

dm

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1 About

dm is a package for functions relating to data management.

2 dm and related functions

2.1 The problem we want to solve

A statistical report sometimes build an *analytical data base* (ADB) from multiple sources, variables that might need to be renamed and (if categorical) recoded (and possibly transformed), the documentation of which is *significantly boring*.

The `dm` functions is an interactive-ish way of creating an ADB which both inspects the chosen variables and “documents” the process.

2.2 The elevator pitch description

1. point to variables (from possibly different sources), one at the time, with `dm` (along with possible renaming, recoding and transformation). This gives a summary of the variable pointed to ¹, and the information is stored in a list somewhere.
2. create the ADB by `dm_create`.
3. get easy-to-print documentation of where variables came from (`dm_doc2latex`) and what recodings have been done (`dm_recode2latex`).

So, the point really is to get (3) “for free” in a way that is connected to the creation of the ADB.

2.3 The stuck-in-an-elevator description

If most variables are picked form the same source, this can be set in options.

```
opts_dm$set('default_db' = 'MyDataBase')
```

If that is done, `dm` only needs a `var` argument, the name of the var you want to add. But you can use

- `var`, name of variable in source
- `name`, optional, if you want a new name for the variable (else it is set to `var`)
- `db`, name of data frame (or similar) where `var` exists (else will look at the default location, if set)
- `recode`, a list that specifies the recoding. This is the `L` argument for the `recode` function that this package provides (see the help for that functions)
- `transf` a function for transforming (this might be something like a character-to-date function like `ymd` from the `lubridate` package)

¹Typically one wants to to this procedure anyway to sanity check all variables that are to be included.

- `comment` if you want to keep some comment about the variable
- `label` if you want to give the variable a "label" (i.e. the value of the label attribute)
- `keep.label` if `var` already has a label in `db`, should this be kept? (only if no `label` is provided)

Then as `dm` is evaluated, information about the variable is printed (to see range, levels and such).

```
dm(var = 'gEndEr', name = 'gender', label = "Perceived Gender")
  ## is followed by information being printed
```

The information about the options is stored in a list (by default "dm.doc" in an environment "dm.envir").² The key is the 'name' element, so as long as that is not changed, you can rerun the function with new arguments if something went wrong

```
dm('gEndEr', 'gender', label = "Biological Gender") ## overwrites
  ## the 'gender' entry
```

Else, kill all documentation and start again

```
dm_doc(kill = TRUE, prompt = FALSE) ## or possibly kill only this
  ## entry dm::dm_doc_set('gender', NULL)
```

The documentation can be accessed

```
myDoc <- dm_doc()
print(myDoc) ## N.B not all information is printed
```

Once all variables are created you can either store the "documentation" (and point to it later) or go on to create the ADB with `dm_create`. Specify a set of individuals (vector of id's) and, if necessary a vector of how individuals are identified in different data frames. If the `doc` argument is not provided it will just look in `dm_doc()`.

```
id_key = c('MyDataBase' = 'id', 'Other1' = 'ID', 'Other2' = 'idno')
ADB <- dm_create(set = MyDataBase$id, id.name = id_key)
```

Now you have an ADB and you can print `dm_doc()` to show where all variables come from. You can get all recodings from

²This is due to it begin poor for functions practice to write to objects in the global environment.

```
lapply(dm_doc(), FUN = function(x) x$recode_table)
```

There is also convenience functions `dm_recode2latex` and `dm_doc2latex` which will print all tables and documentation, respectively, in $\text{L}^{\text{A}}\text{T}_{\text{E}}\text{X}$ format.

