

Package ‘AdhereR’

February 11, 2019

Type Package

Title Adherence to Medications

Version 0.4.1

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Description Computation of adherence to medications from Electronic Health care Data and visualization of individual medication histories and adherence patterns. The package implements a set of S3 classes and functions consistent with current adherence guidelines and definitions. It allows the computation of different measures of adherence (as defined in the literature, but also several original ones), their publication-quality plotting, the estimation of event duration and time to initiation, the interactive exploration of patient medication history and the real-time estimation of adherence given various parameter settings. It scales from very small datasets stored in flat CSV files to very large databases and from single-thread processing on mid-range consumer laptops to parallel processing on large heterogeneous computing clusters. It exposes a standardized interface allowing it to be used from other programming languages and platforms, such as Python.

URL <https://github.com/ddediu/AdhereR>

License GPL (>= 2)

LazyData TRUE

RoxygenNote 6.1.1

Imports lubridate (>= 1.5), parallel (>= 3.0), data.table (>= 1.9), manipulate (>= 1.0), shiny (>= 1.0), shinyWidgets (>= 0.4.4), shinyjs (>= 1.0), V8 (>= 1.5), colourpicker (>= 1.0), viridisLite (>= 0.3), highlight (>= 0.4), clipr (>= 0.4), knitr (>= 1.20), readODS (>= 1.6), readxl (>= 1.2), haven (>= 2.0), DBI (>= 1.0), RMariaDB (>= 1.0.5), RSQLite (>= 2.1)

Depends R (>= 3.0)

Suggests rmarkdown (≥ 1.1), R.rsp (≥ 0.40)

VignetteBuilder knitr, R.rsp

Encoding UTF-8

NeedsCompilation no

Repository CRAN

Date/Publication 2019-02-11 13:43:17 UTC

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callAdhereR	<i>callAdhereR.</i>
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Description

The function encapsulating all the logics that allows AdhereR to be called from any platform using the generic shell mechanism.

Usage

```
callAdhereR(shared.data.directory)
```

Arguments

`shared.data.directory`

A *string* containing the path to the directory where all the exchanged (shared) data (both input and output) is. AdhereR needs read and write access to this directory.

Details

In most cases this should not be done directly by the user, but instead used by an appropriate wrapper on the client platform. It allows transparent use of AdhereR from virtually any platform or programming language for which an appropriate wrapper is provided. For more details see the vignette describing the included reference Python 3 wrapper.

Value

This function displays any messages to the console, tries to also write them to the `Adherer-results.txt` file in the `shared.data.directory` directory, and, when finished, forces R to quit with a given shell error code:

- 0 The processing ended without major errors;
- 1 General error (hopefully there are messages in the `Adherer-results.txt` file);
- 10 The directory `shared.data.directory` does not exist;
- 11 AdhereR does not have read access to the `shared.data.directory` directory;
- 12 AdhereR does not have write access to the `shared.data.directory` directory;
- 13 issues with the parameters file `parameters.log`;
- 14 issues with the data file `dataset.csv`;
- 15 plotting issues;
- 16 interactive plotting issues;
- 17 issues exporting the results.

CMA0

*CMA0 constructor.***Description**

Constructs a basic CMA (continuous multiple-interval measures of medication availability/gaps) object.

Usage

```
CMA0(data = NULL, ID.colname = NA, event.date.colname = NA,
      event.duration.colname = NA, event.daily.dose.colname = NA,
      medication.class.colname = NA, carryover.within.obs.window = NA,
      carryover.into.obs.window = NA, carry.only.for.same.medication = NA,
      consider.dosage.change = NA, followup.window.start = 0,
      followup.window.start.unit = c("days", "weeks", "months", "years")[1],
      followup.window.duration = 365 * 2,
      followup.window.duration.unit = c("days", "weeks", "months",
      "years")[1], observation.window.start = 0,
      observation.window.start.unit = c("days", "weeks", "months",
      "years")[1], observation.window.duration = 365 * 2,
      observation.window.duration.unit = c("days", "weeks", "months",
      "years")[1], date.format = "%m/%d/%Y",
      summary = "Base CMA object", suppress.warnings = FALSE,
      arguments.that.should.not.be.defined = NULL, ...)
```

Arguments

<code>data</code>	A <i>data.frame</i> containing the medication events (prescribing or dispensing) used to compute the CMA. Must contain, at a minimum, the patient unique ID, the event date and duration, and might also contain the daily dosage and medication type (the actual column names are defined in the following four parameters).
<code>ID.colname</code>	A <i>string</i> , the name of the column in <i>data</i> containing the unique patient ID, or NA if not defined.
<code>event.date.colname</code>	A <i>string</i> , the name of the column in <i>data</i> containing the start date of the event (in the format given in the <code>date.format</code> parameter), or NA if not defined.
<code>event.duration.colname</code>	A <i>string</i> , the name of the column in <i>data</i> containing the event duration (in days), or NA if not defined.
<code>event.daily.dose.colname</code>	A <i>string</i> , the name of the column in <i>data</i> containing the prescribed daily dose, or NA if not defined.
<code>medication.class.colname</code>	A <i>string</i> , the name of the column in <i>data</i> containing the classes/types/groups of medication, or NA if not defined.

<code>carryover.within.obs.window</code>	<i>Logical</i> , if TRUE consider the carry-over within the observation window, or NA if not defined.
<code>carryover.into.obs.window</code>	<i>Logical</i> , if TRUE consider the carry-over from before the starting date of the observation window, or NA if not defined.
<code>carry.only.for.same.medication</code>	<i>Logical</i> , if TRUE the carry-over applies only across medications of the same type, or NA if not defined.
<code>consider.dosage.change</code>	<i>Logical</i> , if TRUE the carry-over is adjusted to reflect changes in dosage, or NA if not defined.
<code>followup.window.start</code>	If a Date object, it represents the actual start date of the follow-up window; if a <i>string</i> it is the name of the column in data containing the start date of the follow-up window either as the numbers of <code>followup.window.start.unit</code> units after the first event (the column must be of type <code>numeric</code>) or as actual dates (in which case the column must be of type <code>Date</code>); if a <i>number</i> it is the number of time units defined in the <code>followup.window.start.unit</code> parameter after the begin of the participant's first event; or NA if not defined.
<code>followup.window.start.unit</code>	can be either " <i>days</i> ", " <i>weeks</i> ", " <i>months</i> " or " <i>years</i> ", and represents the time units that <code>followup.window.start</code> refers to (when a number), or NA if not defined.
<code>followup.window.duration</code>	either a <i>number</i> representing the duration of the follow-up window in the time units given in <code>followup.window.duration.unit</code> , or a <i>string</i> giving the column containing these numbers. Should represent a period for which relevant medication events are recorded accurately (e.g. not extend after end of relevant treatment, loss-to-follow-up or change to a health care provider not covered by the database).
<code>followup.window.duration.unit</code>	can be either " <i>days</i> ", " <i>weeks</i> ", " <i>months</i> " or " <i>years</i> ", and represents the time units that <code>followup.window.duration</code> refers to, or NA if not defined.
<code>observation.window.start</code> , <code>observation.window.start.unit</code> , <code>observation.window.duration</code> , <code>observation.w</code>	the definition of the observation window (see the follow-up window parameters above for details).
<code>date.format</code>	A <i>string</i> giving the format of the dates used in the data and the other parameters; see the format parameters of the <code>as.Date</code> function for details (NB, this concerns only the dates given as strings and not as <code>Date</code> objects).
<code>summary</code>	Metadata as a <i>string</i> , briefly describing this CMA.
<code>suppress.warnings</code>	<i>Logical</i> , if TRUE don't show any warnings.
<code>arguments.that.should.not.be.defined</code>	a <i>list</i> of argument names and pre-defined values for which a warning should be thrown if passed to the function.
<code>...</code>	other possible parameters

Details

In most cases this should not be done directly by the user, but it is used internally by the other CMAs.

Value

An S3 object of class `CMAO` with the following fields:

- `data` The actual event (prescribing or dispensing) data, as given by the `data` parameter.
- `ID.colname` the name of the column in `data` containing the unique patient ID, as given by the `ID.colname` parameter.
- `event.date.colname` the name of the column in `data` containing the start date of the event (in the format given in the `date.format` parameter), as given by the `event.date.colname` parameter.
- `event.duration.colname` the name of the column in `data` containing the event duration (in days), as given by the `event.duration.colname` parameter.
- `event.daily.dose.colname` the name of the column in `data` containing the prescribed daily dose, as given by the `event.daily.dose.colname` parameter.
- `medication.class.colname` the name of the column in `data` containing the classes/types/groups of medication, as given by the `medication.class.colname` parameter.
- `carryover.within.obs.window` whether to consider the carry-over within the observation window, as given by the `carryover.within.obs.window` parameter.
- `carryover.into.obs.window` whether to consider the carry-over from before the starting date of the observation window, as given by the `carryover.into.obs.window` parameter.
- `carry.only.for.same.medication` whether the carry-over applies only across medication of the same type, as given by the `carry.only.for.same.medication` parameter.
- `consider.dosage.change` whether the carry-over is adjusted to reflect changes in dosage, as given by the `consider.dosage.change` parameter.
- `followup.window.start` the beginning of the follow-up window, as given by the `followup.window.start` parameter.
- `followup.window.start.unit` the time unit of the `followup.window.start`, as given by the `followup.window.start.unit` parameter.
- `followup.window.duration` the duration of the follow-up window, as given by the `followup.window.duration` parameter.
- `followup.window.duration.unit` the time unit of the `followup.window.duration`, as given by the `followup.window.duration.unit` parameter.
- `observation.window.start` the beginning of the observation window, as given by the `observation.window.start` parameter.
- `observation.window.start.unit` the time unit of the `observation.window.start`, as given by the `observation.window.start.unit` parameter.
- `observation.window.duration` the duration of the observation window, as given by the `observation.window.duration` parameter.
- `observation.window.duration.unit` the time unit of the `observation.window.duration`, as given by the `observation.window.duration.unit` parameter.

- `date.format` the format of the dates, as given by the `date.format` parameter.
- `summary` the metadata, as given by the `summary` parameter.

Examples

```
cma0 <- CMA0(data=med.events,
             ID.colname="PATIENT_ID",
             event.date.colname="DATE",
             event.duration.colname="DURATION",
             event.daily.dose.colname="PERDAY",
             medication.class.colname="CATEGORY",
             followup.window.start=0,
             followup.window.start.unit="days",
             followup.window.duration=2*365,
             followup.window.duration.unit="days",
             observation.window.start=30,
             observation.window.start.unit="days",
             observation.window.duration=365,
             observation.window.duration.unit="days",
             date.format="%m/%d/%Y",
             summary="Base CMA");
```

CMA1

CMA1 and CMA3 constructors.

Description

Constructs a CMA (continuous multiple-interval measures of medication availability/gaps) type 1 or type 3 object.

Usage

```
CMA1(data = NULL, ID.colname = NA, event.date.colname = NA,
      event.duration.colname = NA, followup.window.start = 0,
      followup.window.start.unit = c("days", "weeks", "months", "years")[1],
      followup.window.duration = 365 * 2,
      followup.window.duration.unit = c("days", "weeks", "months",
      "years")[1], observation.window.start = 0,
      observation.window.start.unit = c("days", "weeks", "months",
      "years")[1], observation.window.duration = 365 * 2,
      observation.window.duration.unit = c("days", "weeks", "months",
      "years")[1], date.format = "%m/%d/%Y", summary = NA,
      event.interval.colname = "event.interval",
      gap.days.colname = "gap.days",
      force.NA.CMA.for.failed.patients = TRUE, parallel.backend = c("none",
      "multicore", "snow", "snow(SOCK)", "snow(MPI)", "snow(NWS)")[1],
      parallel.threads = "auto", suppress.warnings = FALSE,
      arguments.that.should.not.be.defined = c(carryover.within.obs.window =
```

```
FALSE, carryover.into.obs.window = FALSE, carry.only.for.same.medication
= FALSE, consider.dosage.change = FALSE), ...)
```

```
CMA3(data = NULL, ID.colname = NA, event.date.colname = NA,
event.duration.colname = NA, followup.window.start = 0,
followup.window.start.unit = c("days", "weeks", "months", "years")[1],
followup.window.duration = 365 * 2,
followup.window.duration.unit = c("days", "weeks", "months",
"years")[1], observation.window.start = 0,
observation.window.start.unit = c("days", "weeks", "months",
"years")[1], observation.window.duration = 365 * 2,
observation.window.duration.unit = c("days", "weeks", "months",
"years")[1], date.format = "%m/%d/%Y", summary = NA,
event.interval.colname = "event.interval",
gap.days.colname = "gap.days",
force.NA.CMA.for.failed.patients = TRUE, parallel.backend = c("none",
"multicore", "snow", "snow(SOCK)", "snow(MPI)", "snow(NWS)")[1],
parallel.threads = "auto", suppress.warnings = FALSE,
arguments.that.should.not.be.defined = c(carryover.within.obs.window =
FALSE, carryover.into.obs.window = FALSE, carry.only.for.same.medication
= FALSE, consider.dosage.change = FALSE), ...)
```

Arguments

<code>data</code>	A data frame containing the events used to compute the CMA. Must contain, at a minimum, the patient unique ID, the event date and duration, and might also contain the daily dosage and medication type (the actual column names are defined in the following four parameters).
<code>ID.colname</code>	A <i>string</i> , the name of the column in <code>data</code> containing the unique patient ID; must be present.
<code>event.date.colname</code>	A <i>string</i> , the name of the column in <code>data</code> containing the start date of the event (in the format given in the <code>date.format</code> parameter); must be present.
<code>event.duration.colname</code>	A <i>string</i> , the name of the column in <code>data</code> containing the event duration (in days); must be present.
<code>followup.window.start</code>	If a Date object, it represents the actual start date of the follow-up window; if a <i>string</i> it is the name of the column in <code>data</code> containing the start date of the follow-up window either as the numbers of <code>followup.window.start.unit</code> units after the first event (the column must be of type <code>numeric</code>) or as actual dates (in which case the column must be of type <code>Date</code>); if a <i>number</i> it is the number of time units defined in the <code>followup.window.start.unit</code> parameter after the begin of the participant's first event; or NA if not defined.
<code>followup.window.start.unit</code>	can be either <i>"days"</i> , <i>"weeks"</i> , <i>"months"</i> or <i>"years"</i> , and represents the time units that <code>followup.window.start</code> refers to (when a number), or NA if not defined.

`followup.window.duration`

either a *number* representing the duration of the follow-up window in the time units given in `followup.window.duration.unit`, or a *string* giving the column containing these numbers. Should represent a period for which relevant medication events are recorded accurately (e.g. not extend after end of relevant treatment, loss-to-follow-up or change to a health care provider not covered by the database).

`followup.window.duration.unit`

can be either *"days"*, *"weeks"*, *"months"* or *"years"*, and represents the time units that `followup.window.duration` refers to, or NA if not defined.

`observation.window.start`, `observation.window.start.unit`, `observation.window.duration`, `observation.w`

the definition of the observation window (see the follow-up window parameters above for details).

`date.format`

A *string* giving the format of the dates used in the data and the other parameters; see the format parameters of the `as.Date` function for details (NB, this concerns only the dates given as strings and not as Date objects).

`summary`

Metadata as a *string*, briefly describing this CMA.

`event.interval.colname`

A *string*, the name of a newly-created column storing the number of days between the start of the current event and the start of the next one; the default value *"event.interval"* should be changed only if there is a naming conflict with a pre-existing *"event.interval"* column in `event.info`.

`gap.days.colname`

A *string*, the name of a newly-created column storing the number of days when medication was not available (i.e., the *"gap days"*); the default value *"gap.days"* should be changed only if there is a naming conflict with a pre-existing *"gap.days"* column in `event.info`.

`force.NA.CMA.for.failed.patients`

Logical describing how the patients for which the CMA estimation fails are treated: if TRUE they are returned with an NA CMA estimate, while for FALSE they are omitted.

`parallel.backend`

Can be *"none"* (the default) for single-threaded execution, *"multicore"* (using `mclapply` in package `parallel`) for multicore processing (NB. not currently implemented on MS Windows and automatically falls back on *"snow"* on this platform), or *"snow"*, *"snow(SOCK)"* (equivalent to *"snow"*), *"snow(MPI)"* or *"snow(NWS)"* specifying various types of SNOW clusters (can be on the local machine or more complex setups – please see the documentation of package `snow` for details; the last two require packages `Rmpi` and `nws`, respectively, not automatically installed with `AdhereR`).

`parallel.threads`

Can be *"auto"* (for `parallel.backend == "multicore"`, defaults to the number of cores in the system as given by `options("cores")`), while for `parallel.backend == "snow"`, defaults to 2), a strictly positive integer specifying the number of parallel threads, or a more complex specification of the SNOW cluster nodes for `parallel.backend == "snow"` (see the documentation of package `snow` for details).

suppress.warnings *Logical*, if TRUE don't show any warnings.

arguments.that.should.not.be.defined a *list* of argument names and pre-defined values for which a warning should be thrown if passed to the function.

... other possible parameters

Details

CMA1 considers the total number of days with medication supplied in all medication events in the observation window, excluding the last event. CMA3 is identical to CMA1 except that it is capped at 100%.

The formula is

$$(\text{number of days supply excluding last}) / (\text{first to last event})$$

Thus, the durations of all events are added up, possibly resulting in an CMA estimate (much) bigger than 1.0 (100%).

[CMA2](#) and CMA1 differ in the inclusion or not of the last event.

Value

An S3 object of class CMA1 (derived from CMA0) with the following fields:

- data The actual event data, as given by the data parameter.
- ID.colname the name of the column in data containing the unique patient ID, as given by the ID.colname parameter.
- event.date.colname the name of the column in data containing the start date of the event (in the format given in the date.format parameter), as given by the event.date.colname parameter.
- event.duration.colname the name of the column in data containing the event duration (in days), as given by the event.duration.colname parameter.
- event.daily.dose.colname the name of the column in data containing the prescribed daily dose, as given by the event.daily.dose.colname parameter.
- medication.class.colname the name of the column in data containing the classes/types/groups of medication, as given by the medication.class.colname parameter.
- followup.window.start the beginning of the follow-up window, as given by the followup.window.start parameter.
- followup.window.start.unit the time unit of the followup.window.start, as given by the followup.window.start.unit parameter.
- followup.window.duration the duration of the follow-up window, as given by the followup.window.duration parameter.
- followup.window.duration.unit the time unit of the followup.window.duration, as given by the followup.window.duration.unit parameter.
- observation.window.start the beginning of the observation window, as given by the observation.window.start parameter.

- `observation.window.start.unit` the time unit of the `observation.window.start`, as given by the `observation.window.start.unit` parameter.
- `observation.window.duration` the duration of the observation window, as given by the `observation.window.duration` parameter.
- `observation.window.duration.unit` the time unit of the `observation.window.duration`, as given by the `observation.window.duration.unit` parameter.
- `date.format` the format of the dates, as given by the `date.format` parameter.
- `summary` the metadata, as given by the `summary` parameter.
- `event.info` the data frame containing the event info (irrelevant for most users; see [compute.event.int.gaps](#) for details).
- `CMA` the data frame containing the actual CMA estimates for each participant (the `ID.colname` column).

See Also

CMAs 1 to 8 are described in:

Vollmer, W. M., Xu, M., Feldstein, A., Smith, D., Waterbury, A., & Rand, C. (2012). Comparison of pharmacy-based measures of medication adherence. *BMC Health Services Research*, **12**, 155. <http://doi.org/10.1186/1472-6963-12-155>.

Examples

```
cma1 <- CMA1(data=med.events,
             ID.colname="PATIENT_ID",
             event.date.colname="DATE",
             event.duration.colname="DURATION",
             followup.window.start=30,
             observation.window.start=30,
             observation.window.duration=365,
             date.format="%m/%d/%Y"
            );
cma3 <- CMA3(data=med.events,
             ID.colname="PATIENT_ID",
             event.date.colname="DATE",
             event.duration.colname="DURATION",
             followup.window.start=30,
             observation.window.start=30,
             observation.window.duration=365,
             date.format="%m/%d/%Y"
            );
```

CMA2

CMA2 and CMA4 constructors.

Description

Constructs a CMA (continuous multiple-interval measures of medication availability/gaps) type 2 or type 4 object.

