description**

saveSccsEraData saves an object of type sccsEraData to folder.

**Usage**

```
saveSccsEraData(sccsEraData, folder)
```

**Arguments**

sccsEraData	An object of type sccsEraData as generated using <a href="#">createSccsEraData</a> .
folder	The name of the folder where the data will be written. The folder should not yet exist.

**Details**

The data will be written to a set of files in the specified folder.

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**Description**

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