2.4 Toy Example

We create some toy data

```
n <- 200
BL <- data.frame(
  id = structure(sprintf("id%d", 1:n),
    label = "identification"),
  aalder = structure(round(rnorm(n, 50, 10)),
    label = "Age at some time",
    foo = 'whatever'),
  vikt = rpois(n, 50),
  gr = sample(c('A', 'A2', 'B', 'C', 'D', 'd1', 'unknown'),
    n, TRUE),
  koon = structure(sample(c('M', 'K'), n, T),
    label = "The Sex"),
  nar = as.Date("2001-01-01") + runif(n, 0, 3650),
  stringsAsFactors = FALSE
)
BL$vikt[sample(1:n, 15)] <- NA
BL$gr[sample(1:n, 10)] <- NA
m <- .9*n
COMP <- data.frame(
  ID = structure(sample(BL$id, m),
    label = "identification"),
  foo = rbinom(m, 1, .2),
  bar = structure(rexp(m, 1/150),
    label = "Time passed")
)
```

There are some functions to help look for relevant variables.

```
db_info(BL) ## prints names and 'label' attributes
```

##	source	variable	label	class
## 1	BL	id	identification	character
## 2	BL	aalder	Age at some time	numeric
## 3	BL	vikt		integer
## 4	BL	gr		character
## 5	BL	koon	The Sex	character
## 6	BL	nar		Date

```
dm_find(pattern = 'time') ## looks in names and labels
```

```
## dm_find found:
```

```
##   source variable          label   class
## 2    BL    aalder Age at some time numeric
## 3    COMP      bar      Time passed numeric
```

Most variables of interest are in BL so set this as default.

```
opts_dm$set('default_db' = 'BL')
```

Next, we add the first variable (and view the output). We've chosen a variable with a fairly complex recoding to also illustrate the use the `recode` argument (also see the help for the `recode` function that is being utilized). `L` is a list where each entry has a vector of levels that will acquire the name of that entry, where the order of entries will be the order of the levels.

```
## 'gr' will be recoded in a more complex way
L <- list('A' = 'A2',
          'B' = NULL, ## placeholder to get the order right
          'CD' = c('C', 'D', 'd1'),
          'Unknown' = c('unknown', NA))
## # this would also work:
## L <- list('A' = c('A', 'A2'),
##          'B' = 'B',
##          'CD' = c('C', 'D', 'd1', 'something not in data'),
##          'Unknown' = c('unknown', NA))
dm('gr', recode = L, label = 'Group')
```

```
## -----
## Adding data base 'BL' entry 'gr' as variable 'gr'
## A variable of class: character
## There are 10 (5 percent) missing
##      and 7 (3.7 percent) unique values
## Since there are less than 20 unique vales we tabulate them:
##
##
##      A      A2      B      C      D      d1 unknown  <NA>
##      22      31      27      28      25      26      31      10
##
## Cross-tabulating the recoding:
##
##      gr
## gr      A  B CD Unknown
##  A      22  0  0      0
```

```
##   A2      31  0  0      0
##   B       0 27  0      0
##   C       0  0 28      0
##   D       0  0 25      0
##   d1      0  0 26      0
##   unknown 0  0  0     31
##   <NA>    0  0  0     10
```

Next, we add some more variables (but hide the output)

```
dm('aalder', 'Age')
dm('nar', 'When', comment = "wtf?")
dm('foo', 'event', db = 'COMP',
    recode = list('No' = '0', 'Yes' = 1),
    label = "An event at some time")
dm('bar', 'time', db = 'COMP', transf = log)
dm('koon', 'Gender',
    recode = list('Male' = 'M', 'Female' = 'K'))
```

When we are done, we create the ADB with

```
ADB <- dm_create(set = BL$id,
                 id.name = c('BL' = 'id', 'COMP' = 'ID'))

## Fixing variable no.1: gr
## Fixing variable no.2: Age
## Fixing variable no.3: When
## Fixing variable no.4: event
## Fixing variable no.5: time
## Fixing variable no.6: Gender

db_info(ADB)

##   source variable          label   class
## 1   ADB         id              factor
## 2   ADB         gr             Group factor
## 3   ADB         Age   Age at some time numeric
## 4   ADB         When              Date
## 5   ADB         event An event at some time factor
## 6   ADB         time   Time passed numeric
## 7   ADB         Gender   The Sex   factor
```