Usage

```
CMA2(data = NULL, ID.colname = NA, event.date.colname = NA,
      event.duration.colname = NA, followup.window.start = 0,
      followup.window.start.unit = c("days", "weeks", "months", "years")[1],
      followup.window.duration = 365 * 2,
      followup.window.duration.unit = c("days", "weeks", "months",
      "years")[1], observation.window.start = 0,
      observation.window.start.unit = c("days", "weeks", "months",
      "years")[1], observation.window.duration = 365 * 2,
      observation.window.duration.unit = c("days", "weeks", "months",
      "years")[1], date.format = "%m/%d/%Y", summary = NA,
      event.interval.colname = "event.interval",
      gap.days.colname = "gap.days",
      force.NA.CMA.for.failed.patients = TRUE, parallel.backend = c("none",
      "multicore", "snow", "snow(SOCK)", "snow(MPI)", "snow(NWS)")[1],
      parallel.threads = "auto", suppress.warnings = FALSE,
      arguments.that.should.not.be.defined = c(carryover.within.obs.window =
      FALSE, carryover.into.obs.window = FALSE, carry.only.for.same.medication
      = FALSE, consider.dosage.change = FALSE), ...)
```

```
CMA4(data = NULL, ID.colname = NA, event.date.colname = NA,
      event.duration.colname = NA, followup.window.start = 0,
      followup.window.start.unit = c("days", "weeks", "months", "years")[1],
      followup.window.duration = 365 * 2,
      followup.window.duration.unit = c("days", "weeks", "months",
      "years")[1], observation.window.start = 0,
      observation.window.start.unit = c("days", "weeks", "months",
      "years")[1], observation.window.duration = 365 * 2,
      observation.window.duration.unit = c("days", "weeks", "months",
      "years")[1], date.format = "%m/%d/%Y", summary = NA,
      event.interval.colname = "event.interval",
      gap.days.colname = "gap.days",
      force.NA.CMA.for.failed.patients = TRUE, parallel.backend = c("none",
      "multicore", "snow", "snow(SOCK)", "snow(MPI)", "snow(NWS)")[1],
      parallel.threads = "auto", suppress.warnings = FALSE,
      arguments.that.should.not.be.defined = c(carryover.within.obs.window =
      FALSE, carryover.into.obs.window = FALSE, carry.only.for.same.medication
      = FALSE, consider.dosage.change = FALSE), ...)
```

Arguments

data	A data.frame containing the events used to compute the CMA. Must contain, at a minimum, the patient unique ID, the event date and duration, and might also contain the daily dosage and medication type (the actual column names are defined in the following four parameters).
ID.colname	A string, the name of the column in data containing the unique patient ID; must be present.

<code>event.date.colname</code>	A <i>string</i> , the name of the column in data containing the start date of the event (in the format given in the <code>date.format</code> parameter); must be present.
<code>event.duration.colname</code>	A <i>string</i> , the name of the column in data containing the event duration (in days); must be present.
<code>followup.window.start</code>	If a Date object, it represents the actual start date of the follow-up window; if a <i>string</i> it is the name of the column in data containing the start date of the follow-up window either as the numbers of <code>followup.window.start.unit</code> units after the first event (the column must be of type <code>numeric</code>) or as actual dates (in which case the column must be of type <code>Date</code>); if a <i>number</i> it is the number of time units defined in the <code>followup.window.start.unit</code> parameter after the begin of the participant's first event; or NA if not defined.
<code>followup.window.start.unit</code>	can be either " <i>days</i> ", " <i>weeks</i> ", " <i>months</i> " or " <i>years</i> ", and represents the time units that <code>followup.window.start</code> refers to (when a number), or NA if not defined.
<code>followup.window.duration</code>	either a <i>number</i> representing the duration of the follow-up window in the time units given in <code>followup.window.duration.unit</code> , or a <i>string</i> giving the column containing these numbers. Should represent a period for which relevant medication events are recorded accurately (e.g. not extend after end of relevant treatment, loss-to-follow-up or change to a health care provider not covered by the database).
<code>followup.window.duration.unit</code>	can be either " <i>days</i> ", " <i>weeks</i> ", " <i>months</i> " or " <i>years</i> ", and represents the time units that <code>followup.window.duration</code> refers to, or NA if not defined.
<code>observation.window.start</code> , <code>observation.window.start.unit</code> , <code>observation.window.duration</code> , <code>observation.window.duration.unit</code>	the definition of the observation window (see the follow-up window parameters above for details).
<code>date.format</code>	A <i>string</i> giving the format of the dates used in the data and the other parameters; see the format parameters of the <code>as.Date</code> function for details (NB, this concerns only the dates given as strings and not as <code>Date</code> objects).
<code>summary</code>	Metadata as a <i>string</i> , briefly describing this CMA.
<code>event.interval.colname</code>	A <i>string</i> , the name of a newly-created column storing the number of days between the start of the current event and the start of the next one; the default value "event.interval" should be changed only if there is a naming conflict with a pre-existing "event.interval" column in <code>event.info</code> .
<code>gap.days.colname</code>	A <i>string</i> , the name of a newly-created column storing the number of days when medication was not available (i.e., the "gap days"); the default value "gap.days" should be changed only if there is a naming conflict with a pre-existing "gap.days" column in <code>event.info</code> .
<code>force.NA.CMA.for.failed.patients</code>	<i>Logical</i> describing how the patients for which the CMA estimation fails are treated: if TRUE they are returned with an NA CMA estimate, while for FALSE they are omitted.

`parallel.backend`

Can be "none" (the default) for single-threaded execution, "multicore" (using `mclapply` in package `parallel`) for multicore processing (NB. not currently implemented on MS Windows and automatically falls back on "snow" on this platform), or "snow", "snow(SOCK)" (equivalent to "snow"), "snow(MPI)" or "snow(NWS)" specifying various types of SNOW clusters (can be on the local machine or more complex setups – please see the documentation of package `snow` for details; the last two require packages `Rmpi` and `nws`, respectively, not automatically installed with `AdhereR`).

`parallel.threads`

Can be "auto" (for `parallel.backend == "multicore"`, defaults to the number of cores in the system as given by `options("cores")`), while for `parallel.backend == "snow"`, defaults to 2), a strictly positive integer specifying the number of parallel threads, or a more complex specification of the SNOW cluster nodes for `parallel.backend == "snow"` (see the documentation of package `snow` for details).

`suppress.warnings`

Logical, if TRUE don't show any warnings.

`arguments.that.should.not.be.defined`

a *list* of argument names and pre-defined values for which a warning should be thrown if passed to the function.

...

other possible parameters

Details

CMA2 considers the total number of days with medication supplied in all medication events in the observation window, including the last event. CMA4 is identical to CMA2 except that it is capped at 100%.

The formula is

$$(number\ of\ days\ supply\ including\ last\ event) / (first\ to\ last\ event)$$

Thus, the durations of all events are added up, possibly resulting in an CMA estimate (much) bigger than 1.0 (100%)

CMA2 and [CMA1](#) differ in the inclusion or not of the last event.

Value

An S3 object of class `CMA2` (derived from `CMA0`) with the following fields:

- `data` The actual event data, as given by the `data` parameter.
- `ID.colname` the name of the column in `data` containing the unique patient ID, as given by the `ID.colname` parameter.
- `event.date.colname` the name of the column in `data` containing the start date of the event (in the format given in the `date.format` parameter), as given by the `event.date.colname` parameter.
- `event.duration.colname` the name of the column in `data` containing the event duration (in days), as given by the `event.duration.colname` parameter.

- `event.daily.dose.colname` the name of the column in data containing the prescribed daily dose, as given by the `event.daily.dose.colname` parameter.
- `medication.class.colname` the name of the column in data containing the classes/types/groups of medication, as given by the `medication.class.colname` parameter.
- `followup.window.start` the beginning of the follow-up window, as given by the `followup.window.start` parameter.
- `followup.window.start.unit` the time unit of the `followup.window.start`, as given by the `followup.window.start.unit` parameter.
- `followup.window.duration` the duration of the follow-up window, as given by the `followup.window.duration` parameter.
- `followup.window.duration.unit` the time unit of the `followup.window.duration`, as given by the `followup.window.duration.unit` parameter.
- `observation.window.start` the beginning of the observation window, as given by the `observation.window.start` parameter.
- `observation.window.start.unit` the time unit of the `observation.window.start`, as given by the `observation.window.start.unit` parameter.
- `observation.window.duration` the duration of the observation window, as given by the `observation.window.duration` parameter.
- `observation.window.duration.unit` the time unit of the `observation.window.duration`, as given by the `observation.window.duration.unit` parameter.
- `date.format` the format of the dates, as given by the `date.format` parameter.
- `summary` the metadata, as given by the `summary` parameter.
- `event.info` the data.frame containing the event info (irrelevant for most users; see `compute.event.int.gaps` for details).
- `CMA` the data.frame containing the actual CMA estimates for each participant (the `ID.colname` column).

See Also

CMAs 1 to 8 are defined in:

Vollmer, W. M., Xu, M., Feldstein, A., Smith, D., Waterbury, A., & Rand, C. (2012). Comparison of pharmacy-based measures of medication adherence. *BMC Health Services Research*, **12**, 155. <http://doi.org/10.1186/1472-6963-12-155>.

Examples

```
## Not run:
cma2 <- CMA2(data=med.events,
             ID.colname="PATIENT_ID",
             event.date.colname="DATE",
             event.duration.colname="DURATION",
             followup.window.start=30,
             observation.window.start=30,
             observation.window.duration=365,
             date.format="%m/%d/%Y")
```

```

    );
cma4 <- CMA4(data=med.events,
             ID.colname="PATIENT_ID",
             event.date.colname="DATE",
             event.duration.colname="DURATION",
             followup.window.start=30,
             observation.window.start=30,
             observation.window.duration=365,
             date.format="%m/%d/%Y"
           );
## End(Not run)

```

CMA5

CMA5 constructor.

Description

Constructs a CMA (continuous multiple-interval measures of medication availability/gaps) type 5 object.

Usage

```

CMA5(data = NULL, ID.colname = NA, event.date.colname = NA,
      event.duration.colname = NA, event.daily.dose.colname = NA,
      medication.class.colname = NA,
      carry.only.for.same.medication = FALSE,
      consider.dosage.change = FALSE, followup.window.start = 0,
      followup.window.start.unit = c("days", "weeks", "months", "years")[1],
      followup.window.duration = 365 * 2,
      followup.window.duration.unit = c("days", "weeks", "months",
      "years")[1], observation.window.start = 0,
      observation.window.start.unit = c("days", "weeks", "months",
      "years")[1], observation.window.duration = 365 * 2,
      observation.window.duration.unit = c("days", "weeks", "months",
      "years")[1], date.format = "%m/%d/%Y", summary = NA,
      event.interval.colname = "event.interval",
      gap.days.colname = "gap.days",
      force.NA.CMA.for.failed.patients = TRUE, parallel.backend = c("none",
      "multicore", "snow", "snow(SOCK)", "snow(MPI)", "snow(NWS)")[1],
      parallel.threads = "auto", suppress.warnings = FALSE,
      arguments.that.should.not.be.defined = c(carryover.within.obs.window =
      TRUE, carryover.into.obs.window = FALSE), ...)

```

Arguments

data A data.frame containing the medication events used to compute the CMA. Must contain, at a minimum, the patient unique ID, the event date and duration, and might also contain the daily dosage and medication type (the actual column names are defined in the following four parameters).

ID.colname	A <i>string</i> , the name of the column in data containing the unique patient ID; must be present.
event.date.colname	A <i>string</i> , the name of the column in data containing the start date of the event (in the format given in the <code>date.format</code> parameter); must be present.
event.duration.colname	A <i>string</i> , the name of the column in data containing the event duration (in days); must be present.
event.daily.dose.colname	A <i>string</i> , the name of the column in data containing the prescribed daily dose, or NA if not defined.
medication.class.colname	A <i>string</i> , the name of the column in data containing the medication type, or NA if not defined.
carry.only.for.same.medication	<i>Logical</i> , if TRUE, the carry-over applies only across medication of the same type.
consider.dosage.change	<i>Logical</i> , if TRUE, the carry-over is adjusted to also reflect changes in dosage.
followup.window.start	If a Date object, it represents the actual start date of the follow-up window; if a <i>string</i> it is the name of the column in data containing the start date of the follow-up window either as the numbers of <code>followup.window.start.unit</code> units after the first event (the column must be of type <code>numeric</code>) or as actual dates (in which case the column must be of type <code>Date</code>); if a <i>number</i> it is the number of time units defined in the <code>followup.window.start.unit</code> parameter after the begin of the participant's first event; or NA if not defined.
followup.window.start.unit	can be either " <i>days</i> ", " <i>weeks</i> ", " <i>months</i> " or " <i>years</i> ", and represents the time units that <code>followup.window.start</code> refers to (when a number), or NA if not defined.
followup.window.duration	either a <i>number</i> representing the duration of the follow-up window in the time units given in <code>followup.window.duration.unit</code> , or a <i>string</i> giving the column containing these numbers. Should represent a period for which relevant medication events are recorded accurately (e.g. not extend after end of relevant treatment, loss-to-follow-up or change to a health care provider not covered by the database).
followup.window.duration.unit	can be either " <i>days</i> ", " <i>weeks</i> ", " <i>months</i> " or " <i>years</i> ", and represents the time units that <code>followup.window.duration</code> refers to, or NA if not defined.
observation.window.start, observation.window.start.unit, observation.window.duration, observation.w	the definition of the observation window (see the follow-up window parameters above for details).
date.format	A <i>string</i> giving the format of the dates used in the data and the other parameters; see the format parameters of the <code>as.Date</code> function for details (NB, this concerns only the dates given as strings and not as <code>Date</code> objects).
summary	Metadata as a <i>string</i> , briefly describing this CMA.

<code>event.interval.colname</code>	A <i>string</i> , the name of a newly-created column storing the number of days between the start of the current event and the start of the next one; the default value "event.interval" should be changed only if there is a naming conflict with a pre-existing "event.interval" column in <code>event.info</code> .
<code>gap.days.colname</code>	A <i>string</i> , the name of a newly-created column storing the number of days when medication was not available (i.e., the "gap days"); the default value "gap.days" should be changed only if there is a naming conflict with a pre-existing "gap.days" column in <code>event.info</code> .
<code>force.NA.CMA.for.failed.patients</code>	<i>Logical</i> describing how the patients for which the CMA estimation fails are treated: if TRUE they are returned with an NA CMA estimate, while for FALSE they are omitted.
<code>parallel.backend</code>	Can be "none" (the default) for single-threaded execution, "multicore" (using <code>mclapply</code> in package <code>parallel</code>) for multicore processing (NB. not currently implemented on MS Windows and automatically falls back on "snow" on this platform), or "snow", "snow(SOCK)" (equivalent to "snow"), "snow(MPI)" or "snow(NWS)" specifying various types of SNOW clusters (can be on the local machine or more complex setups – please see the documentation of package <code>snow</code> for details; the last two require packages <code>Rmpi</code> and <code>nws</code> , respectively, not automatically installed with <code>AdhereR</code>).
<code>parallel.threads</code>	Can be "auto" (for <code>parallel.backend == "multicore"</code> , defaults to the number of cores in the system as given by <code>options("cores")</code>), while for <code>parallel.backend == "snow"</code> , defaults to 2), a strictly positive integer specifying the number of parallel threads, or a more complex specification of the SNOW cluster nodes for <code>parallel.backend == "snow"</code> (see the documentation of package <code>snow</code> for details).
<code>suppress.warnings</code>	<i>Logical</i> , if TRUE don't show any warnings.
<code>arguments.that.should.not.be.defined</code>	a <i>list</i> of argument names and pre-defined values for which a warning should be thrown if passed to the function.
<code>...</code>	other possible parameters

Details

CMA5 assumes that, within the observation window, the medication is used as prescribed and new medication is "banked" until needed (oversupply from previous events is used first, followed new medication supply). It computes days of theoretical use by extracting the total number of gap days from the total time interval between the first and the last event, accounting for carry over for all medication events within the observation window. Thus, it accounts for timing within the observation window, and excludes the remaining supply at the start of the last event within the observation window.

The formula is

$$(\text{number of days of theoretical use}) / (\text{first to last event})$$

Observations:

- the `carry.only.for.same.medication` parameter controls the transmission of carry-over across medication changes, producing a "standard" CMA5 (default value is FALSE), and an "alternative" CMA5b, respectively;
- the `consider.dosage.change` parameter controls if dosage changes are taken into account, i.e. if set as TRUE and a new medication event has a different daily dosage recommendation, carry-over is recomputed assuming medication use according to the new prescribed dosage (default value is FALSE).

Value

An S3 object of class CMA5 (derived from CMA0) with the following fields:

- `data` The actual event data, as given by the `data` parameter.
- `ID.colname` the name of the column in `data` containing the unique patient ID, as given by the `ID.colname` parameter.
- `event.date.colname` the name of the column in `data` containing the start date of the event (in the format given in the `date.format` parameter), as given by the `event.date.colname` parameter.
- `event.duration.colname` the name of the column in `data` containing the event duration (in days), as given by the `event.duration.colname` parameter.
- `event.daily.dose.colname` the name of the column in `data` containing the prescribed daily dose, as given by the `event.daily.dose.colname` parameter.
- `medication.class.colname` the name of the column in `data` containing the classes/types/groups of medication, as given by the `medication.class.colname` parameter.
- `carry.only.for.same.medication` whether the carry-over applies only across medication of the same type, as given by the `carry.only.for.same.medication` parameter.
- `consider.dosage.change` whether the carry-over is adjusted to reflect changes in dosage, as given by the `consider.dosage.change` parameter.
- `followup.window.start` the beginning of the follow-up window, as given by the `followup.window.start` parameter.
- `followup.window.start.unit` the time unit of the `followup.window.start`, as given by the `followup.window.start.unit` parameter.
- `followup.window.duration` the duration of the follow-up window, as given by the `followup.window.duration` parameter.
- `followup.window.duration.unit` the time unit of the `followup.window.duration`, as given by the `followup.window.duration.unit` parameter.
- `observation.window.start` the beginning of the observation window, as given by the `observation.window.start` parameter.
- `observation.window.start.unit` the time unit of the `observation.window.start`, as given by the `observation.window.start.unit` parameter.
- `observation.window.duration` the duration of the observation window, as given by the `observation.window.duration` parameter.

- `observation.window.duration.unit` the time unit of the `observation.window.duration`, as given by the `observation.window.duration.unit` parameter.
- `date.format` the format of the dates, as given by the `date.format` parameter.
- `summary` the metadata, as given by the `summary` parameter.
- `event.info` the data frame containing the event info (irrelevant for most users; see [compute.event.int.gaps](#) for details).
- `CMA` the data frame containing the actual CMA estimates for each participant (the `ID.colname` column).

See Also

CMAs 1 to 8 are defined in:

Vollmer, W. M., Xu, M., Feldstein, A., Smith, D., Waterbury, A., & Rand, C. (2012). Comparison of pharmacy-based measures of medication adherence. *BMC Health Services Research*, **12**, 155. <http://doi.org/10.1186/1472-6963-12-155>.

Examples

```
cma5 <- CMA5(data=med.events,
             ID.colname="PATIENT_ID",
             event.date.colname="DATE",
             event.duration.colname="DURATION",
             event.daily.dose.colname="PERDAY",
             medication.class.colname="CATEGORY",
             carry.only.for.same.medicament=FALSE,
             consider.dosage.change=FALSE,
             followup.window.start=30,
             observation.window.start=30,
             observation.window.duration=365,
             date.format="%m/%d/%Y"
             );
```

CMA6

CMA6 constructor.

Description

Constructs a CMA (continuous multiple-interval measures of medication availability/gaps) type 6 object.

Usage

```
CMA6(data = NULL, ID.colname = NA, event.date.colname = NA,
      event.duration.colname = NA, event.daily.dose.colname = NA,
      medication.class.colname = NA,
      carry.only.for.same.medicament = FALSE,
      consider.dosage.change = FALSE, followup.window.start = 0,
```

```

followup.window.start.unit = c("days", "weeks", "months", "years")[1],
followup.window.duration = 365 * 2,
followup.window.duration.unit = c("days", "weeks", "months",
"years")[1], observation.window.start = 0,
observation.window.start.unit = c("days", "weeks", "months",
"years")[1], observation.window.duration = 365 * 2,
observation.window.duration.unit = c("days", "weeks", "months",
"years")[1], date.format = "%m/%d/%Y", summary = NA,
event.interval.colname = "event.interval",
gap.days.colname = "gap.days",
force.NA.CMA.for.failed.patients = TRUE, parallel.backend = c("none",
"multicore", "snow", "snow(SOCK)", "snow(MPI)", "snow(NWS)")[1],
parallel.threads = "auto", suppress.warnings = FALSE,
arguments.that.should.not.be.defined = c(carryover.within.obs.window =
TRUE, carryover.into.obs.window = FALSE), ...)

