Package 'maic'

October 13, 2022

Type Package

Title Matching-Adjusted Indirect Comparison

Version 0.1.4

Date 2022-04-27

Maintainer Rob Young <rob.young@heor.co.uk>

Depends R (>= 3.0.0)

Description A generalised workflow for generation of subject weights to be used in Matching-Adjusted Indirect Comparison (MAIC) per Signorovitch et al. (2012) <doi:10.1016/j.jval.2012.05.004>, Signorovitch et al (2010) <doi:10.2165/11538370-000000000-00000>. In MAIC, unbiased comparison between outcomes of two trials is facilitated by weighting the subject-level outcomes of one trial with weights derived such that the weighted aggregate measures of the prognostic or effect modifying variables are equal to those of the sample in the comparator trial. The functions and classes included in this package wrap and abstract the process demonstrated in the UK National Institute for Health and Care Excellence Decision Support Unit (NICE DSU)'s example (Phillippo et al, (2016) [see URL]), providing a repeatable and easily specifiable workflow for producing multiple comparison variable sets against a variety of target studies, with preprocessing for a number of aggregate target forms (e.g. mean, median, domain limits).

License GPL-3

Encoding UTF-8

URL https://github.com/heorltd/maic,

https://nicedsu.sites.sheffield.ac.uk/tsds/
population-adjusted-indirect-comparisons-maic-and-stc

Imports Hmisc, matrixStats, weights

RoxygenNote 7.1.2

Suggests testthat

NeedsCompilation no

Author Rob Young [aut, cre]

Repository CRAN Date/Publication 2022-04-27 14:50:07 UTC

R topics documented:

createMAICInput	2
maicMatching	4
maicWeight	6
reportCovariates	7
	10

Index

createMAICInput Construct a MAIC input matrix

Description

From index patient level data and a set of target baseline characteristics, construct the input matrix to the maic.

Usage

```
createMAICInput(index, target, dictionary, matching.variables, x = FALSE)
```

Arguments

index	A matrix or data.frame containing patient-level data			
target	A list containing target summary data			
dictionary	A data frame containing the columns "match.id", "target.variable", "index.variable" and "match.type"			
matching.variables				
	A character vector indicating the match.id to use			
х	Return subject level inputs?			

Details

The dictionary is a data frame containing at least 4 vectors:

- "match.id" the name of the match, used to refer to it in the matching.variables list
- "target.variable" the name of the variable in the target values list use to inform the matching. Use dependent on type
- "index.variable" the name of the variable in the index data frame to match on.
- "match.type" A string indicating the match type to use. The following values are accepted:
 - minimum records with index values lower than the target variable will be assigned 0 weight

- maximum records with index values greater than the target variable will be assigned 0 weight
- median records with index values greater than the target variable will be assigned a value of 1, those lower 0. The target for matching will be a mean of 0.5
- quantile.X Generalisation of the median code. records with index values greater than the target variable will be assigned a value of 1, those lower 0. The target for matching will be a mean of 0.X
- mean records will match index value directly onto target value
- proportion as mean, with index values encoded as 1 = true, 0 = false. If target proportion
 is exclusive (0 or 1 exactly) then excluded members of the index population shall receive
 no weighting.
- sd a matching on the square of the index value on the sum of the square of the target mean and target standard deviation. The target mean is provided by the "supplementary.target.variable"
- var a matching on the square of the index value on the sum of the square of the target mean and the variance specified by the target variable. The target mean is provided by the "supplementary.target.variable"

In addition, the following vector may be necessary:

• "supplementary.target.variable" - The name of the variable in the target values list that provides e.g. the mean for sd and var matching.

and, for estimating some p-values on difference (e.g. in proportion)

• "sample.size.variable" - The name of the variable in the target values list that provides the number of subjects in the sample. Only for reporting.

It is possible to use these match types to match on other variables by pre-processing the input correctly.

Finally, the matching.variables is a list or character vector containing match.ids to be acted upon in this MAIC.