We can view, or get the information

```
## myDoc <- dm_doc()
dm_doc() ## only prints partial information in the doc
```

##	name	var	db	transf	label	comment
## 1	gr	gr	BL		Group	
## 2	Age	aalder	BL		Age at some time	
## 3	When	nar	BL			wtf?
## 4	event	foo	COMP		An event at some time	
## 5	time	bar	COMP	log	Time passed	
## 6	Gender	koon	BL		The Sex	

If we are using L^AT_EX, we can get the code for this with

```
dm_doc2latex(caption = "Variables and their origin.")
```

Table 1: Variables and their origin.

name	var	db	label	comment
gr	gr	BL	Group	
Age	aalder	BL	Age at some time	
When	nar	BL		<i>wtf?</i>
event	foo	COMP	An event at some time	
time	bar	COMP	Time passed	
Gender	koon	BL	The Sex	

and all recode-information with

```
dm_recode2latex()
```

Table 2: Recoding of data base entry **gr** into **gr**.

old ↓ new →	A	B	CD	Unknown
A	22	0	0	0
A2	31	0	0	0
B	0	27	0	0
C	0	0	28	0
D	0	0	25	0
d1	0	0	26	0
unknown	0	0	0	31
NA	0	0	0	10

Else, you can get the information from the 'print' of the `dm_doc()` and the recodings with, respectively,

Table 3: Recoding of data base entry `foo` into `event`.

old ↓ new →	No	Yes
0	148	0
1	0	32

Table 4: Recoding of data base entry `koon` into `Gender`.

old ↓ new →	Male	Female
K	0	83
M	117	0

```
d <- print(dm_doc(), print = FALSE)
lapply(dm_doc(), FUN = function(x) x$recode_table)
```


3 time_match

3.1 The problem we want to solve

Suppose you have a set of units in a data set, and another data set where each unit may occur none or several times, each row associated with a date and one or more variables that contains some kind of information you want to search. For each unit we want to find matches in this information within some specified time range.

The function was specifically written to deal with this situation: the units are selected to be part of some study cohort, possible with different start and end times. Another data set exists which contains the medical records of (some possibly larger) population. At least two tasks are commonly associated with creating an analytical data base:

- Find the medical history of each individual, i.e. look for codes pertaining to different diagnosis that appear before the individual is entered into the cohort.
- Find diagnosis that appear after the individual is entered into the study (and possibly before some end date).

3.2 An example

3.2.1 Generate data

Our cohort will consists of some individuals entering a study during the year 2010, with no longer than one year follow-up.

- Anna enters 2010-01-01, with no medical history or outcomes,
- Barry enter 2010-02-01, with a previous 'foo', but no outcome,
- Christina enters 2010-03-01, with no medical history but a later 'bar',
- David enters 2010-04-01, with a medical history of both 'bar' and 'foo', as well as a later 'quuz',
- Esteban enters 2010-05-01, with no medical history and an outcome 'bar' *after* the end of follow-up.

Moreover, the medical history is to be found in two different variables.

```
POP <- data.frame(  
  id = c('Anna', 'Barry', 'Christina',  
         'David', 'Esteban'),  
  enter = as.Date(c('2010-01-01', '2010-02-01', '2010-03-01',  
                   '2010-04-01', '2010-05-01'))  
)
```

```

RECORDS <- data.frame(
  identity = c('Barry', 'Christina', 'David',
               'David', 'David', 'Esteban',
               'Other', 'Other'),
  what1 = c('headache', 'bar type I', 'nausea',
             'percutaneous foo', 'quuz', '',
             'other foo', 'other bar'),
  what2 = c('mild foo', 'bar type II', 'severe bar',
             'subcutaneous foo', NA, 'bar-ish',
             'yet other foo', 'yet other bar'),
  what.date = as.Date(c('2010-01-07', '2010-07-23', '1998-06-27',
                        '1996-10-12', '2011-01-18', '2011-05-03',
                        '1999-12-01', '2010-06-01'))
)[sample(1:8),]
options('knitr.kable.NA' = '')