```

Arguments

<code>data</code>	A data frame containing the events used to compute the CMA. Must contain, at a minimum, the patient unique ID, the event date and duration, and might also contain the daily dosage and medication type (the actual column names are defined in the following four parameters).
<code>ID.colname</code>	A <i>string</i> , the name of the column in <code>data</code> containing the unique patient ID; must be present.
<code>event.date.colname</code>	A <i>string</i> , the name of the column in <code>data</code> containing the start date of the event (in the format given in the <code>date.format</code> parameter); must be present.
<code>event.duration.colname</code>	A <i>string</i> , the name of the column in <code>data</code> containing the event duration (in days); must be present.
<code>event.daily.dose.colname</code>	A <i>string</i> , the name of the column in <code>data</code> containing the prescribed daily dose, or NA if not defined.
<code>medication.class.colname</code>	A <i>string</i> , the name of the column in <code>data</code> containing the medication type, or NA if not defined.
<code>carry.only.for.same.medication</code>	<i>Logical</i> , if TRUE, the carry-over applies only across medication of the same type.
<code>consider.dosage.change</code>	<i>Logical</i> , if TRUE, the carry-over is adjusted to also reflect changes in dosage.
<code>followup.window.start</code>	If a Date object, it represents the actual start date of the follow-up window; if a <i>string</i> it is the name of the column in <code>data</code> containing the start date of the follow-up window either as the numbers of <code>followup.window.start.unit</code> units after the first event (the column must be of type <code>numeric</code>) or as actual dates (in which case the column must be of type <code>Date</code>); if a <i>number</i> it is the number of time units defined in the <code>followup.window.start.unit</code> parameter after the begin of the participant's first event; or NA if not defined.

- `followup.window.start.unit`
can be either *"days"*, *"weeks"*, *"months"* or *"years"*, and represents the time units that `followup.window.start` refers to (when a number), or NA if not defined.
- `followup.window.duration`
either a *number* representing the duration of the follow-up window in the time units given in `followup.window.duration.unit`, or a *string* giving the column containing these numbers. Should represent a period for which relevant medication events are recorded accurately (e.g. not extend after end of relevant treatment, loss-to-follow-up or change to a health care provider not covered by the database).
- `followup.window.duration.unit`
can be either *"days"*, *"weeks"*, *"months"* or *"years"*, and represents the time units that `followup.window.duration` refers to, or NA if not defined.
- `observation.window.start`, `observation.window.start.unit`, `observation.window.duration`, `observation.window.duration.unit`
the definition of the observation window (see the follow-up window parameters above for details).
- `date.format` A *string* giving the format of the dates used in the data and the other parameters; see the format parameters of the `as.Date` function for details (NB, this concerns only the dates given as strings and not as Date objects).
- `summary` Metadata as a *string*, briefly describing this CMA.
- `event.interval.colname`
A *string*, the name of a newly-created column storing the number of days between the start of the current event and the start of the next one; the default value "event.interval" should be changed only if there is a naming conflict with a pre-existing "event.interval" column in `event.info`.
- `gap.days.colname`
A *string*, the name of a newly-created column storing the number of days when medication was not available (i.e., the "gap days"); the default value "gap.days" should be changed only if there is a naming conflict with a pre-existing "gap.days" column in `event.info`.
- `force.NA.CMA.for.failed.patients`
Logical describing how the patients for which the CMA estimation fails are treated: if TRUE they are returned with an NA CMA estimate, while for FALSE they are omitted.
- `parallel.backend`
Can be "none" (the default) for single-threaded execution, "multicore" (using `mclapply` in package `parallel`) for multicore processing (NB. not currently implemented on MS Windows and automatically falls back on "snow" on this platform), or "snow", "snow(SOCK)" (equivalent to "snow"), "snow(MPI)" or "snow(NWS)" specifying various types of SNOW clusters (can be on the local machine or more complex setups – please see the documentation of package `snow` for details; the last two require packages `Rmpi` and `nws`, respectively, not automatically installed with `AdhereR`).
- `parallel.threads`
Can be "auto" (for `parallel.backend == "multicore"`, defaults to the number of cores in the system as given by `options("cores")`), while for `parallel.backend == "snow"`, defaults to 2), a strictly positive integer specifying the number of

parallel threads, or a more complex specification of the SNOW cluster nodes for `parallel.backend == "snow"` (see the documentation of package `snow` for details).

`suppress.warnings`

Logical, if TRUE don't show any warnings.

`arguments.that.should.not.be.defined`

a *list* of argument names and pre-defined values for which a warning should be thrown if passed to the function.

...

other possible parameters

Details

CMA6 assumes that, within the observation window, the medication is used as prescribed and new medication is "banked" until needed (oversupply from previous events is used first, followed new medication supply). It computes days of theoretical use by extracting the total number of gap days from the total time interval between the first event and the end of the observation window, accounting for carry over for all medication events within the observation window. Thus, it accounts for timing within the observation window, and excludes the remaining supply at the end of the observation window.

The formula is

$$(\text{number of days of theoretical use}) / (\text{first event to end of observation window})$$

Observations:

- the `carry.only.for.same.medication` parameter controls the transmission of carry-over across medication changes, producing a "standard" CMA6 (default value is FALSE), and an "alternative" CMA6b, respectively;
- the `consider.dosage.change` parameter controls if dosage changes are taken into account, i.e. if set as TRUE and a new medication event has a different daily dosage recommendation, carry-over is recomputed assuming medication use according to the new prescribed dosage (default value is FALSE).

Value

An S3 object of class CMA6 (derived from CMA0) with the following fields:

- `data` The actual event data, as given by the `data` parameter.
- `ID.colname` the name of the column in `data` containing the unique patient ID, as given by the `ID.colname` parameter.
- `event.date.colname` the name of the column in `data` containing the start date of the event (in the format given in the `date.format` parameter), as given by the `event.date.colname` parameter.
- `event.duration.colname` the name of the column in `data` containing the event duration (in days), as given by the `event.duration.colname` parameter.
- `event.daily.dose.colname` the name of the column in `data` containing the prescribed daily dose, as given by the `event.daily.dose.colname` parameter.

- `medication.class.colname` the name of the column in data containing the classes/types/groups of medication, as given by the `medication.class.colname` parameter.
- `carry.only.for.same.medication` whether the carry-over applies only across medication of the same type, as given by the `carry.only.for.same.medication` parameter.
- `consider.dosage.change` whether the carry-over is adjusted to reflect changes in dosage, as given by the `consider.dosage.change` parameter.
- `followup.window.start` the beginning of the follow-up window, as given by the `followup.window.start` parameter.
- `followup.window.start.unit` the time unit of the `followup.window.start`, as given by the `followup.window.start.unit` parameter.
- `followup.window.duration` the duration of the follow-up window, as given by the `followup.window.duration` parameter.
- `followup.window.duration.unit` the time unit of the `followup.window.duration`, as given by the `followup.window.duration.unit` parameter.
- `observation.window.start` the beginning of the observation window, as given by the `observation.window.start` parameter.
- `observation.window.start.unit` the time unit of the `observation.window.start`, as given by the `observation.window.start.unit` parameter.
- `observation.window.duration` the duration of the observation window, as given by the `observation.window.duration` parameter.
- `observation.window.duration.unit` the time unit of the `observation.window.duration`, as given by the `observation.window.duration.unit` parameter.
- `date.format` the format of the dates, as given by the `date.format` parameter.
- `summary` the metadata, as given by the `summary` parameter.
- `event.info` the data.frame containing the event info (irrelevant for most users; see `compute.event.int.gaps` for details).
- `CMA` the data.frame containing the actual CMA estimates for each participant (the `ID.colname` column).

See Also

CMA6 1 to 8 are defined in:

Vollmer, W. M., Xu, M., Feldstein, A., Smith, D., Waterbury, A., & Rand, C. (2012). Comparison of pharmacy-based measures of medication adherence. *BMC Health Services Research*, **12**, 155. <http://doi.org/10.1186/1472-6963-12-155>.

Examples

```
cma6 <- CMA6(data=med.events,
             ID.colname="PATIENT_ID",
             event.date.colname="DATE",
             event.duration.colname="DURATION",
             event.daily.dose.colname="PERDAY",
             medication.class.colname="CATEGORY",
             carry.only.for.same.medication=FALSE,
```

```

consider.dosage.change=FALSE,
followup.window.start=30,
observation.window.start=30,
observation.window.duration=365,
date.format="%m/%d/%Y"
);

```

CMA7

CMA7 constructor.

Description

Constructs a CMA (continuous multiple-interval measures of medication availability/gaps) type 7 object.

Usage

```

CMA7(data = NULL, ID.colname = NA, event.date.colname = NA,
      event.duration.colname = NA, event.daily.dose.colname = NA,
      medication.class.colname = NA,
      carry.only.for.same.medicament = FALSE,
      consider.dosage.change = FALSE, followup.window.start = 0,
      followup.window.start.unit = c("days", "weeks", "months", "years")[1],
      followup.window.duration = 365 * 2,
      followup.window.duration.unit = c("days", "weeks", "months",
      "years")[1], observation.window.start = 0,
      observation.window.start.unit = c("days", "weeks", "months",
      "years")[1], observation.window.duration = 365 * 2,
      observation.window.duration.unit = c("days", "weeks", "months",
      "years")[1], date.format = "%m/%d/%Y", summary = NA,
      event.interval.colname = "event.interval",
      gap.days.colname = "gap.days",
      force.NA.CMA.for.failed.patients = TRUE, parallel.backend = c("none",
      "multicore", "snow", "snow(SOCK)", "snow(MPI)", "snow(NWS)")[1],
      parallel.threads = "auto", suppress.warnings = FALSE,
      arguments.that.should.not.be.defined = c(carryover.within.obs.window =
      TRUE, carryover.into.obs.window = TRUE), ...)

```

Arguments

data	A data frame containing the events used to compute the CMA. Must contain, at a minimum, the patient unique ID, the event date and duration, and might also contain the daily dosage and medication type (the actual column names are defined in the following four parameters).
ID.colname	A <i>string</i> , the name of the column in data containing the unique patient ID; must be present.

<code>event.date.colname</code>	A <i>string</i> , the name of the column in data containing the start date of the event (in the format given in the <code>date.format</code> parameter); must be present.
<code>event.duration.colname</code>	A <i>string</i> , the name of the column in data containing the event duration (in days); must be present.
<code>event.daily.dose.colname</code>	A <i>string</i> , the name of the column in data containing the prescribed daily dose, or NA if not defined.
<code>medication.class.colname</code>	A <i>string</i> , the name of the column in data containing the medication type, or NA if not defined.
<code>carry.only.for.same.medication</code>	<i>Logical</i> , if TRUE, the carry-over applies only across medication of the same type.
<code>consider.dosage.change</code>	<i>Logical</i> , if TRUE, the carry-over is adjusted to also reflect changes in dosage.
<code>followup.window.start</code>	If a Date object, it represents the actual start date of the follow-up window; if a <i>string</i> it is the name of the column in data containing the start date of the follow-up window either as the numbers of <code>followup.window.start.unit</code> units after the first event (the column must be of type <code>numeric</code>) or as actual dates (in which case the column must be of type <code>Date</code>); if a <i>number</i> it is the number of time units defined in the <code>followup.window.start.unit</code> parameter after the begin of the participant's first event; or NA if not defined.
<code>followup.window.start.unit</code>	can be either " <i>days</i> ", " <i>weeks</i> ", " <i>months</i> " or " <i>years</i> ", and represents the time units that <code>followup.window.start</code> refers to (when a number), or NA if not defined.
<code>followup.window.duration</code>	either a <i>number</i> representing the duration of the follow-up window in the time units given in <code>followup.window.duration.unit</code> , or a <i>string</i> giving the column containing these numbers. Should represent a period for which relevant medication events are recorded accurately (e.g. not extend after end of relevant treatment, loss-to-follow-up or change to a health care provider not covered by the database).
<code>followup.window.duration.unit</code>	can be either " <i>days</i> ", " <i>weeks</i> ", " <i>months</i> " or " <i>years</i> ", and represents the time units that <code>followup.window.duration</code> refers to, or NA if not defined.
<code>observation.window.start</code> , <code>observation.window.start.unit</code> , <code>observation.window.duration</code> , <code>observation.w</code>	the definition of the observation window (see the follow-up window parameters above for details).
<code>date.format</code>	A <i>string</i> giving the format of the dates used in the data and the other parameters; see the format parameters of the <code>as.Date</code> function for details (NB, this concerns only the dates given as strings and not as Date objects).
<code>summary</code>	Metadata as a <i>string</i> , briefly describing this CMA.
<code>event.interval.colname</code>	A <i>string</i> , the name of a newly-created column storing the number of days between the start of the current event and the start of the next one; the default

value "event.interval" should be changed only if there is a naming conflict with a pre-existing "event.interval" column in `event.info`.

`gap.days.colname`

A *string*, the name of a newly-created column storing the number of days when medication was not available (i.e., the "gap days"); the default value "gap.days" should be changed only if there is a naming conflict with a pre-existing "gap.days" column in `event.info`.

`force.NA.CMA.for.failed.patients`

Logical describing how the patients for which the CMA estimation fails are treated: if TRUE they are returned with an NA CMA estimate, while for FALSE they are omitted.

`parallel.backend`

Can be "none" (the default) for single-threaded execution, "multicore" (using `mclapply` in package `parallel`) for multicore processing (NB. not currently implemented on MS Windows and automatically falls back on "snow" on this platform), or "snow", "snow(SOCK)" (equivalent to "snow"), "snow(MPI)" or "snow(NWS)" specifying various types of SNOW clusters (can be on the local machine or more complex setups – please see the documentation of package `snow` for details; the last two require packages `Rmpi` and `nws`, respectively, not automatically installed with `AdhereR`).

`parallel.threads`

Can be "auto" (for `parallel.backend == "multicore"`, defaults to the number of cores in the system as given by `options("cores")`), while for `parallel.backend == "snow"`, defaults to 2), a strictly positive integer specifying the number of parallel threads, or a more complex specification of the SNOW cluster nodes for `parallel.backend == "snow"` (see the documentation of package `snow` for details).

`suppress.warnings`

Logical, if TRUE don't show any warnings.

`arguments.that.should.not.be.defined`

a *list* of argument names and pre-defined values for which a warning should be thrown if passed to the function.

...

other possible parameters

Details

CMA7 assumes that, within and before the observation window, the medication is used as prescribed and new medication is "banked" until needed (oversupply from previous events is used first, followed new medication supply). It computes days of theoretical use by extracting the total number of gap days from the total time interval between the start and the end of the observation window, accounting for carry over for all medication events within and before the observation window. All medication events in the follow up window before observation window are considered for carry-over calculation. Thus, it accounts for timing within and before the observation window, and excludes the remaining supply at the end of the observation window.

The formula is

$$(\text{number of days of theoretical use}) / (\text{start to end of observation window})$$

Observations:

- the `carry.only.for.same.medication` parameter controls the transmission of carry-over across medication changes, producing a "standard" CMA7 (default value is FALSE), and an "alternative" CMA7b, respectively;
- the `consider.dosage.change` parameter controls if dosage changes are taken into account, i.e. if set as TRUE and a new medication event has a different daily dosage recommendation, carry-over is recomputed assuming medication use according to the new prescribed dosage (default value is FALSE).

Value

An S3 object of class CMA7 (derived from CMA0) with the following fields:

- `data` The actual event data, as given by the `data` parameter.
- `ID.colname` the name of the column in `data` containing the unique patient ID, as given by the `ID.colname` parameter.
- `event.date.colname` the name of the column in `data` containing the start date of the event (in the format given in the `date.format` parameter), as given by the `event.date.colname` parameter.
- `event.duration.colname` the name of the column in `data` containing the event duration (in days), as given by the `event.duration.colname` parameter.
- `event.daily.dose.colname` the name of the column in `data` containing the prescribed daily dose, as given by the `event.daily.dose.colname` parameter.
- `medication.class.colname` the name of the column in `data` containing the classes/types/groups of medication, as given by the `medication.class.colname` parameter.
- `carry.only.for.same.medication` whether the carry-over applies only across medication of the same type, as given by the `carry.only.for.same.medication` parameter.
- `consider.dosage.change` whether the carry-over is adjusted to reflect changes in dosage, as given by the `consider.dosage.change` parameter.
- `followup.window.start` the beginning of the follow-up window, as given by the `followup.window.start` parameter.
- `followup.window.start.unit` the time unit of the `followup.window.start`, as given by the `followup.window.start.unit` parameter.
- `followup.window.duration` the duration of the follow-up window, as given by the `followup.window.duration` parameter.
- `followup.window.duration.unit` the time unit of the `followup.window.duration`, as given by the `followup.window.duration.unit` parameter.
- `observation.window.start` the beginning of the observation window, as given by the `observation.window.start` parameter.
- `observation.window.start.unit` the time unit of the `observation.window.start`, as given by the `observation.window.start.unit` parameter.
- `observation.window.duration` the duration of the observation window, as given by the `observation.window.duration` parameter.

- `observation.window.duration.unit` the time unit of the `observation.window.duration`, as given by the `observation.window.duration.unit` parameter.
- `date.format` the format of the dates, as given by the `date.format` parameter.
- `summary` the metadata, as given by the `summary` parameter.
- `event.info` the data frame containing the event info (irrelevant for most users; see [compute.event.int.gaps](#) for details).
- `CMA` the data frame containing the actual CMA estimates for each participant (the `ID.colname` column).

Examples

```
cma7 <- CMA7(data=med.events,
             ID.colname="PATIENT_ID",
             event.date.colname="DATE",
             event.duration.colname="DURATION",
             event.daily.dose.colname="PERDAY",
             medication.class.colname="CATEGORY",
             carry.only.for.same.medicament=FALSE,
             consider.dosage.change=FALSE,
             followup.window.start=30,
             observation.window.start=30,
             observation.window.duration=365,
             date.format="%m/%d/%Y"
            );
```

CMA8

CMA8 constructor.

Description

Constructs a CMA (continuous multiple-interval measures of medication availability/gaps) type 8 object.