Value

An object of class maic.input

Examples

```
target <- c("Air.Flow" = 60,
                "Water.Temp" = 21,
                "Prop.Acid.Conc.LT.90" = 0.7,
                "min.air.flow" = 55)
stackloss$match.conc.lt.90 <-
    ifelse(stackloss$Acid.Conc. < 90, 1, 0)
dict <- data.frame(
               "match.id" =
                c("airflow", "watertemp",
```

```
"acidconc", "min.airflow"),
  "target.variable" =
    c("Air.Flow", "Water.Temp",
      "Prop.Acid.Conc.LT.90", "min.air.flow"),
  "index.variable" =
    c("Air.Flow", "Water.Temp",
      "match.conc.lt.90", "Air.Flow"),
  "match.type" =
    c("mean", "mean", "proportion", "min"),
  stringsAsFactors = FALSE)
ipmat <- createMAICInput(</pre>
  index = stackloss,
  target = target,
  dictionary = dict,
  matching.variables =
    c("airflow", "watertemp",
      "acidconc", "min.airflow"))
wts <- maicWeight(ipmat)</pre>
rcv <- reportCovariates(</pre>
  stackloss, target, dict,
  matching.variables =
    c("airflow", "watertemp",
    "acidconc", "min.airflow"),
  wts)
```

maicMatching calculate MAIC weights

Description

From index patient level data and a set of target baseline characteristics, calculate MAIC weights.

Usage

```
maicMatching(
    index,
    target,
    dictionary,
    matching.variables,
    reporting.variables = NULL
)
```

Arguments

index	A matrix or data.frame containing patient-level data
target	A list containing target summary data

4

maicMatching

dictionary	A data frame containing the columns "match.id", "target.variable", "index.variable"		
	and "match.type"		
matching.variables			
	A character vector indicating the match.id to use		
reporting.variables			
	A optional character vector of matches to report upon (defaults to matching.variables)		

Details

The dictionary is a data frame containing at least 4 vectors:

- "match.id" the name of the match, used to refer to it in the matching.variables list
- "target.variable" the name of the variable in the target values list use to inform the matching. Use dependent on type
- "index.variable" the name of the variable in the index data frame to match on.
- "match.type" A string indicating the match type to use. The following values are accepted:
 - minimum records with index values lower than the target variable will be discarded
 - maximum records with index values greater than the target variable will be discarded
 - median records with index values greater than the target variable will be assigned a value of 1, those lower 0, and those equal 0.5. The target for matching will be a mean of 0.5
 - quantile.X Generalisation of the median code. records with index values greater than the target variable will be assigned a value of 1, those lower 0. The target for matching will be a mean of 0.X
 - mean records will match index value directly onto target value
 - proportion as mean, with index values encoded as 1 = true, 0 = false. If target proportion is exclusive (0 or 1 exactly) then excluded members of the index population shall receive no weighting.
 - sd a matching on the square of the index value on the sum of the square of the target mean and target standard deviation. The target mean is provided by the "supplementary.target.variable"
 - var a matching on the square of the index value on the sum of the square of the target mean and the variance specified by the target variable. The target mean is provided by the "supplementary.target.variable"

In addition, the following vector may be necessary:

• "supplementary.target.variable" - The name of the variable in the target values list that provides e.g. the mean for sd and var matching.

It is possible to use these match types to match on other variables, e.g. variance, by pre-processing the input correctly.

Finally, the matching.variables is a list or character vector containing match.ids to be acted upon in this MAIC.

Value

An object of class MaicAnalysis, with components weights and aggregate, containing the weights vector and the covariate aggregate data respectively

Examples

```
target <- c("Air.Flow" = 60,</pre>
             "Water.Temp" = 21,
             "Prop.Acid.Conc.LT.90" = 0.7,
             "min.air.flow" = 55)
stackloss$match.conc.lt.90 <-</pre>
  ifelse(stackloss$Acid.Conc. < 90, 1, 0)</pre>
dict <- data.frame(</pre>
  "match.id" =
    c("airflow", "watertemp",
    "acidconc", "min.airflow"),
  "target.variable" =
    c("Air.Flow", "Water.Temp",
      "Prop.Acid.Conc.LT.90", "min.air.flow"),
  "index.variable" =
    c("Air.Flow", "Water.Temp",
      "match.conc.lt.90", "Air.Flow"),
  "match.type" =
    c("mean", "mean", "proportion", "min"),
  stringsAsFactors = FALSE)
weightObj <- maicMatching(</pre>
  index = stackloss,
  target = target,
  dictionary = dict,
  matching.variables =
    c("airflow", "watertemp",
      "acidconc", "min.airflow"))
```

maicWeight

Calculate MAIC weights

Description

This function calculates the weights to apply to records for Matching-Adjusted Indirect Comparison (MAIC), from either a raw input matrix or a maic.input object

Usage

maicWeight(x, opt = TRUE, keep.x = TRUE, ...)