```

The data is tabulated below

```

POP

##      id      enter
## 1   Anna 2010-01-01
## 2   Barry 2010-02-01
## 3 Christina 2010-03-01
## 4   David 2010-04-01
## 5 Esteban 2010-05-01

RECORDS

##  identity      what1      what2  what.date
## 1   Barry      headache      mild foo 2010-01-07
## 4   David percutaneous foo subcutaneous foo 1996-10-12
## 5   David              quuz              <NA> 2011-01-18
## 8   Other      other bar      yet other bar 2010-06-01
## 3   David              nausea      severe bar 1998-06-27
## 6 Esteban              bar-ish 2011-05-03
## 2 Christina      bar type I      bar type II 2010-07-23
## 7   Other      other foo      yet other foo 1999-12-01

```

3.2.2 Medical history

Now we'll find the medical history of this cohort. We will need to point to the relevant variables in the different data sets, in `RECORDS` we need to point to `identity`, `date` and `what.date`. In `POP` we need to point to `id` and specify the search interval 'begin' and 'end'. In this case, we search as far back as we can,

which will happen if we set 'begin' to NULL. We'll search all the way until the beginning of the study (which is coded in `enter` in the data frame).

There are options for the output format, but typically we want a stacked long format (which will be default).

```
searchString <- c('Foo' = 'foo', 'Bar' = 'bar', 'Quuz' = 'quuz')

tm <- time_match(
  pattern = searchString, ## what to search for
  x = c('what1', 'what2'), ## search variables in 'data'
  data = RECORDS, ## data set to search in
  id = 'identity', ## name of id variable in 'data'
  date = 'what.date', ## name of date variable in 'data'
  units = POP, ## data set, or vector, containing individuals
  units.id = 'id', ## name of id variable in 'units'
  begin = NULL, ## earliest date to search from
  end = 'enter', ## name of latest date to search,
  ## long = TRUE, ## long output format is default
  ## stack = TRUE, ## stacked results are default
  verbose = FALSE ## give calculation progress info?
)
```

`time_match` will return a data frame with many variables and, with this configuration, at least one row per individual and search string, and possibly as many as one per search string times variable searched in. Output (names are fixed):

- `id` the identifier
- `begin` first date searched from (inclusive)
- `end` last date searched until (inclusive)
- `date` the date of the match
- `event` indicator for match
- `time` days between `begin` and `date`
- `match` that which matched
- `match.in` name of variable of match
- `pattern` pattern searched for
- `alias` name of pattern searched for
- `first.id` indicator for the first match for each individual and pattern
- `first.id_date` indicator for the first match for each individual, date and pattern

A few of these are tabulated below.

```
tm[, c('id', 'event', 'alias', 'match', 'match.in', 'first.id')]
```

```
id event alias match match.in first.id 1 Barry 1 Foo mild foo what2 1 2
David 1 Foo percutaneous foo what1 1 3 David 1 Foo subcutaneous foo what2
0 4 Christina 0 Foo ;NA; ;NA; 1 5 Esteban 0 Foo ;NA; ;NA; 1 6 Anna 0 Foo
;NA; ;NA; 1 7 David 1 Bar severe bar what2 1 8 Barry 0 Bar ;NA; ;NA; 1 9
Christina 0 Bar ;NA; ;NA; 1 10 Esteban 0 Bar ;NA; ;NA; 1 11 Anna 0 Bar
;NA; ;NA; 1 12 Barry 0 Quuz ;NA; ;NA; 1 13 Christina 0 Quuz ;NA; ;NA;
1 14 David 0 Quuz ;NA; ;NA; 1 15 Esteban 0 Quuz ;NA; ;NA; 1 16 Anna 0
Quuz ;NA; ;NA; 1
```