Usage

```
CMA8(data = NULL, ID.colname = NA, event.date.colname = NA,
      event.duration.colname = NA, event.daily.dose.colname = NA,
      medication.class.colname = NA,
      carry.only.for.same.medicament = FALSE,
      consider.dosage.change = FALSE, followup.window.start = 0,
      followup.window.start.unit = c("days", "weeks", "months", "years")[1],
      followup.window.duration = 365 * 2,
      followup.window.duration.unit = c("days", "weeks", "months",
      "years")[1], observation.window.start = 0,
      observation.window.start.unit = c("days", "weeks", "months",
      "years")[1], observation.window.duration = 365 * 2,
      observation.window.duration.unit = c("days", "weeks", "months",
```

```
"years")[1], date.format = "%m/%d/%Y", summary = NA,
event.interval.colname = "event.interval",
gap.days.colname = "gap.days",
force.NA.CMA.for.failed.patients = TRUE, parallel.backend = c("none",
"multicore", "snow", "snow(SOCK)", "snow(MPI)", "snow(NWS)")[1],
parallel.threads = "auto", suppress.warnings = FALSE,
arguments.that.should.not.be.defined = c(carryover.within.obs.window =
TRUE, carryover.into.obs.window = TRUE), ...)
```

Arguments

<code>data</code>	A data frame containing the events used to compute the CMA. Must contain, at a minimum, the patient unique ID, the event date and duration, and might also contain the daily dosage and medication type (the actual column names are defined in the following four parameters).
<code>ID.colname</code>	A <i>string</i> , the name of the column in data containing the unique patient ID; must be present.
<code>event.date.colname</code>	A <i>string</i> , the name of the column in data containing the start date of the event (in the format given in the <code>date.format</code> parameter); must be present.
<code>event.duration.colname</code>	A <i>string</i> , the name of the column in data containing the event duration (in days); must be present.
<code>event.daily.dose.colname</code>	A <i>string</i> , the name of the column in data containing the prescribed daily dose, or NA if not defined.
<code>medication.class.colname</code>	A <i>string</i> , the name of the column in data containing the medication type, or NA if not defined.
<code>carry.only.for.same.medication</code>	<i>Logical</i> , if TRUE, the carry-over applies only across medication of the same type.
<code>consider.dosage.change</code>	<i>Logical</i> , if TRUE, the carry-over is adjusted to also reflect changes in dosage.
<code>followup.window.start</code>	If a Date object, it represents the actual start date of the follow-up window; if a <i>string</i> it is the name of the column in data containing the start date of the follow-up window either as the numbers of <code>followup.window.start.unit</code> units after the first event (the column must be of type <code>numeric</code>) or as actual dates (in which case the column must be of type <code>Date</code>); if a <i>number</i> it is the number of time units defined in the <code>followup.window.start.unit</code> parameter after the begin of the participant's first event; or NA if not defined.
<code>followup.window.start.unit</code>	can be either <i>"days"</i> , <i>"weeks"</i> , <i>"months"</i> or <i>"years"</i> , and represents the time units that <code>followup.window.start</code> refers to (when a number), or NA if not defined.
<code>followup.window.duration</code>	either a <i>number</i> representing the duration of the follow-up window in the time units given in <code>followup.window.start.unit</code> , or a <i>string</i> giving the column containing these numbers. Should represent a period for which relevant

	medication events are recorded accurately (e.g. not extend after end of relevant treatment, loss-to-follow-up or change to a health care provider not covered by the database).
<code>followup.window.duration.unit</code>	can be either <i>"days"</i> , <i>"weeks"</i> , <i>"months"</i> or <i>"years"</i> , and represents the time units that <code>followup.window.duration</code> refers to, or NA if not defined.
<code>observation.window.start</code> , <code>observation.window.start.unit</code> , <code>observation.window.duration</code> , <code>observation.w</code>	the definition of the observation window (see the follow-up window parameters above for details).
<code>date.format</code>	A <i>string</i> giving the format of the dates used in the data and the other parameters; see the format parameters of the <code>as.Date</code> function for details (NB, this concerns only the dates given as strings and not as Date objects).
<code>summary</code>	Metadata as a <i>string</i> , briefly describing this CMA.
<code>event.interval.colname</code>	A <i>string</i> , the name of a newly-created column storing the number of days between the start of the current event and the start of the next one; the default value <i>"event.interval"</i> should be changed only if there is a naming conflict with a pre-existing <i>"event.interval"</i> column in <code>event.info</code> .
<code>gap.days.colname</code>	A <i>string</i> , the name of a newly-created column storing the number of days when medication was not available (i.e., the <i>"gap days"</i>); the default value <i>"gap.days"</i> should be changed only if there is a naming conflict with a pre-existing <i>"gap.days"</i> column in <code>event.info</code> .
<code>force.NA.CMA.for.failed.patients</code>	<i>Logical</i> describing how the patients for which the CMA estimation fails are treated: if TRUE they are returned with an NA CMA estimate, while for FALSE they are omitted.
<code>parallel.backend</code>	Can be <i>"none"</i> (the default) for single-threaded execution, <i>"multicore"</i> (using <code>mclapply</code> in package <code>parallel</code>) for multicore processing (NB. not currently implemented on MS Windows and automatically falls back on <i>"snow"</i> on this platform), or <i>"snow"</i> , <i>"snow(SOCK)"</i> (equivalent to <i>"snow"</i>), <i>"snow(MPI)"</i> or <i>"snow(NWS)"</i> specifying various types of SNOW clusters (can be on the local machine or more complex setups – please see the documentation of package <code>snow</code> for details; the last two require packages <code>Rmpi</code> and <code>nws</code> , respectively, not automatically installed with <code>AdhereR</code>).
<code>parallel.threads</code>	Can be <i>"auto"</i> (for <code>parallel.backend == "multicore"</code> , defaults to the number of cores in the system as given by <code>options("cores")</code>), while for <code>parallel.backend == "snow"</code> , defaults to 2), a strictly positive integer specifying the number of parallel threads, or a more complex specification of the SNOW cluster nodes for <code>parallel.backend == "snow"</code> (see the documentation of package <code>snow</code> for details).
<code>suppress.warnings</code>	<i>Logical</i> , if TRUE don't show any warnings.
<code>arguments.that.should.not.be.defined</code>	a <i>list</i> of argument names and pre-defined values for which a warning should be thrown if passed to the function.

... other possible parameters

Details

CMA8 is similar to CMA6 in that it assumes that, within the observation window, the medication is used as prescribed and new medication is "banked" until needed (oversupply from previous events is used first, followed new medication supply). Unlike CMA6 it accounts for carry-over from before the window - but in a different way from CMA7: by adding a time lag at the start of the observation window equal to the duration of carry-over from before. It is designed for situations when an event with a hypothesized causal effect on adherence happens at the start of the observation window (e.g. enrolment in an intervention study); in this case, it may be that the existing supply is not part of the relationship under study (e.g. it delays the actual start of the study for that participant) and needs to be excluded by shortening the time interval examined. The end of the observation window remains the same. Thus, CMA8 computes days of theoretical use by extracting the total number of gap days from the total time interval between the lagged start and the end of the observation window, accounting for carry over for all medication events within the observation window. All medication events in the follow up window before observation window are considered for carry-over calculation. Thus, as CMA7, it accounts for timing within the observation window, as well as before (different adjustment than CMA7), and excludes the remaining supply at the end of the observation window.

The formula is

$$(number\ of\ days\ of\ theoretical\ use) / (lagged\ start\ to\ end\ of\ observation\ window)$$

Observations:

- the `carry.only.for.same.medication` parameter controls the transmission of carry-over across medication changes, producing a "standard" CMA8 (default value is FALSE), and an "alternative" CMA8b, respectively;
- the `consider.dosage.change` parameter controls if dosage changes are taken into account, i.e. if set as TRUE and a new medication event has a different daily dosage recommendation, carry-over is recomputed assuming medication use according to the new prescribed dosage (default value is FALSE).

Value

An S3 object of class CMA8 (derived from CMA0) with the following fields:

- `data` The actual event data, as given by the `data` parameter.
- `ID.colname` the name of the column in `data` containing the unique patient ID, as given by the `ID.colname` parameter.
- `event.date.colname` the name of the column in `data` containing the start date of the event (in the format given in the `date.format` parameter), as given by the `event.date.colname` parameter.
- `event.duration.colname` the name of the column in `data` containing the event duration (in days), as given by the `event.duration.colname` parameter.
- `event.daily.dose.colname` the name of the column in `data` containing the prescribed daily dose, as given by the `event.daily.dose.colname` parameter.

- `medication.class.colname` the name of the column in data containing the classes/types/groups of medication, as given by the `medication.class.colname` parameter.
- `carry.only.for.same.medication` whether the carry-over applies only across medication of the same type, as given by the `carry.only.for.same.medication` parameter.
- `consider.dosage.change` whether the carry-over is adjusted to reflect changes in dosage, as given by the `consider.dosage.change` parameter.
- `followup.window.start` the beginning of the follow-up window, as given by the `followup.window.start` parameter.
- `followup.window.start.unit` the time unit of the `followup.window.start`, as given by the `followup.window.start.unit` parameter.
- `followup.window.duration` the duration of the follow-up window, as given by the `followup.window.duration` parameter.
- `followup.window.duration.unit` the time unit of the `followup.window.duration`, as given by the `followup.window.duration.unit` parameter.
- `observation.window.start` the beginning of the observation window, as given by the `observation.window.start` parameter.
- `observation.window.start.unit` the time unit of the `observation.window.start`, as given by the `observation.window.start.unit` parameter.
- `observation.window.duration` the duration of the observation window, as given by the `observation.window.duration` parameter.
- `observation.window.duration.unit` the time unit of the `observation.window.duration`, as given by the `observation.window.duration.unit` parameter.
- `date.format` the format of the dates, as given by the `date.format` parameter.
- `summary` the metadata, as given by the `summary` parameter.
- `event.info` the data.frame containing the event info (irrelevant for most users; see `compute.event.int.gaps` for details).
- `CMA` the data.frame containing the actual CMA estimates for each participant (the `ID.colname` column).

See Also

CMA8 1 to 8 are defined in:

Vollmer, W. M., Xu, M., Feldstein, A., Smith, D., Waterbury, A., & Rand, C. (2012). Comparison of pharmacy-based measures of medication adherence. *BMC Health Services Research*, **12**, 155. <http://doi.org/10.1186/1472-6963-12-155>.

Examples

```
cma8 <- CMA8(data=med.events,
             ID.colname="PATIENT_ID",
             event.date.colname="DATE",
             event.duration.colname="DURATION",
             event.daily.dose.colname="PERDAY",
             medication.class.colname="CATEGORY",
             carry.only.for.same.medication=FALSE,
```

```

consider.dosage.change=FALSE,
followup.window.start=30,
observation.window.start=30,
observation.window.duration=365,
date.format="%m/%d/%Y"
);

```

CMA9

CMA9 constructor.

Description

Constructs a CMA (continuous multiple-interval measures of medication availability/gaps) type 9 object.

Usage

```

CMA9(data = NULL, ID.colname = NA, event.date.colname = NA,
      event.duration.colname = NA, event.daily.dose.colname = NA,
      medication.class.colname = NA,
      carry.only.for.same.medicament = FALSE,
      consider.dosage.change = FALSE, followup.window.start = 0,
      followup.window.start.unit = c("days", "weeks", "months", "years")[1],
      followup.window.duration = 365 * 2,
      followup.window.duration.unit = c("days", "weeks", "months",
      "years")[1], observation.window.start = 0,
      observation.window.start.unit = c("days", "weeks", "months",
      "years")[1], observation.window.duration = 365 * 2,
      observation.window.duration.unit = c("days", "weeks", "months",
      "years")[1], date.format = "%m/%d/%Y", summary = NA,
      event.interval.colname = "event.interval",
      gap.days.colname = "gap.days",
      force.NA.CMA.for.failed.patients = TRUE, parallel.backend = c("none",
      "multicore", "snow", "snow(SOCK)", "snow(MPI)", "snow(NWS)")[1],
      parallel.threads = "auto", suppress.warnings = FALSE,
      arguments.that.should.not.be.defined = c(carryover.within.obs.window =
      TRUE, carryover.into.obs.window = TRUE), ...)

```

Arguments

data	A data frame containing the events used to compute the CMA. Must contain, at a minimum, the patient unique ID, the event date and duration, and might also contain the daily dosage and medication type (the actual column names are defined in the following four parameters).
ID.colname	A <i>string</i> , the name of the column in data containing the unique patient ID; must be present.

<code>event.date.colname</code>	A <i>string</i> , the name of the column in data containing the start date of the event (in the format given in the <code>date.format</code> parameter); must be present.
<code>event.duration.colname</code>	A <i>string</i> , the name of the column in data containing the event duration (in days); must be present.
<code>event.daily.dose.colname</code>	A <i>string</i> , the name of the column in data containing the prescribed daily dose, or NA if not defined.
<code>medication.class.colname</code>	A <i>string</i> , the name of the column in data containing the medication type, or NA if not defined.
<code>carry.only.for.same.medication</code>	<i>Logical</i> , if TRUE, the carry-over applies only across medication of the same type.
<code>consider.dosage.change</code>	<i>Logical</i> , if TRUE, the carry-over is adjusted to also reflect changes in dosage.
<code>followup.window.start</code>	If a Date object, it represents the actual start date of the follow-up window; if a <i>string</i> it is the name of the column in data containing the start date of the follow-up window either as the numbers of <code>followup.window.start.unit</code> units after the first event (the column must be of type <code>numeric</code>) or as actual dates (in which case the column must be of type <code>Date</code>); if a <i>number</i> it is the number of time units defined in the <code>followup.window.start.unit</code> parameter after the begin of the participant's first event; or NA if not defined.
<code>followup.window.start.unit</code>	can be either " <i>days</i> ", " <i>weeks</i> ", " <i>months</i> " or " <i>years</i> ", and represents the time units that <code>followup.window.start</code> refers to (when a number), or NA if not defined.
<code>followup.window.duration</code>	either a <i>number</i> representing the duration of the follow-up window in the time units given in <code>followup.window.duration.unit</code> , or a <i>string</i> giving the column containing these numbers. Should represent a period for which relevant medication events are recorded accurately (e.g. not extend after end of relevant treatment, loss-to-follow-up or change to a health care provider not covered by the database).
<code>followup.window.duration.unit</code>	can be either " <i>days</i> ", " <i>weeks</i> ", " <i>months</i> " or " <i>years</i> ", and represents the time units that <code>followup.window.duration</code> refers to, or NA if not defined.
<code>observation.window.start</code> , <code>observation.window.start.unit</code> , <code>observation.window.duration</code> , <code>observation.w</code>	the definition of the observation window (see the follow-up window parameters above for details).
<code>date.format</code>	A <i>string</i> giving the format of the dates used in the data and the other parameters; see the format parameters of the <code>as.Date</code> function for details (NB, this concerns only the dates given as strings and not as Date objects).
<code>summary</code>	Metadata as a <i>string</i> , briefly describing this CMA.
<code>event.interval.colname</code>	A <i>string</i> , the name of a newly-created column storing the number of days between the start of the current event and the start of the next one; the default

value "event.interval" should be changed only if there is a naming conflict with a pre-existing "event.interval" column in `event.info`.

`gap.days.colname`

A *string*, the name of a newly-created column storing the number of days when medication was not available (i.e., the "gap days"); the default value "gap.days" should be changed only if there is a naming conflict with a pre-existing "gap.days" column in `event.info`.

`force.NA.CMA.for.failed.patients`

Logical describing how the patients for which the CMA estimation fails are treated: if TRUE they are returned with an NA CMA estimate, while for FALSE they are omitted.

`parallel.backend`

Can be "none" (the default) for single-threaded execution, "multicore" (using `mclapply` in package `parallel`) for multicore processing (NB. not currently implemented on MS Windows and automatically falls back on "snow" on this platform), or "snow", "snow(SOCK)" (equivalent to "snow"), "snow(MPI)" or "snow(NWS)" specifying various types of SNOW clusters (can be on the local machine or more complex setups – please see the documentation of package `snow` for details; the last two require packages `Rmpi` and `nws`, respectively, not automatically installed with `AdhereR`).

`parallel.threads`

Can be "auto" (for `parallel.backend == "multicore"`, defaults to the number of cores in the system as given by `options("cores")`), while for `parallel.backend == "snow"`, defaults to 2), a strictly positive integer specifying the number of parallel threads, or a more complex specification of the SNOW cluster nodes for `parallel.backend == "snow"` (see the documentation of package `snow` for details).

`suppress.warnings`

Logical, if TRUE don't show any warnings.

`arguments.that.should.not.be.defined`

a *list* of argument names and pre-defined values for which a warning should be thrown if passed to the function.

...

other possible parameters

Details

CMA9 is similar to CMA7 and CMA8 in that it accounts for carry-over within and before the observation window assuming that new medication is "banked" until needed (oversupply from previous events is used first, followed new medication supply). Yet, unlike these previous CMAs, it does not assume the medication is used as prescribed; in longitudinal studies with multiple CMA measures, this assumption may introduce additional variation in CMA estimates depending on when the observation window starts in relation to the previous medication event. A shorter time distance from the previous event (and longer to the first event in the observation window) results in higher values even if the number of gap days is the same, and it may also be that the patient has had a similar use pattern for that time interval, rather than perfect adherence followed by no medication use. CMA9 applies a different adjustment: it computes a ratio of days supply over each interval between two prescriptions and considers this applies for each day of that interval, up to 100% (moving oversupply to the

next event interval). All medication events in the follow up window before observation window are considered for carry-over calculation. The last interval ends at the end of the follow-up window. Thus, it accounts for timing within the observation window, as well as before (but differently from CMA7 and CMA8), and excludes the remaining supply at the end of the observation window, if any.

The formula is

(number of days in the observation window, each weighted by the ratio of days supply applicable to the event interval) / (

Observations:

- the `carry.only.for.same.medication` parameter controls the transmission of carry-over across medication changes, producing a "standard" CMA7 (default value is FALSE), and an "alternative" CMA7b, respectively;
- the `consider.dosage.change` parameter controls if dosage changes are taken into account, i.e. if set as TRUE and a new medication event has a different daily dosage recommendation, carry-over is recomputed assuming medication use according to the new prescribed dosage (default value is FALSE).

Value

An S3 object of class CMA9 (derived from CMA0) with the following fields:

- `data` The actual event data, as given by the `data` parameter.
- `ID.colname` the name of the column in `data` containing the unique patient ID, as given by the `ID.colname` parameter.
- `event.date.colname` the name of the column in `data` containing the start date of the event (in the format given in the `date.format` parameter), as given by the `event.date.colname` parameter.
- `event.duration.colname` the name of the column in `data` containing the event duration (in days), as given by the `event.duration.colname` parameter.
- `event.daily.dose.colname` the name of the column in `data` containing the prescribed daily dose, as given by the `event.daily.dose.colname` parameter.
- `medication.class.colname` the name of the column in `data` containing the classes/types/groups of medication, as given by the `medication.class.colname` parameter.
- `carry.only.for.same.medication` whether the carry-over applies only across medication of the same type, as given by the `carry.only.for.same.medication` parameter.
- `consider.dosage.change` whether the carry-over is adjusted to reflect changes in dosage, as given by the `consider.dosage.change` parameter.
- `followup.window.start` the beginning of the follow-up window, as given by the `followup.window.start` parameter.
- `followup.window.start.unit` the time unit of the `followup.window.start`, as given by the `followup.window.start.unit` parameter.
- `followup.window.duration` the duration of the follow-up window, as given by the `followup.window.duration` parameter.
- `followup.window.duration.unit` the time unit of the `followup.window.duration`, as given by the `followup.window.duration.unit` parameter.

- `observation.window.start` the beginning of the observation window, as given by the `observation.window.start` parameter.
- `observation.window.start.unit` the time unit of the `observation.window.start`, as given by the `observation.window.start.unit` parameter.
- `observation.window.duration` the duration of the observation window, as given by the `observation.window.duration` parameter.
- `observation.window.duration.unit` the time unit of the `observation.window.duration`, as given by the `observation.window.duration.unit` parameter.
- `date.format` the format of the dates, as given by the `date.format` parameter.
- `summary` the metadata, as given by the `summary` parameter.
- `event.info` the data frame containing the event info (irrelevant for most users; see `compute.event.int.gaps` for details).
- `CMA` the data frame containing the actual CMA estimates for each participant (the `ID.colname` column).

Examples

```
cma9 <- CMA9(data=med.events,
             ID.colname="PATIENT_ID",
             event.date.colname="DATE",
             event.duration.colname="DURATION",
             event.daily.dose.colname="PERDAY",
             medication.class.colname="CATEGORY",
             carry.only.for.same.medicament=FALSE,
             consider.dosage.change=FALSE,
             followup.window.start=30,
             observation.window.start=30,
             observation.window.duration=365,
             date.format="%m/%d/%Y"
             );
```

CMA_per_episode

CMA_per_episode constructor.

Description

Applies a given CMA to each treatment episode and constructs a `CMA_per_episode` object.

Usage

```
CMA_per_episode(CMA.to.apply, data, ID.colname = NA,
               event.date.colname = NA, event.duration.colname = NA,
               event.daily.dose.colname = NA, medication.class.colname = NA,
               carry.only.for.same.medicament = NA, consider.dosage.change = NA,
               medication.change.means.new.treatment.episode = TRUE,
               dosage.change.means.new.treatment.episode = FALSE,
```

```

maximum.permissible.gap = 180,
maximum.permissible.gap.unit = c("days", "weeks", "months", "years",
"percent")[1], followup.window.start = 0,
followup.window.start.unit = c("days", "weeks", "months", "years")[1],
followup.window.duration = 365 * 2,
followup.window.duration.unit = c("days", "weeks", "months",
"years")[1], observation.window.start = 0,
observation.window.start.unit = c("days", "weeks", "months",
"years")[1], observation.window.duration = 365 * 2,
observation.window.duration.unit = c("days", "weeks", "months",
"years")[1], date.format = "%m/%d/%Y",
summary = "CMA per treatment episode",
event.interval.colname = "event.interval",
gap.days.colname = "gap.days",
force.NA.CMA.for.failed.patients = TRUE, parallel.backend = c("none",
"multicore", "snow", "snow(SOCK)", "snow(MPI)", "snow(NWS)")[1],
parallel.threads = "auto", suppress.warnings = FALSE, ...)