Arguments

х	Either a maic.input object or a MAIC input matrix
opt	return the optim object as attribute
keep.x	return the input matrix as an attribute
	Optional arguments to optim

6

reportCovariates

Value

A numeric vector of weights corresponding to the rows in the input matrix

Examples

```
target <- c("Air.Flow" = 60,</pre>
           "Water.Temp" = 21,
           "Prop.Acid.Conc.LT.90" = 0.7,
           "min.air.flow" = 55)
stackloss$match.conc.lt.90 <-</pre>
 ifelse(stackloss$Acid.Conc. < 90, 1, 0)</pre>
dict <- data.frame(</pre>
  "match.id" =
   c("airflow", "watertemp",
      "acidconc", "min.airflow"),
  "target.variable" =
    c("Air.Flow", "Water.Temp",
      "Prop.Acid.Conc.LT.90", "min.air.flow"),
  "index.variable" =
    c("Air.Flow", "Water.Temp",
      "match.conc.lt.90", "Air.Flow"),
  "match.type" =
    c("mean", "mean", "proportion", "min"),
 stringsAsFactors = FALSE)
ipmat <- createMAICInput(</pre>
 index = stackloss,
 target = target,
 dictionary = dict,
 matching.variables =
   c("airflow", "watertemp",
      "acidconc", "min.airflow"))
wts <- maicWeight(ipmat)</pre>
rcv <- reportCovariates(</pre>
 stackloss, target, dict,
 matching.variables =
   c("airflow", "watertemp",
      "acidconc", "min.airflow"),
 wts)
```

reportCovariates Calculate the rebalanced covariates

Description

This function calculates the raw, target and achieved covariates given a set of weights. Note that for mean values, bootstrapped standard errors are used and so downstream values (such as p-values for difference) may differ from run to run if the random number stream is not consistent

Usage

```
reportCovariates(
    index,
    target,
    dictionary,
    matching.variables,
    weights,
    tidy = TRUE,
    var.method = c("ML", "unbiased")
)
```

Arguments

index	A matrix or data.frame containing patient-level data			
target	A list containing target summary data			
dictionary	A data frame containing the columns "match.id", "target.variable", "index.variable" and "match.type"			
matching.variables				
	A character vector indicating the match.id to use			
weights	A numeric vector with weights corresponding to the index data rows			
tidy	A boolean - return as a data frame (otherwise list)			
var.method	Estimator type passed through to wtd.var. Defaults to ML, as Bessel's correction not used in weights generation.			

Value

An object of class maic.covariates

Examples

```
target <- c("Air.Flow" = 60,
                "Water.Temp" = 21,
                "Prop.Acid.Conc.LT.90" = 0.7,
                "min.air.flow" = 55)
stackloss$match.conc.lt.90 <-
    ifelse(stackloss$Acid.Conc. < 90, 1, 0)
dict <- data.frame(
                "match.id" =
                c("airflow", "watertemp",
                "acidconc", "min.airflow"),
```

reportCovariates

```
"target.variable" =
    c("Air.Flow", "Water.Temp",
      "Prop.Acid.Conc.LT.90", "min.air.flow"),
  "index.variable" =
    c("Air.Flow", "Water.Temp",
      "match.conc.lt.90", "Air.Flow"),
  "match.type" =
    c("mean", "mean", "proportion", "min"),
  stringsAsFactors = FALSE)
ipmat <- createMAICInput(</pre>
  index = stackloss,
  target = target,
  dictionary = dict,
  matching.variables =
    c("airflow", "watertemp",
    "acidconc", "min.airflow"))
wts <- maicWeight(ipmat)</pre>
rcv <- reportCovariates(</pre>
  stackloss, target, dict,
  matching.variables =
    c("airflow", "watertemp",
    "acidconc", "min.airflow"),
  wts)
```

Index

createMAICInput, 2

maicMatching, 4
maicWeight, 6

optim, <mark>6</mark>

reportCovariates, 7

wtd.var,<mark>8</mark>