For the history, we typically only care whether at least one instance of each search term is found. Also, we might want to transform this to a wide format.

```
tmp <- subset(tm, first.id == 1, select = c('id', 'event', 'alias'))
(medhist <- reshape(tmp, idvar = 'id', timevar = c('alias'), direction = 'wide'))

##           id event.Foo event.Bar event.Quuz
## 1      Barry         1         0         0
## 2      David         1         1         0
## 4 Christina         0         0         0
## 5      Esteban         0         0         0
## 6        Anna         0         0         0

names(medhist) <- gsub("event", "prior", names(medhist), fixed = TRUE)
```

Now, we have a data frame containing the relevant medical history

```
medhist

##           id prior.Foo prior.Bar prior.Quuz
## 1      Barry         1         0         0
## 2      David         1         1         0
## 4 Christina         0         0         0
## 5      Esteban         0         0         0
## 6        Anna         0         0         0
```

3.2.3 Outcomes

Next, we'll look at outcomes. Since the end of study is variable, we'll have to create this variable, let's call it `endofstudy`. Note: if we want to start the search one day after `enter` we'll have to create this delayed start as a variable. Search dates are inclusive.

```
POP$endofstudy <- POP$enter + 365
tm2 <- time_match(pattern = searchString, x = c('what1', 'what2'),
  data = RECORDS, id = 'identity', date = 'what.date',
  units = POP, units.id = 'id',
  begin = 'enter', ## earliest date to search from
  end = 'endofstudy', ## name of latest date to search,
  verbose = FALSE)
```

For the outcomes, we probably care about more things, especially time-to-event. The event and time variables now serve as right-censored data for each outcome.

```
tm2[, c('id', 'event', 'time', 'alias', 'match', 'match.in')]

##           id event time alias      match match.in
## 1      Barry      0  365   Foo      <NA>      <NA>
## 2  Christina      0  365   Foo      <NA>      <NA>
## 3      David      0  365   Foo      <NA>      <NA>
## 4    Esteban      0  365   Foo      <NA>      <NA>
## 5       Anna      0  365   Foo      <NA>      <NA>
## 6  Christina      1  144   Bar bar type I  what1
## 7  Christina      1  144   Bar bar type II what2
## 8      Barry      0  365   Bar      <NA>      <NA>
## 9      David      0  365   Bar      <NA>      <NA>
## 10   Esteban      0  365   Bar      <NA>      <NA>
## 11      Anna      0  365   Bar      <NA>      <NA>
## 12   David      1  292  Quuz    quuz    what1
## 13   Barry      0  365  Quuz      <NA>      <NA>
## 14  Christina      0  365  Quuz      <NA>      <NA>
## 15   Esteban      0  365  Quuz      <NA>      <NA>
## 16      Anna      0  365  Quuz      <NA>      <NA>
```

We'll assume that we only care about the first instance of each outcome.

```
tmp2 <- subset(tm2, first.id == 1, select = c('id', 'event', 'time', 'alias'))
(outcomes <- reshape(tmp2, idvar = 'id', timevar = c('alias'), direction = 'wide'))

##           id event.Foo time.Foo event.Bar time.Bar event.Quuz time.Quuz
## 1      Barry      0      365      0      365      0      365
## 2  Christina      0      365      1      144      0      365
## 3      David      0      365      0      365      1      292
## 4    Esteban      0      365      0      365      0      365
## 5       Anna      0      365      0      365      0      365

names(outcomes) <- gsub("event", "ev", names(outcomes), fixed = TRUE)
names(outcomes) <- gsub("time", "t", names(outcomes), fixed = TRUE)
```

Now, we have a data frame containing the relevant medical history

outcomes							
##	id	ev.Foo	t.Foo	ev.Bar	t.Bar	ev.Quuz	t.Quuz
## 1	Barry	0	365	0	365	0	365
## 2	Christina	0	365	1	144	0	365
## 3	David	0	365	0	365	1	292
## 4	Esteban	0	365	0	365	0	365
## 5	Anna	0	365	0	365	0	365

3.3 Other output

I'm thinking... maybe not use these?