```

Arguments

- `CMA.to.apply` A *string* giving the name of the CMA function (1 to 9) that will be computed for each treatment episode.
- `data` A *data.frame* containing the events (prescribing or dispensing) used to compute the CMA. Must contain, at a minimum, the patient unique ID, the event date and duration, and might also contain the daily dosage and medication type (the actual column names are defined in the following four parameters).
- `ID.colname` A *string*, the name of the column in *data* containing the unique patient ID; must be present.
- `event.date.colname` A *string*, the name of the column in *data* containing the start date of the event (in the format given in the `date.format` parameter); must be present.
- `event.duration.colname` A *string*, the name of the column in *data* containing the event duration (in days); must be present.
- `event.daily.dose.colname` A *string*, the name of the column in *data* containing the prescribed daily dose, or NA if not defined.
- `medication.class.colname` A *string*, the name of the column in *data* containing the medication type, or NA if not defined.
- `carry.only.for.same.medication` *Logical*, if TRUE, the carry-over applies only across medication of the same type; valid only for CMAs 5 to 9, in which case it is coupled (i.e., the same value is used for computing the treatment episodes and the CMA on each treatment episode).
- `consider.dosage.change` *Logical*, if TRUE, the carry-over is adjusted to also reflect changes in dosage; valid only for CMAs 5 to 9, in which case it is coupled (i.e., the same value

	is used for computing the treatment episodes and the CMA on each treatment episode).
medication.change.means.new.treatment.episode	<i>Logical</i> , should a change in medication automatically start a new treatment episode?
dosage.change.means.new.treatment.episode	<i>Logical</i> , should a change in dosage automatically start a new treatment episode?
maximum.permissible.gap	The <i>number</i> of units given by maximum.permissible.gap.unit representing the maximum duration of permissible gaps between treatment episodes (can also be a percent, see maximum.permissible.gap.unit for details).
maximum.permissible.gap.unit	can be either " <i>days</i> ", " <i>weeks</i> ", " <i>months</i> ", " <i>years</i> " or " <i>percent</i> ", and represents the time units that maximum.permissible.gap refers to; if <i>percent</i> , then maximum.permissible.gap is interpreted as a percent (can be greater than 100%) of the duration of the current prescription.
followup.window.start	If a Date object, it represents the actual start date of the follow-up window; if a <i>string</i> it is the name of the column in data containing the start date of the follow-up window either as the numbers of followup.window.start.unit units after the first event (the column must be of type numeric) or as actual dates (in which case the column must be of type Date); if a <i>number</i> it is the number of time units defined in the followup.window.start.unit parameter after the begin of the participant's first event; or NA if not defined.
followup.window.start.unit	can be either " <i>days</i> ", " <i>weeks</i> ", " <i>months</i> " or " <i>years</i> ", and represents the time units that followup.window.start refers to (when a number), or NA if not defined.
followup.window.duration	either a <i>number</i> representing the duration of the follow-up window in the time units given in followup.window.duration.unit, or a <i>string</i> giving the column containing these numbers. Should represent a period for which relevant medication events are recorded accurately (e.g. not extend after end of relevant treatment, loss-to-follow-up or change to a health care provider not covered by the database).
followup.window.duration.unit	can be either " <i>days</i> ", " <i>weeks</i> ", " <i>months</i> " or " <i>years</i> ", and represents the time units that followup.window.duration refers to, or NA if not defined.
observation.window.start, observation.window.start.unit, observation.window.duration, observation.w	the definition of the observation window (see the follow-up window parameters above for details).
date.format	A <i>string</i> giving the format of the dates used in the data and the other parameters; see the format parameters of the as.Date function for details (NB, this concerns only the dates given as strings and not as Date objects).
summary	Metadata as a <i>string</i> , briefly describing this CMA.
event.interval.colname	A <i>string</i> , the name of a newly-created column storing the number of days between the start of the current event and the start of the next one; the default

value "event.interval" should be changed only if there is a naming conflict with a pre-existing "event.interval" column in `event.info`.

`gap.days.colname`

A *string*, the name of a newly-created column storing the number of days when medication was not available (i.e., the "gap days"); the default value "gap.days" should be changed only if there is a naming conflict with a pre-existing "gap.days" column in `event.info`.

`force.NA.CMA.for.failed.patients`

Logical describing how the patients for which the CMA estimation fails are treated: if TRUE they are returned with an NA CMA estimate, while for FALSE they are omitted.

`parallel.backend`

Can be "none" (the default) for single-threaded execution, "multicore" (using `mclapply` in package `parallel`) for multicore processing (NB. not currently implemented on MS Windows and automatically falls back on "snow" on this platform), or "snow", "snow(SOCK)" (equivalent to "snow"), "snow(MPI)" or "snow(NWS)" specifying various types of SNOW clusters (can be on the local machine or more complex setups – please see the documentation of package `snow` for details; the last two require packages `Rmpi` and `nws`, respectively, not automatically installed with `AdhereR`).

`parallel.threads`

Can be "auto" (for `parallel.backend == "multicore"`, defaults to the number of cores in the system as given by `options("cores")`), while for `parallel.backend == "snow"`, defaults to 2), a strictly positive integer specifying the number of parallel threads, or a more complex specification of the SNOW cluster nodes for `parallel.backend == "snow"` (see the documentation of package `snow` for details).

`suppress.warnings`

Logical, if TRUE don't show any warnings.

...

other possible parameters

Details

`CMA_per_episode` first identifies the treatment episodes for the whole follow-up window (using the `compute.treatment.episodes` function), and then computes the given "simple" CMA for each treatment episode that intersects with the observation window. NB: the CMA is computed for the period of the episode that is part of the observations window; thus, if an episode starts earlier or ends later than the observation window, CMA will be computed for a section of that episode. Thus, as opposed to the "simple" CMAs 1 to 9, it returns a set of CMAs, with possibly more than one element.

It is highly similar to `CMA_sliding_window` which computes a CMA for a set of sliding windows.

Value

An S3 object of class `CMA_per_episode` with the following fields:

- `data` The actual event data, as given by the `data` parameter.

- `ID.colname` the name of the column in data containing the unique patient ID, as given by the `ID.colname` parameter.
- `event.date.colname` the name of the column in data containing the start date of the event (in the format given in the `date.format` parameter), as given by the `event.date.colname` parameter.
- `event.duration.colname` the name of the column in data containing the event duration (in days), as given by the `event.duration.colname` parameter.
- `event.daily.dose.colname` the name of the column in data containing the prescribed daily dose, as given by the `event.daily.dose.colname` parameter.
- `medication.class.colname` the name of the column in data containing the classes/types/groups of medication, as given by the `medication.class.colname` parameter.
- `carry.only.for.same.medication` whether the carry-over applies only across medication of the same type, as given by the `carry.only.for.same.medication` parameter.
- `consider.dosage.change` whether the carry-over is adjusted to reflect changes in dosage, as given by the `consider.dosage.change` parameter.
- `followup.window.start` the beginning of the follow-up window, as given by the `followup.window.start` parameter.
- `followup.window.start.unit` the time unit of the `followup.window.start`, as given by the `followup.window.start.unit` parameter.
- `followup.window.duration` the duration of the follow-up window, as given by the `followup.window.duration` parameter.
- `followup.window.duration.unit` the time unit of the `followup.window.duration`, as given by the `followup.window.duration.unit` parameter.
- `observation.window.start` the beginning of the observation window, as given by the `observation.window.start` parameter.
- `observation.window.start.unit` the time unit of the `observation.window.start`, as given by the `observation.window.start.unit` parameter.
- `observation.window.duration` the duration of the observation window, as given by the `observation.window.duration` parameter.
- `observation.window.duration.unit` the time unit of the `observation.window.duration`, as given by the `observation.window.duration.unit` parameter.
- `date.format` the format of the dates, as given by the `date.format` parameter.
- `summary` the metadata, as given by the `summary` parameter.
- `event.info` the data frame containing the event info (irrelevant for most users; see [compute.event.int.gaps](#) for details).
- `computed.CMA` the class name of the computed CMA.
- `CMA` the data frame containing the actual CMA estimates for each participant (the `ID.colname` column) and treatment episode, with columns:
 - `ID.colname` the patient ID as given by the `ID.colname` parameter.
 - `episode.ID` the unique treatment episode ID (within patients).
 - `episode.start` the treatment episode's start date (as a Date object).
 - `end.episode.gap.days` the corresponding gap days of the last event in this episode.

- `episode.duration` the treatment episode's duration in days.
- `episode.end` the treatment episode's end date (as a Date object).
- `CMA` the treatment episode's estimated CMA.

See Also

`CMA_sliding_window` is very similar, computing a "simple" CMA for each of a set of same-size sliding windows. The "simple" CMAs that can be computed comprise [CMA1](#), [CMA2](#), [CMA3](#), [CMA4](#), [CMA5](#), [CMA6](#), [CMA7](#), [CMA8](#), [CMA9](#), as well as user-defined classes derived from [CMA0](#) that have a `CMA` component giving the estimated CMA per patient as a `data.frame`.

Examples

```
## Not run:
cmaE <- CMA_per_episode(CMA="CMA1",
                        data=med.events,
                        ID.colname="PATIENT_ID",
                        event.date.colname="DATE",
                        event.duration.colname="DURATION",
                        event.daily.dose.colname="PERDAY",
                        medication.class.colname="CATEGORY",
                        carry.only.for.same.medicament=FALSE,
                        consider.dosage.change=FALSE,
                        followup.window.start=0,
                        observation.window.start=0,
                        observation.window.duration=365,
                        date.format="%m/%d/%Y"
                        );
## End(Not run)
```

`CMA_sliding_window` *CMA_sliding_window constructor.*

Description

Applies a given CMA to each sliding window and constructs a `CMA_sliding_window` object.

Usage

```
CMA_sliding_window(CMA.to.apply, data, ID.colname = NA,
                  event.date.colname = NA, event.duration.colname = NA,
                  event.daily.dose.colname = NA, medication.class.colname = NA,
                  carry.only.for.same.medicament = NA, consider.dosage.change = NA,
                  followup.window.start = 0, followup.window.start.unit = c("days",
                  "weeks", "months", "years")[1], followup.window.duration = 365 * 2,
                  followup.window.duration.unit = c("days", "weeks", "months",
                  "years")[1], observation.window.start = 0,
                  observation.window.start.unit = c("days", "weeks", "months",
```

```

"years")[1], observation.window.duration = 365 * 2,
observation.window.duration.unit = c("days", "weeks", "months",
"years")[1], sliding.window.start = 0,
sliding.window.start.unit = c("days", "weeks", "months", "years")[1],
sliding.window.duration = 90,
sliding.window.duration.unit = c("days", "weeks", "months",
"years")[1], sliding.window.step.duration = 30,
sliding.window.step.unit = c("days", "weeks", "months", "years")[1],
sliding.window.no.steps = NA, date.format = "%m/%d/%Y",
summary = "CMA per sliding window",
event.interval.colname = "event.interval",
gap.days.colname = "gap.days",
force.NA.CMA.for.failed.patients = TRUE, parallel.backend = c("none",
"multicore", "snow", "snow(SOCK)", "snow(MPI)", "snow(NWS)")[1],
parallel.threads = "auto", suppress.warnings = FALSE, ...)

```

Arguments

CMA.to.apply	A <i>string</i> giving the name of the CMA function (1 to 9) that will be computed for each treatment episode.
data	A <i>data.frame</i> containing the events used to compute the CMA. Must contain, at a minimum, the patient unique ID, the event date and duration, and might also contain the daily dosage and medication type (the actual column names are defined in the following four parameters).
ID.colname	A <i>string</i> , the name of the column in data containing the unique patient ID; must be present.
event.date.colname	A <i>string</i> , the name of the column in data containing the start date of the event (in the format given in the <code>date.format</code> parameter); must be present.
event.duration.colname	A <i>string</i> , the name of the column in data containing the event duration (in days); must be present.
event.daily.dose.colname	A <i>string</i> , the name of the column in data containing the prescribed daily dose, or NA if not defined.
medication.class.colname	A <i>string</i> , the name of the column in data containing the medication type, or NA if not defined.
carry.only.for.same.medication	<i>Logical</i> , if TRUE, the carry-over applies only across medication of the same type.
consider.dosage.change	<i>Logical</i> , if TRUE, the carry-over is adjusted to also reflect changes in dosage.
followup.window.start	If a <i>Date</i> object, it represents the actual start date of the follow-up window; if a <i>string</i> it is the name of the column in data containing the start date of the follow-up window either as the numbers of <code>followup.window.start.unit</code> units after the first event (the column must be of type <code>numeric</code>) or as actual dates (in which

- case the column must be of type Date); if a *number* it is the number of time units defined in the `followup.window.start.unit` parameter after the begin of the participant's first event; or NA if not defined.
- `followup.window.start.unit`
can be either *"days"*, *"weeks"*, *"months"* or *"years"*, and represents the time units that `followup.window.start` refers to (when a number), or NA if not defined.
- `followup.window.duration`
either a *number* representing the duration of the follow-up window in the time units given in `followup.window.duration.unit`, or a *string* giving the column containing these numbers. Should represent a period for which relevant medication events are recorded accurately (e.g. not extend after end of relevant treatment, loss-to-follow-up or change to a health care provider not covered by the database).
- `followup.window.duration.unit`
can be either *"days"*, *"weeks"*, *"months"* or *"years"*, and represents the time units that `followup.window.duration` refers to, or NA if not defined.
- `observation.window.start`, `observation.window.start.unit`, `observation.window.duration`, `observation.window.duration.unit`
the definition of the observation window (see the follow-up window parameters above for details).
- `sliding.window.start`, `sliding.window.start.unit`, `sliding.window.duration`, `sliding.window.duration.unit`
the definition of the first sliding window (see the follow-up window parameters above for details).
- `sliding.window.step.duration`, `sliding.window.step.unit`
if not missing (NA), these give the step (or "jump") to the right of the sliding window in time units.
- `sliding.window.no.steps`
a *integer* specifying the desired number of sliding windows to cover the observation window (if possible); trumps `sliding.window.step.duration` and `sliding.window.step.unit`.
- `date.format` A *string* giving the format of the dates used in the data and the other parameters; see the format parameters of the `as.Date` function for details (NB, this concerns only the dates given as strings and not as Date objects).
- `summary` Metadata as a *string*, briefly describing this CMA.
- `event.interval.colname`
A *string*, the name of a newly-created column storing the number of days between the start of the current event and the start of the next one; the default value "event.interval" should be changed only if there is a naming conflict with a pre-existing "event.interval" column in `event.info`.
- `gap.days.colname`
A *string*, the name of a newly-created column storing the number of days when medication was not available (i.e., the "gap days"); the default value "gap.days" should be changed only if there is a naming conflict with a pre-existing "gap.days" column in `event.info`.
- `force.NA.CMA.for.failed.patients`
Logical describing how the patients for which the CMA estimation fails are treated: if TRUE they are returned with an NA CMA estimate, while for FALSE they are omitted.

`parallel.backend`

Can be "none" (the default) for single-threaded execution, "multicore" (using `mclapply` in package `parallel`) for multicore processing (NB. not currently implemented on MS Windows and automatically falls back on "snow" on this platform), or "snow", "snow(SOCK)" (equivalent to "snow"), "snow(MPI)" or "snow(NWS)" specifying various types of SNOW clusters (can be on the local machine or more complex setups – please see the documentation of package `snow` for details; the last two require packages `Rmpi` and `nws`, respectively, not automatically installed with `AdhereR`).

`parallel.threads`

Can be "auto" (for `parallel.backend == "multicore"`, defaults to the number of cores in the system as given by `options("cores")`), while for `parallel.backend == "snow"`, defaults to 2), a strictly positive integer specifying the number of parallel threads, or a more complex specification of the SNOW cluster nodes for `parallel.backend == "snow"` (see the documentation of package `snow` for details).

`suppress.warnings`

Logical, if TRUE don't show any warnings.

...

other possible parameters

Details

`CMA_sliding_window` first computes a set of fixed-size (possibly partly overlapping) sliding windows, each sliding to the right by a fixed timelag, and then, for each of them, it computes the given "simple" CMA. Thus, as opposed to the "simple" CMAs 1 to 9, it returns a set of CMAs, with possibly more than one element.

It is highly similar to [CMA_per_episode](#) which computes a CMA for a set of treatment episodes.

Value

An S3 object of class `CMA_sliding_window` with the following fields:

- `data` The actual event data, as given by the `data` parameter.
- `ID.colname` the name of the column in `data` containing the unique patient ID, as given by the `ID.colname` parameter.
- `event.date.colname` the name of the column in `data` containing the start date of the event (in the format given in the `date.format` parameter), as given by the `event.date.colname` parameter.
- `event.duration.colname` the name of the column in `data` containing the event duration (in days), as given by the `event.duration.colname` parameter.
- `event.daily.dose.colname` the name of the column in `data` containing the prescribed daily dose, as given by the `event.daily.dose.colname` parameter.
- `medication.class.colname` the name of the column in `data` containing the classes/types/groups of medication, as given by the `medication.class.colname` parameter.
- `carry.only.for.same.medication` whether the carry-over applies only across medication of the same type, as given by the `carry.only.for.same.medication` parameter.

- `consider.dosage.change` whether the carry-over is adjusted to reflect changes in dosage, as given by the `consider.dosage.change` parameter.
- `followup.window.start` the beginning of the follow-up window, as given by the `followup.window.start` parameter.
- `followup.window.start.unit` the time unit of the `followup.window.start`, as given by the `followup.window.start.unit` parameter.
- `followup.window.duration` the duration of the follow-up window, as given by the `followup.window.duration` parameter.
- `followup.window.duration.unit` the time unit of the `followup.window.duration`, as given by the `followup.window.duration.unit` parameter.
- `observation.window.start` the beginning of the observation window, as given by the `observation.window.start` parameter.
- `observation.window.start.unit` the time unit of the `observation.window.start`, as given by the `observation.window.start.unit` parameter.
- `observation.window.duration` the duration of the observation window, as given by the `observation.window.duration` parameter.
- `observation.window.duration.unit` the time unit of the `observation.window.duration`, as given by the `observation.window.duration.unit` parameter.
- `date.format` the format of the dates, as given by the `date.format` parameter.
- `summary` the metadata, as given by the `summary` parameter.
- `event.info` the data.frame containing the event info (irrelevant for most users; see [compute.event.int.gaps](#) for details).
- `computed.CMA` the class name of the computed CMA.
- `CMA` the data.frame containing the actual CMA estimates for each participant (the `ID.colname` column) and sliding window, with columns:
 - `ID.colname` the patient ID as given by the `ID.colname` parameter.
 - `window.ID` the unique window ID (within patients).
 - `window.start` the window's start date (as a Date object).
 - `window.end` the window's end date (as a Date object).
 - `CMA` the window's estimated CMA.

See Also

[CMA_per_episode](#) is very similar, computing a "simple" CMA for each of the treatment episodes. The "simple" CMAs that can be computed comprise [CMA1](#), [CMA2](#), [CMA3](#), [CMA4](#), [CMA5](#), [CMA6](#), [CMA7](#), [CMA8](#), [CMA9](#), as well as user-defined classes derived from [CMA0](#) that have a CMA component giving the estimated CMA per patient as a data.frame.

Examples

```
## Not run:
cmaW <- CMA_sliding_window(CMA="CMA1",
                           data=med.events,
                           ID.colname="PATIENT_ID",
```

```

event.date.colname="DATE",
event.duration.colname="DURATION",
event.daily.dose.colname="PERDAY",
medication.class.colname="CATEGORY",
carry.only.for.same.medications=FALSE,
consider.dosage.change=FALSE,
followup.window.start=0,
observation.window.start=0,
observation.window.duration=365,
sliding.window.start=0,
sliding.window.start.unit="days",
sliding.window.duration=90,
sliding.window.duration.unit="days",
sliding.window.step.duration=7,
sliding.window.step.unit="days",
sliding.window.no.steps=NA,
date.format="%m/%d/%Y"
);
## End(Not run)

```

```
compute.event.int.gaps
```

Gap Days and Event (prescribing or dispensing) Intervals.

Description

For a given event (prescribing or dispensing) database, compute the gap days and event intervals in various scenarios.

Usage

```

compute.event.int.gaps(data, ID.colname = NA, event.date.colname = NA,
  event.duration.colname = NA, event.daily.dose.colname = NA,
  medication.class.colname = NA,
  event.interval.colname = "event.interval",
  gap.days.colname = "gap.days", carryover.within.obs.window = FALSE,
  carryover.into.obs.window = FALSE,
  carry.only.for.same.medications = FALSE,
  consider.dosage.change = FALSE, followup.window.start = 0,
  followup.window.start.unit = c("days", "weeks", "months", "years")[1],
  followup.window.duration = 365 * 2,
  followup.window.duration.unit = c("days", "weeks", "months",
  "years")[1], observation.window.start = 0,
  observation.window.start.unit = c("days", "weeks", "months",
  "years")[1], observation.window.duration = 365 * 2,
  observation.window.duration.unit = c("days", "weeks", "months",
  "years")[1], date.format = "%m/%d/%Y",
  keep.window.start.end.dates = FALSE,

```

```

remove.events.outside.followup.window = TRUE,
keep.event.interval.for.all.events = FALSE,
parallel.backend = c("none", "multicore", "snow", "snow(SOCK)",
"snow(MPI)", "snow(NWS)")[1], parallel.threads = "auto",
suppress.warnings = FALSE, return.data.table = FALSE, ...)