3.3.1 Wide and stacked

With a wide, stacked output, we get one row per individual and search. We get some information on the first match - all information from the long stacked format, except `first.id` and `first.id_date` - and some summary information on all matches:

- `events` which counts the matches,
- `matches` which concatenates the matches, and
- `matches.info` which stores a concatenation of *match:math.in:date* for all matches.

```
tm3 <- time_match(pattern = searchString, x = c('what1', 'what2'),
  data = RECORDS, id = 'identity', date = 'what.date',
  units = POP, units.id = 'id', begin = 'enter',
  end = 'endofstudy',
  long = FALSE, ## wide output format
  stack = TRUE, ## stack
  verbose = FALSE
)
str(tm3)

## 'data.frame': 15 obs. of 13 variables:
## $ id : Factor w/ 5 levels "Anna","Barry",...: 1 2 3 4 5 1 2 3 4 5 ...
## $ begin : Date, format: "2010-01-01" "2010-02-01" ...
## $ end : Date, format: "2011-01-01" "2011-02-01" ...
## $ date : Date, format: "2011-01-01" "2011-02-01" ...
## $ event : num 0 0 0 0 0 0 0 1 0 0 ...
## $ time : num 365 365 365 365 365 365 365 144 365 365 ...
## $ match : chr NA NA NA NA ...
## $ match.in : chr NA NA NA NA ...
## $ pattern : chr "foo" "foo" "foo" "foo" ...
## $ alias : chr "Foo" "Foo" "Foo" "Foo" ...
## $ events : num 0 0 0 0 0 0 0 2 0 0 ...
## $ matches : chr NA NA NA NA ...
## $ matches.info: chr NA NA NA NA ...
```

Also, selected info tabulated below.

```
val <- c('id', 'alias', 'event', 'time', 'events', 'matches.info')
tm3[, val]
```

##		id	alias	event	time	events	
## 1		Anna	Foo	0	365	0	
## 2		Barry	Foo	0	365	0	
## 3		Christina	Foo	0	365	0	
## 4		David	Foo	0	365	0	
## 5		Esteban	Foo	0	365	0	
## 6		Anna	Bar	0	365	0	
## 7		Barry	Bar	0	365	0	
## 8		Christina	Bar	1	144	2	
## 9		David	Bar	0	365	0	
## 10		Esteban	Bar	0	365	0	
## 11		Anna	Quuz	0	365	0	
## 12		Barry	Quuz	0	365	0	
## 13		Christina	Quuz	0	365	0	
## 14		David	Quuz	1	292	1	
## 15		Esteban	Quuz	0	365	0	
##							matches.info
## 1							<NA>
## 2							<NA>
## 3							<NA>
## 4							<NA>
## 5							<NA>
## 6							<NA>
## 7							<NA>
## 8	bar type I:	what1:2010-07-23	bar type II:	what2:2010-07-23			
## 9							<NA>
## 10							<NA>
## 11							<NA>
## 12							<NA>
## 13							<NA>
## 14						quuz:what1:2011-01-18	
## 15							<NA>

3.3.2 Wide and unstacked

With a wide and unstacked output, we get all variables (from the wide stacked output) *for each search term* - except `id`, `begin` and `end` which are the same for all rows - with the name of the search term as suffix (or a naming scheme if no names are supplied).