```

Arguments

data	A data.frame containing the events used to compute the CMA. Must contain, at a minimum, the patient unique ID, the event date and duration, and might also contain the daily dosage and medication type (the actual column names are defined in the following four parameters); the CMA constructors call this parameter data.
ID.colname	A <i>string</i> , the name of the column in data containing the unique patient ID; must be present.
event.date.colname	A <i>string</i> , the name of the column in data containing the start date of the event (in the format given in the date.format parameter); must be present.
event.duration.colname	A <i>string</i> , the name of the column in data containing the event duration (in days); must be present.
event.daily.dose.colname	A <i>string</i> , the name of the column in data containing the prescribed daily dose, or NA if not defined.
medication.class.colname	A <i>string</i> , the name of the column in data containing the classes/types/groups of medication, or NA if not defined.
event.interval.colname	A <i>string</i> , the name of a newly-created column storing the number of days between the start of the current event and the start of the next one; the default value "event.interval" should be changed only if there is a naming conflict with a pre-existing "event.interval" column in event.info.
gap.days.colname	A <i>string</i> , the name of a newly-created column storing the number of days when medication was not available (i.e., the "gap days"); the default value "gap.days" should be changed only if there is a naming conflict with a pre-existing "gap.days" column in event.info.
carryover.within.obs.window	<i>Logical</i> , if TRUE consider the carry-over within the observation window, or NA if not defined.
carryover.into.obs.window	<i>Logical</i> , if TRUE consider the carry-over from before the starting date of the observation window, or NA if not defined.
carry.only.for.same.medication	<i>Logical</i> , if TRUE the carry-over applies only across medication of the same type, or NA if not defined.

consider.dosage.change

Logical, if TRUE the carry-over is adjusted to reflect changes in dosage, or NA if not defined.

followup.window.start

If a *Date* object, it represents the actual start date of the follow-up window; if a *string* it is the name of the column in data containing the start date of the follow-up window either as the numbers of `followup.window.start.unit` units after the first event (the column must be of type `numeric`) or as actual dates (in which case the column must be of type `Date`); if a *number* it is the number of time units defined in the `followup.window.start.unit` parameter after the begin of the participant's first event.

followup.window.start.unit

can be either *"days"*, *"weeks"*, *"months"* or *"years"*, and represents the time units that `followup.window.start` refers to (when a number), or NA if not defined.

followup.window.duration

either a *number* representing the duration of the follow-up window in the time units given in `followup.window.duration.unit`, or a *string* giving the column containing these numbers. Should represent a period for which relevant medication events are recorded accurately (e.g. not extend after end of relevant treatment, loss-to-follow-up or change to a health care provider not covered by the database).

followup.window.duration.unit

can be either *"days"*, *"weeks"*, *"months"* or *"years"*, and represents the time units that `followup.window.duration` refers to, or NA if not defined.

observation.window.start, observation.window.start.unit, observation.window.duration, observation.w

the definition of the observation window (see the follow-up window parameters above for details).

date.format

A *string* giving the format of the dates used in the data and the other parameters; see the format parameters of the `as.Date` function for details (NB, this concerns only the dates given as strings and not as `Date` objects).

keep.window.start.end.dates

Logical, should the computed start and end dates of the windows be kept?

remove.events.outside.followup.window

Logical, should the events that fall outside the follow-up window be removed from the results?

keep.event.interval.for.all.events

Logical, should the computed event intervals be kept for all events, or NA'ed for those outside the OW?

parallel.backend

Can be *"none"* (the default) for single-threaded execution, *"multicore"* (using `mclapply` in package `parallel`) for multicore processing (NB. not currently implemented on MS Windows and automatically falls back on *"snow"* on this platform), or *"snow"*, *"snow(SOCK)"* (equivalent to *"snow"*), *"snow(MPI)"* or *"snow(NWS)"* specifying various types of SNOW clusters (can be on the local machine or more complex setups – please see the documentation of package `snow` for details; the last two require packages `Rmpi` and `nws`, respectively, not automatically installed with `AdhereR`).

parallel.threads	Can be "auto" (for parallel.backend == "multicore", defaults to the number of cores in the system as given by options("cores"), while for parallel.backend == "snow", defaults to 2), a strictly positive integer specifying the number of parallel threads, or a more complex specification of the SNOW cluster nodes for parallel.backend == "snow" (see the documentation of package snow for details).
suppress.warnings	<i>Logical</i> , if TRUE don't show any warnings.
return.data.table	<i>Logical</i> , if TRUE return a data.table object, otherwise a data.frame.
...	extra arguments.

Details

This should in general not be called directly by the user, but is provided as a basis for the extension to new CMAs.

Value

A data.frame or data.table extending the event.info parameter with:

- event.interval Or any other name given in event.interval.colname, containing the number of days between the start of the current event and the start of the next one.
- gap.days Or any other name given in gap.days.colname, containing the number of days when medication was not available for the current event (i.e., the "gap days").
- .FU.START.DATE, .FU.END.DATE if kept, the actual start and end dates of the follow-up window (after adjustments due to the various parameters).
- .OBS.START.DATE, .OBS.END.DATE if kept, the actual start and end dates of the observation window (after adjustments due to the various parameters).
- .EVENT.STARTS.BEFORE.OBS.WINDOW if kept, TRUE if the current event starts before the start of the observation window.
- .TDIFF1, .TDIFF2 if kept, various auxiliary time differences (in days).
- .EVENT.STARTS.AFTER.OBS.WINDOW if kept, TRUE if the current event starts after the end of the observation window.
- .CARRY.OVER.FROM.BEFORE if kept, the carry-over (if any) from the previous events.
- .EVENT.WITHIN.FU.WINDOW if kept, TRUE if the current event is within the follow-up window.

```
compute.treatment.episodes
```

Compute Treatment Episodes.

Description

For a given event (prescribing or dispensing) database, compute the treatment episodes for each patient in various scenarios.

Usage

```
compute.treatment.episodes(data, ID.colname = NA,
  event.date.colname = NA, event.duration.colname = NA,
  event.daily.dose.colname = NA, medication.class.colname = NA,
  carryover.within.obs.window = TRUE,
  carry.only.for.same.medicament = TRUE, consider.dosage.change = TRUE,
  medication.change.means.new.treatment.episode = TRUE,
  dosage.change.means.new.treatment.episode = FALSE,
  maximum.permissible.gap = 90,
  maximum.permissible.gap.unit = c("days", "weeks", "months", "years",
  "percent")[1], followup.window.start = 0,
  followup.window.start.unit = c("days", "weeks", "months", "years")[1],
  followup.window.duration = 365 * 2,
  followup.window.duration.unit = c("days", "weeks", "months",
  "years")[1], event.interval.colname = "event.interval",
  gap.days.colname = "gap.days", date.format = "%m/%d/%Y",
  parallel.backend = c("none", "multicore", "snow", "snow(SOCK)",
  "snow(MPI)", "snow(NWS)")[1], parallel.threads = "auto",
  suppress.warnings = FALSE, return.data.table = FALSE, ...)
```

Arguments

<code>data</code>	A data frame containing the events used to compute the CMA. Must contain, at a minimum, the patient unique ID, the event date and duration, and might also contain the daily dosage and medication type (the actual column names are defined in the following four parameters); the CMA constructors call this parameter <code>data</code> .
<code>ID.colname</code>	A <i>string</i> , the name of the column in <code>data</code> containing the unique patient ID, or NA if not defined.
<code>event.date.colname</code>	A <i>string</i> , the name of the column in <code>data</code> containing the start date of the event (in the format given in the <code>date.format</code> parameter), or NA if not defined.
<code>event.duration.colname</code>	A <i>string</i> , the name of the column in <code>data</code> containing the event duration (in days), or NA if not defined.

`event.daily.dose.colname`
 A *string*, the name of the column in data containing the prescribed daily dose, or NA if not defined.

`medication.class.colname`
 A *string*, the name of the column in data containing the classes/types/groups of medication, or NA if not defined.

`carryover.within.obs.window`
Logical, if TRUE consider the carry-over within the observation window, or NA if not defined.

`carry.only.for.same.medication`
Logical, if TRUE the carry-over applies only across medication of the same type, or NA if not defined.

`consider.dosage.change`
Logical, if TRUE the carry-over is adjusted to reflect changes in dosage, or NA if not defined.

`medication.change.means.new.treatment.episode`
Logical, should a change in medication automatically start a new treatment episode?

`dosage.change.means.new.treatment.episode`
Logical, should a change in dosage automatically start a new treatment episode?

`maximum.permissible.gap`
 The *number* of units given by `maximum.permissible.gap.unit` representing the maximum duration of permissible gaps between treatment episodes (can also be a percent, see `maximum.permissible.gap.unit` for details).

`maximum.permissible.gap.unit`
 can be either "*days*", "*weeks*", "*months*", "*years*" or "*percent*", and represents the time units that `maximum.permissible.gap` refers to; if *percent*, then `maximum.permissible.gap` is interpreted as a percent (can be greater than 100%) of the duration of the current prescription.

`followup.window.start`
 If a Date object it is the actual start date of the follow-up window; if a *string* it is the name of the column in data containing the start date of the follow-up window; if a *number* it is the number of time units defined in the `followup.window.start.unit` parameter after the begin of the participant's first event; or NA if not defined.

`followup.window.start.unit`
 can be either "*days*", "*weeks*", "*months*" or "*years*", and represents the time units that `followup.window.start` refers to (when a number), or NA if not defined.

`followup.window.duration`
 a *number* representing the duration of the follow-up window in the time units given in `followup.window.duration.unit`, or NA if not defined.

`followup.window.duration.unit`
 can be either "*days*", "*weeks*", "*months*" or "*years*", and represents the time units that `followup.window.duration` refers to, or NA if not defined.

`event.interval.colname`
 A *string*, the name of a newly-created column storing the number of days between the start of the current event and the start of the next one; the default value "event.interval" should be changed only if there is a naming conflict with a pre-existing "event.interval" column in `event.info`.

gap.days.colname	A <i>string</i> , the name of a newly-created column storing the number of days when medication was not available (i.e., the "gap days"); the default value "gap.days" should be changed only if there is a naming conflict with a pre-existing "gap.days" column in <code>event.info</code> .
date.format	A <i>string</i> giving the format of the dates used in the data and the other parameters; see the format parameters of the <code>as.Date</code> function for details (NB, this concerns only the dates given as strings and not as Date objects).
parallel.backend	Can be "none" (the default) for single-threaded execution, "multicore" (using <code>mclapply</code> in package <code>parallel</code>) for multicore processing (NB, not currently implemented on MS Windows and automatically falls back on "snow" on this platform), or "snow", "snow(SOCK)" (equivalent to "snow"), "snow(MPI)" or "snow(NWS)" specifying various types of SNOW clusters (can be on the local machine or more complex setups – please see the documentation of package <code>snow</code> for details; the last two require packages <code>Rmpi</code> and <code>nws</code> , respectively, not automatically installed with <code>AdhereR</code>).
parallel.threads	Can be "auto" (for <code>parallel.backend == "multicore"</code> , defaults to the number of cores in the system as given by <code>options("cores")</code>), while for <code>parallel.backend == "snow"</code> , defaults to 2), a strictly positive integer specifying the number of parallel threads, or a more complex specification of the SNOW cluster nodes for <code>parallel.backend == "snow"</code> (see the documentation of package <code>snow</code> for details).
suppress.warnings	<i>Logical</i> , if TRUE don't show any warnings.
return.data.table	<i>Logical</i> , if TRUE return a <code>data.table</code> object, otherwise a <code>data.frame</code> .
...	extra arguments.

Details

This should in general not be called directly by the user, but is provided as a basis for the extension to new CMAs.

For the last treatment episode, the gap is considered only when longer than the maximum permissible gap. Please note the following:

- episode starts at first medication event for a particular medication,
- episode ends on the day when the last supply of that medication finished or if a period longer than the permissible gap preceded the next medication event, or at the end of the FUW,
- end episode gap days represents either the number of days after the end of the treatment episode (if medication changed, or if a period longer than the permissible gap preceded the next medication event) or at the end of (and within) the episode, i.e. the number of days after the last supply finished (if no other medication event followed until the end of the FUW),
- the duration of the episode is the interval between the episode start and episode end (and may include the gap days at the end, in the latter condition described above),

- the number of gap days after the end of the episode can be computed as all values larger than the permissible gap and 0 otherwise,
- if medication change starts new episode, then previous episode ends when the last supply is finished (irrespective of the length of gap compared to a maximum permissible gap); any days before the date of the new medication supply are considered a gap; this maintains consistency with gaps between episodes (whether they are constructed based on the maximum permissible gap rule or the medication change rule).

Value

A `data.frame` or `data.table` with the following columns (or `NULL` if no treatment episodes could be computed):

- `patid` the patient ID.
- `episode.ID` the episode unique ID (increasing sequentially).
- `episode.start` the episode start date.
- `end.episode.gap.days` the corresponding gap days of the last event in this episode.
- `episode.duration` the episode duration in days.
- `episode.end` the episode end date.

compute_event_durations

Computation of event durations.

Description

Computes event durations based on dispensing, prescription, and hospitalization data and returns a `data.frame` which can be used with the CMA constructors in `AdhereR`.

Usage

```
compute_event_durations(dispatch.data = NULL, presc.data = NULL,
  hosp.data = NULL, ID.colname, presc.date.colname, disp.date.colname,
  date.format = "%d.%m.%Y", medication.class.colnames,
  total.dose.colname, presc.daily.dose.colname, presc.duration.colname,
  visit.colname, force.init.presc = FALSE, force.presc.renew = FALSE,
  split.on.dosage.change = TRUE, trt.interruption = c("continue",
  "discard", "carryover")[1], suppress.warnings = FALSE,
  return.data.table = FALSE, ...)
```

Arguments

disp.data	A <code>data.frame</code> or <code>data.table</code> containing the dispensing events. Must contain, at a minimum, the patient unique ID, one medication identifier, the dispensing date, and total dispensed dose, and might also contain additional columns to identify and group medications (the actual column names are defined in the <code>medication.class.colnames</code> parameter).
presc.data	A <code>data.frame</code> containing the prescribing events. Must contain, at a minimum, the same unique patient ID and medication identifier(s) as the dispensing data, the prescription date, the daily prescribed dose, and the prescription duration. Optionally, it might also contain a visit number.
hosp.data	Optional, <code>NULL</code> or a <code>data.frame</code> containing the hospitalization periods (or other situations where patients may not use their own supply, e.g. during incarcerations). Must contain the same unique patient ID as dispensing and prescription data, and the start and end dates of the hospitalizations with the exact column names <code>DATE.IN</code> and <code>DATE.OUT</code> .
ID.colname	A <i>string</i> , the name of the column in <code>disp.data</code> , <code>presc.data</code> , and <code>hosp.data</code> containing the unique patient ID.
presc.date.colname	A <i>string</i> , the name of the column in <code>presc.data</code> containing the prescription date (in the format given in the <code>date.format</code> parameter).
disp.date.colname	A <i>string</i> , the name of the column in <code>disp.data</code> containing the dispensing date (in the format given in the <code>date.format</code> parameter).
date.format	A <i>string</i> giving the format of the dates used in the data and the other parameters; see the format parameters of the <code>as.Date</code> function for details (NB, this concerns only the dates given as strings and not as <code>Date</code> objects).
medication.class.colnames	A <code>Vector</code> of <i>strings</i> , the name(s) of the column(s) in <code>disp.data</code> and <code>presc.data</code> containing the classes/types/groups of medication.
total.dose.colname	A <i>string</i> , the name of the column in <code>disp.data</code> containing the total dispensed dose as numeric (e.g. 500 for 10 tablets of 50 mg).
presc.daily.dose.colname	A <i>string</i> , the name of the column in <code>presc.data</code> containing the daily prescribed dose as numeric (e.g. 50 for 50 mg once per day, or 25 for 50 mg once ever 2 days).
presc.duration.colname	A <i>string</i> , the name of the column in <code>presc.data</code> containing the duration of the prescription as numeric or <code>NA</code> if duration is unknown.
visit.colname	A <i>string</i> , the name of the column in <code>presc.data</code> containing the number of the visit or a new column name if the prescribing data does not contain such a column.
force.init.presc	<i>Logical</i> . If <code>TRUE</code> advance the date of the first prescription event to the date of the first dispensing event, if the first prescription event is after the first dispensing

	event for a specific medication. Only if the first prescription event is not limited in duration (as indicated in the <code>presc.duration.colname</code>).
<code>force.presc.renew</code>	<i>Logical</i> or <i>string</i> . If TRUE require a new prescription for all medications for every prescription event (visit), otherwise prescriptions end on the first visit without renewal. If <i>string</i> , the name of the column containing the <code>Logical</code> for each medication class separately.
<code>split.on.dosage.change</code>	<i>Logical</i> or <i>string</i> . If TRUE split the dispensing event on days with dosage change and create a new event with the new dosage for the remaining supply. If <i>string</i> , the name of the column containing the <code>Logical</code> for each medication class separately. Important if carryover should be considered later on.
<code>trt.interruption</code>	can be either <i>continue</i> , <i>discard</i> , or <i>carryover</i> , and indicates how to handle supplies affected by treatment interruptions or hospitalizations. With <i>continue</i> , interruptions have no effect on durations and dispensing start dates. With <i>discard</i> , supplies are truncated at the beginning of an interruption or hospitalization and the remaining supply is discarded. With <i>carryover</i> , supplies are truncated at the beginning of an interruption or hospitalization and a new dispensing start event with the remaining supply is created after the end of the interruption or hospitalization.
<code>suppress.warnings</code>	<i>Logical</i> , if TRUE don't show any warnings.
<code>return.data.table</code>	<i>Logical</i> , if TRUE return a <code>data.table</code> object, otherwise a <code>data.frame</code> .
<code>...</code>	other possible parameters.

Details

Computation of CMAs requires a supply duration for medications dispensed to patients. If medications are not supplied for fixed durations but as a quantity that may last for various durations based on the prescribed dose, the supply duration has to be calculated based on dispensed and prescribed doses. Treatments may be interrupted and resumed at later times, for which existing supplies may or may not be taken into account. Patients may be hospitalized or incarcerated, and may not use their own supplies during these periods. This function calculates supply durations, taking into account the aforementioned situations and providing various parameters for flexible adjustments.

Value

A `data.frame` or `data.table` with the following columns:

- `ID.colname` the unique patient ID, as given by the `ID.colname` parameter.
- `medication.class.colnames` the column(s) with classes/types/groups of medication, as given by the `medication.class.colnames` parameter.
- `total.dose.colname` the total dispensed dose, as given by the `total.dose.colname` parameter.
- `disp.date.colname` the date of the dispensing event, as given by the `disp.date.colname` parameter.

- DISP.START the start date of the dispensing event, either the same as in disp.date.colname or a later date in case of dosage changes or treatment interruptions/hospitalizations.
- presc.daily.dose.colname the prescribed daily dose, as given by the presc.daily.dose.colname parameter.
- DURATION the calculated duration of the supply, based on the total dispensed dose and the prescribed daily dose, starting from the DISP.START date.
- START.PRESC the start date of the prescription episode.
- END.PRESC the end date of the prescription episode.
- HOSP.DURATION the number of days during the current supply period affected by hospitalizations.
- tot.presc.interruptions the total number of prescription interruptions per patient for a specific medication.
- tot.dosage.changes the total number of dosage changes per patient for a specific medication.

Examples

```
event_durations <- compute_event_durations(disp.data = durcomp.dispensing[1:3,],
                                           presc.data = durcomp.prescribing,
                                           hosp.data = durcomp.hospitalisation,
                                           ID.colname = "ID",
                                           presc.date.colname = "DATE.PRESC",
                                           disp.date.colname = "DATE.DISP",
                                           date.format = "%Y-%m-%d",
                                           medication.class.colnames = c("ATC.CODE",
                                                                              "UNIT", "FORM"),
                                           total.dose.colname = "TOTAL.DOSE",
                                           presc.daily.dose.colname = "DAILY.DOSE",
                                           presc.duration.colname = "PRESC.DURATION",
                                           visit.colname = "VISIT",
                                           force.init.presc = TRUE,
                                           force.presc.renew = TRUE,
                                           split.on.dosage.change = TRUE,
                                           trt.interruption = "continue",
                                           suppress.warnings = FALSE,
                                           return.data.table = TRUE);
```

durcomp.dispensing *Example dispensing events for 16 patients.*

Description

A sample dataset containing dispensing events (one per row) for 16 patients over a period of roughly 24 months (1794 events in total). This is the appropriate format to compute event durations with the `compute_event_durations` function. Each row represent an individual dispensing record for a specific dose of a specific medication for a patient at a given date. Visit number and Duration are optional, and more than one column to group medications can be supplied (such as Form or Unit).

Usage

```
durcomp.dispensing
```

Format

A data frame with 1794 rows and 6 variables:

ID *integer* here; patient unique identifier. Can also be *string*.

DATE.DISP *Date* here; the dispensing event date, by default in the yyyy-mm-dd format. Can also be *string*.

ATC.CODE *character*; the medication type, according to the WHO ATC classification system. This can be a researcher-defined classification depending on study aims (e.g., based on therapeutic use, mechanism of action, chemical molecule, or pharmaceutical formulation). The `compute_event_durations` function will match prescribed medication to dispensed medications based on this variable.

UNIT *integer*; the unit of the dispensed dose. This is optional and can be used as a separate variable to match between prescription and dispensing events.

FORM *character*; the galenic form of the dispensed preparation. This is optional and can be used as a separate variable to match between prescription and dispensing events.

TOTAL.DOSE *numeric*; the total dispensed dose supplied at this medication event (e.g., 5000 for 10 tablets of 500 mg).

```
durcomp.hospitalisation
```

Example hospitalization events for 10 patients.

Description

A sample dataset containing hospitalization periods (one per row) for 10 patients over a period of roughly 18 months (28 events in total). This is the appropriate format to compute event durations with the `compute_event_durations` function. Each row represent an individual hospitalization period of a patient for whom event durations should be calculated. Besides hospitalizations, this could cover other situations where patients may not use their own supply, e.g. during incarcerations. All column names must match the format provided in this example.

Usage

```
durcomp.hospitalisation
```

Format

A data frame with 28 rows and 3 variables:

ID *integer* here; patient unique identifier. Can also be *string*.

DATE.IN *Date* here; the start of the hospitalization period, by default in the yyyy-mm-dd format. Can also be *string*.

DATE.OUT *Date*; the end of the hospitalization period, by default in the yyyy-mm-dd format. Can also be *string*.

durcomp.prescribing *Example prescription events for 16 patients.*

Description

A sample dataset containing prescription events (one per row) for 16 patients over a period of roughly 15 months (1502 events in total). This is the appropriate format to compute event durations with the `compute_event_durations` function. Each row represent an individual prescription record for a specific dose of a specific medication for a patient at a given date. Visit number and Duration are optional, and more than one column to group medications can be supplied (such as Form or Unit).

Usage

durcomp.prescribing

Format

A data table with 1502 rows and 8 variables:

ID *integer* here; patient unique identifier. Can also be *string*.

DATE.PRESC *Date* here; the prescription event date, by default in the yyyy-mm-dd format. Can also be *string*.

VISIT *integer*; the consecutive number of the prescription instances. This column is optional and will be generated internally when not supplied. It is used to identify treatment interruptions.

ATC.CODE *character*; the medication type, according to the WHO ATC classification system. This can be a researcher-defined classification depending on study aims (e.g., based on therapeutic use, mechanism of action, chemical molecule, or pharmaceutical formulation). The `compute_event_durations` function will match prescribed medication to dispensed medications based on this variable.

FORM *character*; the galenic form of the prescribed preparation. This is optional and can be used as a separate variable to match between prescription and dispensing events.

UNIT *integer*; the unit of the prescribed dose. This is optional and can be used as a separate variable to match between prescription and dispensing events.

PRESC.DURATION *numeric*; the duration (in days) for which the prescription is intended. Can be NA if the prescription is continuous without a fixed end date.

DAILY.DOSE *numeric*; the daily dose prescribed during this event (e.g., 50 for 1 tablet of 50 mg per day or 25 for 1 tablet of 50 mg every two days).

```
getCallerWrapperLocation
    getCallerWrapperLocation.
```

Description

This function returns the full path to where the various wrappers that can call AdhereR are installed.

Usage

```
getCallerWrapperLocation(callig.platform = c("python3")[1],
    full.path = FALSE)
```

Arguments

`callig.platform` A *string* specifying the desired wrapper. Currently it can be "python3".

`full.path` A *logical* specifying if the returned path should also include the wrapper's main file name.

Details

In most cases, these wrappers are one or more files in the calling language that may be directly used as such. For more details see the vignette describing the included reference Python 3 wrapper.

Value

The full path to the requested wrapper or NULL if none exists.

```
getCMA Access the actual CMA estimate from a CMA object.
```

Description

Retrieve the actual CMA estimate(s) encapsulated in a simple, per episode, or sliding window CMA object.

Usage

```
getCMA(x)
```

Arguments

`x` a CMA object.

Value

a *data.frame* containing the CMA estimate(s).

Examples

```
cma1 <- CMA1(data=med.events,
             ID.colname="PATIENT_ID",
             event.date.colname="DATE",
             event.duration.colname="DURATION",
             followup.window.start=30,
             observation.window.start=30,
             observation.window.duration=365,
             date.format="%m/%d/%Y"
           );
getCMA(cma1);
## Not run:
cmaE <- CMA_per_episode(CMA="CMA1",
                       data=med.events,
                       ID.colname="PATIENT_ID",
                       event.date.colname="DATE",
                       event.duration.colname="DURATION",
                       event.daily.dose.colname="PERDAY",
                       medication.class.colname="CATEGORY",
                       carry.only.for.same.medications=FALSE,
                       consider.dosage.change=FALSE,
                       followup.window.start=0,
                       observation.window.start=0,
                       observation.window.duration=365,
                       date.format="%m/%d/%Y"
                     );
getCMA(cmaE);
## End(Not run)
```

med.events

Example medication events records for 100 patients.

Description

An artificial dataset containing medication events (one per row) for 100 patients (1080 events in total). This is the dataset format appropriate for medication adherence analyses performed with the R package *AdhereR*. Medication events represent individual records of prescribing or dispensing a specific medication for a patient at a given date. Dosage and medication type is optional (only needed if calculation of adherence or persistence takes into account changes in dosage and type of medication).

Usage

med.events

Format

A data frame with 1080 rows and 5 variables:

PATIENT_ID *integer* here; patient unique identifier. Can also be *string*.

DATE *character*; the medication event date, by default in the mm/dd/yyyy format. It may represent a prescribing or dispensing date.

PERDAY *integer*; the daily dosage prescribed for the medication supplied at this medication event (i.e. how many doses should be taken daily according to the prescription). This column is optional, as it is not considered in all functions but may be relevant for specific research or clinical contexts. All values should be > 0.

CATEGORY *character*; the medication type, here two placeholder labels, 'medA' and 'medB'. This is a researcher-defined classification depending on study aims (e.g., based on therapeutic use, mechanism of action, chemical molecule, or pharmaceutical formulation). This column is optional, as it is not considered in all functions but may be relevant for specific research or clinical contexts.

DURATION *integer*; the medication event duration in days (i.e. how many days the medication supplied would last if used as prescribed); may be available in the extraction or computed based on quantity supplied (the number of doses prescribed or dispensed on that occasion) and daily dosage. All values should be > 0.

plot.CMA0

Plot CMA0 objects.

Description

Plots the events (prescribing or dispensing) data encapsulated in a basic CMA0 object.

Usage

```
## S3 method for class 'CMA0'
plot(x, ..., patients.to.plot = NULL, duration = NA,
      align.all.patients = FALSE, show.period = c("dates", "days")[2],
      period.in.days = 90, show.legend = TRUE, legend.x = "right",
      legend.y = "bottom", legend.bkg.opacity = 0.5, legend.cex = 0.75,
      legend.cex.title = 1, cex = 1, cex.axis = 0.75, cex.lab = 1,
      col.cats = rainbow, unspecified.category.label = "drug",
      lty.event = "solid", lwd.event = 2, pch.start.event = 15,
      pch.end.event = 16, print.dose = FALSE, cex.dose = 0.75,
      print.dose.outline.col = "white", print.dose.centered = FALSE,
      plot.dose = FALSE, lwd.event.max.dose = 8,
      plot.dose.lwd.across.medication.classes = FALSE,
      col.continuation = "black", lty.continuation = "dotted",
      lwd.continuation = 1, col.na = "lightgray",
      highlight.followup.window = TRUE, followup.window.col = "green",
      highlight.observation.window = TRUE,
```

```

observation.window.col = "yellow", observation.window.density = 35,
observation.window.angle = -30, observation.window.opacity = 0.3,
bw.plot = FALSE, print.CMA = TRUE, plot.CMA = TRUE,
CMA.plot.ratio = 0.1, min.plot.size.in.characters.horiz = 10,
min.plot.size.in.characters.vert = 0.5, max.patients.to.plot = 100)

```

Arguments

x	A CMA0 or derived object, representing the CMA to plot
...	other possible parameters
patients.to.plot	A vector of <i>strings</i> containing the list of patient IDs to plot (a subset of those in the cma object), or NULL for all
duration	A <i>number</i> , the total duration (in days) of the whole period to plot; in NA it is automatically determined from the event data such that the whole dataset fits.
align.all.patients	<i>Logical</i> , should all patients be aligned (i.e., the actual dates are discarded and all plots are relative to the earliest date)?
show.period	A <i>string</i> , if "dates" show the actual dates at the regular grid intervals, while for "days" (the default) shows the days since the beginning; if align.all.patients == TRUE, show.period is taken as "days".
period.in.days	The <i>number</i> of days at which the regular grid is drawn (or 0 for no grid).
show.legend	<i>Logical</i> , should the legend be drawn?
legend.x	The position of the legend on the x axis; can be "left", "right" (default), or a <i>numeric</i> value.
legend.y	The position of the legend on the y axis; can be "bottom" (default), "top", or a <i>numeric</i> value.
legend.bkg.opacity	A <i>number</i> between 0.0 and 1.0 specifying the opacity of the legend background.
cex, cex.axis, cex.lab, legend.cex, legend.cex.title	<i>numeric</i> values specifying the cex of the various types of text.
col.cats	A <i>color</i> or a <i>function</i> that specifies the single colour or the colour palette used to plot the different medication; by default rainbow, but we recommend, whenever possible, a colorblind-friendly palette such as viridis or colorblind_pal.
unspecified.category.label	A <i>string</i> giving the name of the unspecified (generic) medication category.
lty.event, lwd.event, pch.start.event, pch.end.event	The style of the event (line style, width, and start and end symbols).
print.dose	<i>Logical</i> , should the daily dose be printed as text?
cex.dose	<i>Numeric</i> , if daily dose is printed, what text size to use?
print.dose.outline.col	If NA, don't print dose text with outline, otherwise a color name/code for the outline.
print.dose.centered	<i>Logical</i> , print the daily dose centered on the segment or slightly below it?

plot.dose	<i>Logical</i> , should the daily dose be indicated through segment width?
lwd.event.max.dose	<i>Numeric</i> , the segment width corresponding to the maximum daily dose (must be \geq lwd.event but not too big either).
plot.dose.lwd.across.medication.classes	<i>Logical</i> , if TRUE, the line width of the even is scaled relative to all medication classes (i.e., relative to the global minimum and maximum doses), otherwise it is scale relative only to its medication class.
col.continuation, lty.continuation, lwd.continuation	The style of the "continuation" lines connecting consecutive events (colour, line style and width).
col.na	The colour used for missing event data.
highlight.followup.window	<i>Logical</i> , should the follow-up window be plotted?
followup.window.col	The follow-up window's colour.
highlight.observation.window	<i>Logical</i> , should the observation window be plotted?
observation.window.col, observation.window.density, observation.window.angle, observation.window.op	Attributes of the observation window (colour, shading density, angle and opacity).
bw.plot	<i>Logical</i> , should the plot use grayscale only (i.e., the <code>gray.colors</code> function)?
print.CMA	<i>Logical</i> , should the CMA values be printed?
plot.CMA	<i>Logical</i> , should the CMA values be represented graphically?
CMA.plot.ratio	A <i>number</i> , the proportion of the total horizontal plot space to be allocated to the CMA plot. <i>Numeric</i> , the minimum size of the plotting surface in characters; horizontally (<code>min.plot.size.in.characters.horiz</code>) refers to the the whole duration of the events to plot; vertically (<code>min.plot.size.in.characters.vert</code>) refers to a single event.
min.plot.size.in.characters.horiz, min.plot.size.in.characters.vert	<i>Numeric</i> , the minimum size of the plotting surface in characters; horizontally (<code>min.plot.size.in.characters.horiz</code>) refers to the the whole duration of the events to plot; vertically (<code>min.plot.size.in.characters.vert</code>) refers to a single event.
max.patients.to.plot	<i>Numeric</i> , the maximum patients to attempt to plot.

Details

The x-axis represents time (either in days since the earliest date or as actual dates), with consecutive events represented as ascending on the y-axis.

Each event is represented as a segment with style `lty.event` and line width `lwd.event` starting with a `pch.start.event` and ending with a `pch.end.event` character, coloured with a unique color as given by `col.cats`, extending from its start date until its end date. Consecutive events are thus represented on consecutive levels of the y-axis and are connected by a "continuation" line with `col.continuation` colour, `lty.continuation` style and `lwd.continuation` width; these

continuation lines are purely visual guides helping to perceive the sequence of events, and carry no information about the availability of medication in this interval.

When several patients are displayed on the same plot, they are organized vertically, and alternating bands (white and gray) help distinguish consecutive patients. Implicitly, all patients contained in the `cma` object will be plotted, but the `patients.to.plot` parameter allows the selection of a subset of patients.

Examples

```
cma0 <- CMA0(data=med.events,
             ID.colname="PATIENT_ID",
             event.date.colname="DATE",
             event.duration.colname="DURATION",
             event.daily.dose.colname="PERDAY",
             medication.class.colname="CATEGORY",
             followup.window.start=0,
             followup.window.start.unit="days",
             followup.window.duration=2*365,
             followup.window.duration.unit="days",
             observation.window.start=30,
             observation.window.start.unit="days",
             observation.window.duration=365,
             observation.window.duration.unit="days",
             date.format="%m/%d/%Y",
             summary="Base CMA");
plot(cma0, patients.to.plot=c("1","2"));
```

plot.CMA1

Plot CMA0-derived objects.

Description

Plots the event data and estimated CMA encapsulated in objects derived from `CMA0`.

Usage

```
## S3 method for class 'CMA1'
plot(x, ..., patients.to.plot = NULL, duration = NA,
     align.all.patients = FALSE, align.first.event.at.zero = TRUE,
     show.period = c("dates", "days")[2], period.in.days = 90,
     show.legend = TRUE, legend.x = "right", legend.y = "bottom",
     legend.bkg.opacity = 0.5, legend.cex = 0.75, legend.cex.title = 1,
     cex = 1, cex.axis = 0.75, cex.lab = 1, show.cma = TRUE,
     col.cats = rainbow, unspecified.category.label = "drug",
     lty.event = "solid", lwd.event = 2, pch.start.event = 15,
     pch.end.event = 16, show.event.intervals = TRUE,
     col.na = "lightgray", print.CMA = TRUE, CMA.cex = 0.5,
     plot.CMA = TRUE, CMA.plot.ratio = 0.1, CMA.plot.col = "lightgreen",
```

```

CMA.plot.border = "darkgreen", CMA.plot.bkg = "aquamarine",
CMA.plot.text = CMA.plot.border, highlight.followup.window = TRUE,
followup.window.col = "green", highlight.observation.window = TRUE,
observation.window.col = "yellow", observation.window.density = 35,
observation.window.angle = -30, observation.window.opacity = 0.3,
show.real.obs.window.start = TRUE, real.obs.window.density = 35,
real.obs.window.angle = 30, print.dose = FALSE, cex.dose = 0.75,
print.dose.outline.col = "white", print.dose.centered = FALSE,
plot.dose = FALSE, lwd.event.max.dose = 8,
plot.dose.lwd.across.medication.classes = FALSE, bw.plot = FALSE,
min.plot.size.in.characters.horiz = 10,
min.plot.size.in.characters.vert = 0.5, max.patients.to.plot = 100)

## S3 method for class 'CMA2'
plot(...)

## S3 method for class 'CMA3'
plot(...)

## S3 method for class 'CMA4'
plot(...)

## S3 method for class 'CMA5'
plot(...)

## S3 method for class 'CMA6'
plot(...)

## S3 method for class 'CMA7'
plot(...)

## S3 method for class 'CMA8'
plot(...)

## S3 method for class 'CMA9'
plot(...)

```

Arguments

x	A CMA0 or derived object, representing the CMA to plot
...	other possible parameters
patients.to.plot	A vector of <i>strings</i> containing the list of patient IDs to plot (a subset of those in the cma object), or NULL for all
duration	A <i>number</i> , the total duration (in days) of the whole period to plot; in NA it is automatically determined from the event data such that the whole dataset fits.
align.all.patients	<i>Logical</i> , should all patients be aligned (i.e., the actual dates are discarded and

	all plots are relative to the earliest date)?
align.first.event.at.zero	<i>Logical</i> , should the first event be placed at the origin of the time axis (at 0)?
show.period	A <i>string</i> , if "dates" show the actual dates at the regular grid intervals, while for "days" (the default) shows the days since the beginning; if align.all.patients == TRUE, show.period is taken as "days".
period.in.days	The <i>number</i> of days at which the regular grid is drawn (or 0 for no grid).
show.legend	<i>Logical</i> , should the legend be drawn?
legend.x	The position of the legend on the x axis; can be "left", "right" (default), or a <i>numeric</i> value.
legend.y	The position of the legend on the y axis; can be "bottom" (default), "top", or a <i>numeric</i> value.
legend.bkg.opacity	A <i>number</i> between 0.0 and 1.0 specifying the opacity of the legend background.
cex, cex.axis, cex.lab, legend.cex, legend.cex.title, CMA.cex	<i>numeric</i> values specifying the cex of the various types of text.
show.cma	<i>Logical</i> , should the CMA type be shown in the title?
col.cats	A <i>color</i> or a <i>function</i> that specifies the single colour or the colour palette used to plot the different medication; by default rainbow, but we recommend, whenever possible, a colorblind-friendly palette such as viridis or colorblind_pal.
unspecified.category.label	A <i>string</i> giving the name of the unspecified (generic) medication category.
lty.event, lwd.event, pch.start.event, pch.end.event	The style of the event (line style, width, and start and end symbols).
show.event.intervals	<i>Logical</i> , should the actual event intervals be shown?
col.na	The colour used for missing event data.
print.CMA	<i>Logical</i> , should the CMA values be printed?
plot.CMA	<i>Logical</i> , should the CMA values be represented graphically?
CMA.plot.ratio	A <i>number</i> , the proportion of the total horizontal plot space to be allocated to the CMA plot.
CMA.plot.col, CMA.plot.border, CMA.plot.bkg, CMA.plot.text	<i>Strings</i> giving the colours of the various components of the CMA plot.
highlight.followup.window	<i>Logical</i> , should the follow-up window be plotted?
followup.window.col	The follow-up window's colour.
highlight.observation.window	<i>Logical</i> , should the observation window be plotted?
observation.window.col, observation.window.density, observation.window.angle, observation.window.opacity	Attributes of the observation window (colour, shading density, angle and opacity).

`show.real.obs.window.start`, `real.obs.window.density`, `real.obs.window.angle`
 For some CMAs, the observation window might be adjusted, in which case should it be plotted and with that attributes?

`print.dose` *Logical*, should the daily dose be printed as text?

`cex.dose` *Numeric*, if daily dose is printed, what text size to use?

`print.dose.outline.col`
 If NA, don't print dose text with outline, otherwise a color name/code for the outline.

`print.dose.centered`
Logical, print the daily dose centered on the segment or slightly below it?

`plot.dose` *Logical*, should the daily dose be indicated through segment width?

`lwd.event.max.dose`
Numeric, the segment width corresponding to the maximum daily dose (must be \geq `lwd.event` but not too big either).

`plot.dose.lwd.across.medication.classes`
Logical, if TRUE, the line width of the even is scaled relative to all medication classes (i.e., relative to the global minimum and maximum doses), otherwise it is scale relative only to its medication class.

`bw.plot` *Logical*, should the plot use grayscale only (i.e., the `gray.colors` function)?

`min.plot.size.in.characters.horiz`, `min.plot.size.in.characters.vert`
Numeric, the minimum size of the plotting surface in characters; horizontally (`min.plot.size.in.characters.horiz`) refers to the the whole duration of the events to plot; vertically (`min.plot.size.in.characters.vert`) refers to a single event.

`max.patients.to.plot`
Numeric, the maximum patients to attempt to plot.

Details

Please note that this function plots objects inheriting from `CMA0` but not objects of type `CMA0` itself (these are plotted by `plot.CMA0`).