Below you can see the structure

```
tm4 <- time_match(pattern = searchString, x = c('what1', 'what2'),
  data = RECORDS, id = 'identity', date = 'what.date',
  units = POP, units.id = 'id', begin = 'enter',
  end = 'endofstudy',
  long = FALSE, ## wide output format
  stack = FALSE, ## don't stack
  verbose = FALSE
)
str(tm4)

## 'data.frame': 5 obs. of 33 variables:
## $ id : Factor w/ 5 levels "Anna","Barry",...: 1 2 3 4 5
## $ begin : Date, format: "2010-01-01" "2010-02-01" ...
## $ end : Date, format: "2011-01-01" "2011-02-01" ...
## $ date.Foo : Date, format: "2011-01-01" "2011-02-01" ...
## $ event.Foo : num 0 0 0 0 0
## $ time.Foo : num 365 365 365 365 365
## $ match.Foo : logi NA NA NA NA NA
## $ match.in.Foo : logi NA NA NA NA NA
## $ pattern.Foo : chr "foo" "foo" "foo" "foo" ...
## $ alias.Foo : chr "Foo" "Foo" "Foo" "Foo" ...
## $ events.Foo : num 0 0 0 0 0
## $ matches.Foo : logi NA NA NA NA NA
## $ matches.info.Foo : logi NA NA NA NA NA
## $ date.Bar : Date, format: "2011-01-01" "2011-02-01" ...
## $ event.Bar : num 0 0 1 0 0
## $ time.Bar : num 365 365 144 365 365
## $ match.Bar : Factor w/ 15 levels "", "bar type I",...: NA NA 2 NA NA
## $ match.in.Bar : Factor w/ 2 levels "what1","what2": NA NA 1 NA NA
## $ pattern.Bar : chr "bar" "bar" "bar" "bar" ...
## $ alias.Bar : chr "Bar" "Bar" "Bar" "Bar" ...
## $ events.Bar : num 0 0 2 0 0
## $ matches.Bar : chr NA NA "bar type I bar type II" NA ...
## $ matches.info.Bar : chr NA NA "bar type I:what1:2010-07-23 bar type II:what2:2010-07-23" ...
## $ date.Quuz : Date, format: "2011-01-01" "2011-02-01" ...
## $ event.Quuz : num 0 0 0 1 0
## $ time.Quuz : num 365 365 365 292 365
## $ match.Quuz : Factor w/ 8 levels "", "bar type I",...: NA NA NA 8 NA
```

```
## $ match.in.Quuz : Factor w/ 2 levels "what1","what2": NA NA NA 1 NA
## $ pattern.Quuz : chr "quuz" "quuz" "quuz" "quuz" ...
## $ alias.Quuz : chr "Quuz" "Quuz" "Quuz" "Quuz" ...
## $ events.Quuz : num 0 0 0 1 0
## $ matches.Quuz : chr NA NA NA "quuz" ...
## $ matches.info.Quuz: chr NA NA NA "quuz:what1:2011-01-18" ...
```

Also, selected info on the Bar- and Quuz outcome tabulated.

```
val <- c('id', names(tm4)[grepl("event|time", names(tm4))])
tm4[, val[!grepl("Foo", val)]]
```

	id	event.Bar	time.Bar	events.Bar	event.Quuz	time.Quuz	events.Quuz
## 1	Anna	0	365	0	0	365	0
## 2	Barry	0	365	0	0	365	0
## 3	Christina	1	144	2	0	365	0
## 4	David	0	365	0	1	292	1
## 5	Esteban	0	365	0	0	365	0

4 Other functions

4.1 cdate

`cdate` is a function to handle dates of the form “20010700” or “20010000”, which can appear as dates of death when the precise date is unknown. If nothing else is known this function will replace an unknown

- day of the month with the midpoint of that month, and
- month (and day) with the midpoint of that year.

Sometimes, there is another date for an individual when the individual was known to be alive. In the applications of most interest to the author it is an admission, or discharge, date to a hospital, and thus we believe that the individual did *not* die at the hospital (else the date of death would be known), therefore the replacement date will be the midpoint of whatever remains of the unknown period, e.g if we encounter date of death as “20010100” with a known hospital discharge at 2001-01-21, we will interpret it as 2001-01-26.

Examples:

```
cdate(x = c("20010101", "20010100", "20010000"))  
## [1] "2001-01-01" "2001-01-16" "2001-07-02"  
  
cdate(x = c("20010101", "20010100", "20010000"),  
      low.bound = as.Date(c("1999-01-01", "2001-01-21", "2001-06-20")))  
## [1] "2001-01-01" "2001-01-26" "2001-09-25"
```