The x-axis represents time (either in days since the earliest date or as actual dates), with consecutive events represented as ascending on the y-axis.

Each event is represented as a segment with style `lty.event` and line width `lwd.event` starting with a `pch.start.event` and ending with a `pch.end.event` character, coloured with a unique color as given by `col.cats`, extending from its start date until its end date. Superimposed on these are shown the event intervals and gap days as estimated by the particular CMA method, more precisely plotting the start and end of the available events as solid filled-in rectangles, and the event gaps as shaded rectangles.

The follow-up and the observation windows are plotted as an empty rectangle and as shaded rectangle, respectively (for some CMAs the observation window might be adjusted in which case the adjustment may also be plotted using a different shading).

The CMA estimates can be visually represented as well in the left side of the figure using bars (sometimes the estimates can go above 100%, in which case the maximum possible bar filling is adjusted to reflect this).

When several patients are displayed on the same plot, they are organized vertically, and alternating bands (white and gray) help distinguish consecutive patients. Implicitly, all patients contained in

the cma object will be plotted, but the `patients.to.plot` parameter allows the selection of a subset of patients.

Finally, the y-axis shows the patient ID and possibly the CMA estimate as well.

Examples

```
cma1 <- CMA1(data=med.events,
             ID.colname="PATIENT_ID",
             event.date.colname="DATE",
             event.duration.colname="DURATION",
             followup.window.start=30,
             observation.window.start=30,
             observation.window.duration=365,
             date.format="%m/%d/%Y"
            );
plot(cma1, patients.to.plot=c("1", "2"));
```

plot.CMA_per_episode *Plot CMA_per_episode and CMA_sliding_window objects.*

Description

Plots the event data and the estimated CMA per treatment episode and sliding window, respectively.

Usage

```
## S3 method for class 'CMA_per_episode'
plot(x, ..., patients.to.plot = NULL,
     duration = NA, align.all.patients = FALSE,
     align.first.event.at.zero = TRUE, show.period = c("dates",
     "days")[2], period.in.days = 90, show.legend = TRUE,
     legend.x = "right", legend.y = "bottom", legend.bkg.opacity = 0.5,
     cex = 1, cex.axis = 0.75, cex.lab = 1, show.cma = TRUE,
     col.cats = rainbow, unspecified.category.label = "drug",
     lty.event = "solid", lwd.event = 2, pch.start.event = 15,
     pch.end.event = 16, col.na = "lightgray",
     col.continuation = "black", lty.continuation = "dotted",
     lwd.continuation = 1, print.CMA = TRUE, CMA.cex = 0.5,
     plot.CMA = TRUE, plot.CMA.as.histogram = TRUE,
     CMA.plot.ratio = 0.1, CMA.plot.col = "lightgreen",
     CMA.plot.border = "darkgreen", CMA.plot.bkg = "aquamarine",
     CMA.plot.text = CMA.plot.border, highlight.followup.window = TRUE,
     followup.window.col = "green", highlight.observation.window = TRUE,
     observation.window.col = "yellow", observation.window.density = 35,
     observation.window.angle = -30, show.real.obs.window.start = TRUE,
     real.obs.window.density = 35, real.obs.window.angle = 30,
     bw.plot = FALSE)
```

```
## S3 method for class 'CMA_sliding_window'
plot(...)
```

Arguments

x A CMA0 or derived object, representing the CMA to plot

... other parameters (to be passed to the estimation and plotting of the simple CMA)

patients.to.plot A vector of *strings* containing the list of patient IDs to plot (a subset of those in the *cma* object), or NULL for all

duration A *number*, the total duration (in days) of the whole period to plot; in NA it is automatically determined from the event data such that the whole dataset fits.

align.all.patients *Logical*, should all patients be aligned (i.e., the actual dates are discarded and all plots are relative to the earliest date)?

align.first.event.at.zero *Logical*, should the first event be placed at the origin of the time axis (at 0)?

show.period A *string*, if "dates" show the actual dates at the regular grid intervals, while for "days" (the default) shows the days since the beginning; if `align.all.patients == TRUE`, `show.period` is taken as "days".

period.in.days The *number* of days at which the regular grid is drawn (or 0 for no grid).

show.legend *Logical*, should the legend be drawn?

legend.x The position of the legend on the x axis; can be "left", "right" (default), or a *numeric* value.

legend.y The position of the legend on the y axis; can be "bottom" (default), "top", or a *numeric* value.

legend.bkg.opacity A *number* between 0.0 and 1.0 specifying the opacity of the legend background.

cex, cex.axis, cex.lab *numeric* values specifying the cex of the various types of text.

show.cma *Logical*, should the CMA type be shown in the title?

col.cats A *color* or a *function* that specifies the single colour or the colour palette used to plot the different medication; by default `rainbow`, but we recommend, whenever possible, a colorblind-friendly palette such as `viridis` or `colorblind_pal`.

unspecified.category.label A *string* giving the name of the unspecified (generic) medication category.

lty.event, lwd.event, pch.start.event, pch.end.event The style of the event (line style, width, and start and end symbols).

col.na The colour used for missing event data.

col.continuation, lty.continuation, lwd.continuation The color, style and width of the continuation lines connecting consecutive events.

print.CMA *Logical*, should the CMA values be printed?

CMA.cex ... and, if printed, what cex (*numeric*) to use?

plot.CMA	<i>Logical</i> , should the CMA values be represented graphically?
plot.CMA.as.histogram	<i>Logical</i> , should the CMA plot be a histogram or a (truncated) density plot? Please note that it is TRUE by default for CMA_per_episode and FALSE for CMA_sliding_window, because usually there are more sliding windows than episodes. Also, the density estimate cannot be estimated for less than three different values.
CMA.plot.ratio	A <i>number</i> , the proportion of the total horizontal plot space to be allocated to the CMA plot.
CMA.plot.col, CMA.plot.border, CMA.plot.bkg, CMA.plot.text	<i>Strings</i> giving the colours of the various components of the CMA plot.
highlight.followup.window	<i>Logical</i> , should the follow-up window be plotted?
followup.window.col	The follow-up window colour.
highlight.observation.window	<i>Logical</i> , should the observation window be plotted?
observation.window.col, observation.window.density, observation.window.angle	Attributes of the observation window (colour, shading density and angle).
show.real.obs.window.start, real.obs.window.density, real.obs.window.angle	For some CMAs, the observation window might be adjusted, in which case should it be plotted and with that attributes?
bw.plot	<i>Logical</i> , should the plot use grayscale only (i.e., the gray.colors function)?

Details

The x-axis represents time (either in days since the earliest date or as actual dates), with consecutive events represented as ascending on the y-axis.

Each event is represented as a segment with style `lty.event` and line width `lwd.event` starting with a `pch.start.event` and ending with a `pch.end.event` character, coloured with a unique color as given by `col.cats`, extending from its start date until its end date. Consecutive events are thus represented on consecutive levels of the y-axis and are connected by a "continuation" line with `col.continuation` colour, `lty.continuation` style and `lwd.continuation` width; these continuation lines are purely visual guides helping to perceive the sequence of events, and carry no information about the availability of medicine in this interval.

Above these, the treatment episodes or the sliding windows are represented in a stacked manner from the earliest (left, bottom of the stack) to the latest (right, top of the stack), each showing the CMA as percent fill (capped at 100% even if CMA values may be higher) and also as text.

The follow-up and the observation windows are plotted as empty an rectangle and as shaded rectangle, respectively (for some CMAs the observation window might be adjusted in which case the adjustment may also be plotted using a different shading).

The kernel density ("smoothed histogram") of the CMA estimates across treatment episodes/sliding windows (if more than 2) can be visually represented as well in the left side of the figure (NB, their horizontal scales may be different across patients).

When several patients are displayed on the same plot, they are organized vertically, and alternating bands (white and gray) help distinguish consecutive patients. Implicitly, all patients contained in

the `cma` object will be plotted, but the `patients.to.plot` parameter allows the selection of a subset of patients.

Finally, the y-axis shows the patient ID and possibly the CMA estimate as well.

Any not explicitly defined arguments are passed to the simple CMA estimation and plotting function; therefore, for more info about possible estimation parameters please see the help for the appropriate simple CMA, and for possible aesthetic tweaks, please see the help for their plotting.

See Also

See the simple CMA estimation [CMA1](#) to [CMA9](#) and plotting [plot.CMA1](#) functions for extra parameters.

Examples

```
## Not run:
cmaW <- CMA_sliding_window(CMA=CMA1,
                           data=med.events,
                           ID.colname="PATIENT_ID",
                           event.date.colname="DATE",
                           event.duration.colname="DURATION",
                           event.daily.dose.colname="PERDAY",
                           medication.class.colname="CATEGORY",
                           carry.only.for.same.medications=FALSE,
                           consider.dosage.change=FALSE,
                           followup.window.start=0,
                           observation.window.start=0,
                           observation.window.duration=365,
                           sliding.window.start=0,
                           sliding.window.start.unit="days",
                           sliding.window.duration=90,
                           sliding.window.duration.unit="days",
                           sliding.window.step.duration=7,
                           sliding.window.step.unit="days",
                           sliding.window.no.steps=NA,
                           date.format="%m/%d/%Y"
                           );
plot(cmaW, patients.to.plot=c("1","2"));
cmaE <- CMA_per_episode(CMA=CMA1,
                        data=med.events,
                        ID.colname="PATIENT_ID",
                        event.date.colname="DATE",
                        event.duration.colname="DURATION",
                        event.daily.dose.colname="PERDAY",
                        medication.class.colname="CATEGORY",
                        carry.only.for.same.medications=FALSE,
                        consider.dosage.change=FALSE,
                        followup.window.start=0,
                        observation.window.start=0,
                        observation.window.duration=365,
                        date.format="%m/%d/%Y"
                        );
```

```
plot(cmaE, patients.to.plot=c("1","2"));
## End(Not run)
```

plot_interactive_cma *Interactive exploration CMA computation.*

Description

Interactively plot a given patient's data, allowing the real-time exploration of the various CMAs and their parameters. It can use Rstudio's manipulate library or Shiny.

Usage

```
plot_interactive_cma(data = NULL, ID = NULL, cma.class = c("simple",
  "per episode", "sliding window")[1], print.full.params = FALSE,
  ID.colname = NA, event.date.colname = NA,
  event.duration.colname = NA, event.daily.dose.colname = NA,
  medication.class.colname = NA, date.format = "%m/%d/%Y",
  followup.window.start.max = 5 * 365, followup.window.duration.max = 5
  * 365, observation.window.start.max = followup.window.start.max,
  observation.window.duration.max = followup.window.duration.max,
  align.all.patients = FALSE, align.first.event.at.zero = TRUE,
  maximum.permissible.gap.max = 2 * 365,
  sliding.window.start.max = followup.window.start.max,
  sliding.window.duration.max = 2 * 365,
  sliding.window.step.duration.max = 2 * 365, backend = c("shiny",
  "rstudio")[1], use.system.browser = FALSE,
  get.colnames.fnc = function(d) names(d),
  get.patients.fnc = function(d, idcol) unique(d[[idcol]]),
  get.data.for.patients.fnc = function(patientid, d, idcol, cols = NA,
  maxrows = NA) d[d[[idcol]] %in% patientid, ])
```

Arguments

- | | |
|------|---|
| data | Usually a data.frame containing the events (prescribing or dispensing) used to compute the CMA. Must contain, at a minimum, the patient unique ID, the event date and duration, and might also contain the daily dosage and medication type (the actual column names are defined in the following four parameters). Alternatively, this can be any other data source (for example, a connection to a database), in which case the user must redefine the arguments <code>get.colnames.fnc</code> , <code>get.patients.fnc</code> and <code>get.data.for.patients.fnc</code> appropriately. Currently, this works only when using Shiny for interactive rendering. For a working example, please see the vignette describing the interfacing with databases. |
| ID | The ID (as given in the <code>ID.colname</code> column) of the patient whose data to interactively plot (if absent, pick the first one); please not that this can be interactively selected during plotting. |

<code>cma.class</code>	The type of CMAs to plot; can be "simple" (CMA0 to CMA9), "per episode", or "sliding window".
<code>print.full.params</code>	A <i>logical</i> specifying if the values of all the parameters used to generate the current plot should be printed in the console (if <i>TRUE</i> , it can generate extremely verbose output!).
<code>ID.colname</code>	A <i>string</i> , the name of the column in data containing the unique patient ID, or NA if not defined.
<code>event.date.colname</code>	A <i>string</i> , the name of the column in data containing the start date of the event (in the format given in the <code>date.format</code> parameter), or NA if not defined.
<code>event.duration.colname</code>	A <i>string</i> , the name of the column in data containing the event duration (in days), or NA if not defined.
<code>event.daily.dose.colname</code>	A <i>string</i> , the name of the column in data containing the prescribed daily dose, or NA if not defined.
<code>medication.class.colname</code>	A <i>string</i> , the name of the column in data containing the classes/types/groups of medication, or NA if not defined.
<code>date.format</code>	A <i>string</i> giving the format of the dates used in the data and the other parameters; see the format parameters of the <code>as.Date</code> function for details (NB, this concerns only the dates given as strings and not as Date objects).
<code>followup.window.start.max</code>	The maximum number of days when the follow-up window can start.
<code>followup.window.duration.max</code>	The maximum duration of the follow-up window in days.
<code>observation.window.start.max</code>	The maximum number of days when the observation window can start.
<code>observation.window.duration.max</code>	The maximum duration of the observation window in days.
<code>align.all.patients</code>	Should the patients be aligned?
<code>align.first.event.at.zero</code>	Should the first event be put at zero?
<code>maximum.permissible.gap.max</code>	The maximum permissible gap in days.
<code>sliding.window.start.max</code>	The maximum number of days when the sliding windows can start.
<code>sliding.window.duration.max</code>	The maximum duration of the sliding windows in days.
<code>sliding.window.step.duration.max</code>	The maximum sliding window step in days.
<code>backend</code>	The plotting backend to use; "shiny" (the default) tries to use the Shiny framework, while "rstudio" uses the manipulate RStudio capability.

```

use.system.browser
    For shiny, use the system browser?
get.colnames.fnc
    A function taking as parameter the data source and returning the column names.
    Must be overridden when the data source is not derived from a data.frame.
get.patients.fnc
    A function taking as parameter the data source and the patient ID column name,
    and returns the list of all patient IDs. Must be overridden when the data source
    is not derived from a data.frame.
get.data.for.patients.fnc
    A function taking as parameter a (set of) patient ID(s), the data source, and
    the patient ID column name, and returns the list of all patient IDs. Must be
    overridden when the data source is not derived from a data.frame.
...
    Extra arguments.

```

Details

The `manipulate` is kept for backward compatibility only, as it is much more limited than Shiny and will receive no new development in the future. Shiny currently allows the use of any other data source besides a default (and usual) `data.frame` (or derived), such a connection to an SQL database. In this case, the user *must* redefine the three argument functions `get.colnames.fnc`, `get.patients.fnc` and `get.data.for.patients.fnc` which collectively define an interface for listing the column names, all the patient IDs, and for retrieving the actual data for a (set of) patient ID(s). A fully worked example is described in the vignette detailing the access to standard databases storing the patient information. For more info please see the online vignette https://htmlpreview.github.io/?https://github.com/ddediu/AdhereR/blob/master/online-only-doc/adherer_interactive_plots/adherer_interactive_plots.html.

Value

Nothing

See Also

The online vignette https://htmlpreview.github.io/?https://github.com/ddediu/AdhereR/blob/master/online-only-doc/adherer_interactive_plots/adherer_interactive_plots.html.

Examples

```

## Not run:
plot_interactive_cma(med.events,
                    ID.colname="PATIENT_ID",
                    event.date.colname="DATE",
                    event.duration.colname="DURATION",
                    event.daily.dose.colname="PERDAY",
                    medication.class.colname="CATEGORY");
## End(Not run)

```

print.CMA0	<i>Print CMA0 (and derived) objects.</i>
------------	--

Description

Prints and summarizes a basic CMA0, or derived, object.

Usage

```
## S3 method for class 'CMA0'  
print(x, ..., inline = FALSE, format = c("text",  
    "latex", "markdown"), print.params = TRUE, print.data = TRUE,  
    exclude.params = c("event.info"), skip.header = FALSE,  
    cma.type = class(cma)[1])  
  
## S3 method for class 'CMA1'  
print(...)  
  
## S3 method for class 'CMA2'  
print(...)  
  
## S3 method for class 'CMA3'  
print(...)  
  
## S3 method for class 'CMA4'  
print(...)  
  
## S3 method for class 'CMA5'  
print(...)  
  
## S3 method for class 'CMA6'  
print(...)  
  
## S3 method for class 'CMA7'  
print(...)  
  
## S3 method for class 'CMA8'  
print(...)  
  
## S3 method for class 'CMA9'  
print(...)  
  
## S3 method for class 'CMA_per_episode'  
print(x, ..., inline = FALSE,  
    format = c("text", "latex", "markdown"), print.params = TRUE,  
    print.data = TRUE, exclude.params = c("event.info"),  
    skip.header = FALSE, cma.type = class(cma)[1])
```

```
## S3 method for class 'CMA_sliding_window'
print(...)
```

Arguments

x	A CMA0 or derived object, representing the CMA to print.
...	other possible parameters
inline	<i>Logical</i> , should print inside a line of text or as a separate, extended object?
format	A <i>string</i> , the type of output: plain text ("text"; default), LaTeX ("latex") or R Markdown ("markdown").
print.params	<i>Logical</i> , should print the parameters?
print.data	<i>Logical</i> , should print a summary of the data?
exclude.params	A vector of <i>strings</i> , the names of the object fields to exclude from printing (usually, internal information irrelevant to the end-user).
skip.header	<i>Logical</i> , should the header be printed?
cma.type	A <i>string</i> , used to override the reported object's class.

Details

Can produce output for the console (text), R Markdown or LaTeX, showing various types of information.

Examples

```
cma0 <- CMA0(data=med.events,
             ID.colname="PATIENT_ID",
             event.date.colname="DATE",
             event.duration.colname="DURATION",
             event.daily.dose.colname="PERDAY",
             medication.class.colname="CATEGORY",
             followup.window.start=0,
             followup.window.start.unit="days",
             followup.window.duration=2*365,
             followup.window.duration.unit="days",
             observation.window.start=30,
             observation.window.start.unit="days",
             observation.window.duration=365,
             observation.window.duration.unit="days",
             date.format="%m/%d/%Y",
             summary="Base CMA");

cma0;
print(cma0, format="markdown");
cma1 <- CMA1(data=med.events,
             ID.colname="PATIENT_ID",
             event.date.colname="DATE",
             event.duration.colname="DURATION",
             followup.window.start=30,
             observation.window.start=30,
```

```

        observation.window.duration=365,
        date.format="%m/%d/%Y"
    );
cma1;

```

time_to_initiation *Computation of initiation times.*

Description

Computes the time between the first prescription event and the first dispensing event for each medication class.

Usage

```

time_to_initiation(presc.data = NULL, disp.data = NULL,
  ID.colname = NA, presc.start.colname = NA, disp.date.colname = NA,
  medication.class.colnames = NA, date.format = "%d.%m.%Y",
  suppress.warnings = FALSE, return.data.table = FALSE, ...)

```

Arguments

- | | |
|---------------------------|---|
| presc.data | A <i>data.frame</i> or <i>data.table</i> containing the prescription events. Must contain, at a minimum, the patient unique ID, one medication identifier, and the start date of the prescription, and might also contain additional columns to identify and group medications (the actual column names are defined in the <code>medication.class.colnames</code> parameter). |
| disp.data | A <i>data.frame</i> or <i>data.table</i> containing the dispensing events. Must contain, at a minimum, the patient unique ID, one medication identifier, the dispensing date, and might also contain additional columns to identify and group medications (the actual column names are defined in the <code>medication.class.colnames</code> parameter). |
| ID.colname | A <i>string</i> , the name of the column in <code>presc.data</code> and <code>disp.data</code> containing the unique patient ID, or NA if not defined. |
| presc.start.colname | A <i>string</i> , the name of the column in <code>presc.data</code> containing the prescription date (in the format given in the <code>date.format</code> parameter), or NA if not defined. |
| disp.date.colname | A <i>string</i> , the name of the column in <code>disp.data</code> containing the dispensing date (in the format given in the <code>date.format</code> parameter), or NA if not defined. |
| medication.class.colnames | A <i>Vector of strings</i> , the name(s) of the column(s) in data containing the classes/types/groups of medication, or NA if not defined. |
| date.format | A <i>string</i> giving the format of the dates used in the data and the other parameters; see the format parameters of the as.Date function for details (NB, this concerns only the dates given as strings and not as Date objects). |

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