

Package ‘medAMM’

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Type Package

Title Mediation Analysis with Multiple Mediators using a Weighted Approach

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Description Provides functions to calculate the proportion of the exposure effect explained by individual mediators, as well as by all the mediators together. Only appropriate for a binary outcome and a binary exposure; mediators and confounders may be binary and/or continuous. Implements the weighting approach of VanderWeele TJ, Vansteelandt S (2014). “Mediation Analysis with Multiple Mediators.” *Epidemiol Methods*. 2(1):95-115.

License GPL

Imports survey, ggplot2, stats

NeedsCompilation no

Depends R (>= 3.5.0)

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causal.calc.all	<i>Main function for causal mediation analysis</i>
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Description

Main function for causal mediation analysis. Calculates the proportion of the exposure effect explained by each given mediator, as well as by all the mediators together. Implements the weighting approach of VanderWeele TJ, Vansteelandt S. *Mediation Analysis with Multiple Mediators*. *Epidemiol Methods*. 2014;2(1):95-115. doi:10.1515/em-2012-0010.

Usage

```
causal.calc.all(datas, confounders, outcome, mainpred, mediator.names,
survey = FALSE, survey.diagnose = FALSE, conf = TRUE, plot = TRUE, plot.labels
= NULL, plot.order = NULL)
```

Arguments

<code>datas</code>	dataset; if <code>survey=TRUE</code> this must be a <code>svydesign</code> object
<code>confounders</code>	names of confounders, should be provided as a vector of strings that corresponds to the variable names in <code>datas</code> e.g. <code>c("age","sex")</code>
<code>outcome</code>	names of outcome, should be provided as a string that corresponds to the variable name in <code>datas</code>
<code>mainpred</code>	names of main predictor or exposure, should be provided as a string that corresponds to the variable name in <code>datas</code>
<code>mediator.names</code>	names of mediators, should be provided as a vector of strings that corresponds to the variable names in <code>datas</code> e.g. <code>c("income","health")</code>
<code>survey</code>	TRUE or FALSE, if the data is a <code>svydesign</code> object; default is FALSE
<code>survey.diagnose</code>	TRUE OR FALSE, if the user would like a summary of the interim weights printed to diagnose possible extreme weights; default is FALSE
<code>conf</code>	TRUE OR FALSE, if confidence intervals are wanted, which are obtained via bootstrapping and are computationally intensive; default is TRUE
<code>plot</code>	TRUE OR FALSE, if plot is wanted, which requires confidence intervals obtained via bootstrapping and is computationally intensive; default is TRUE
<code>plot.labels</code>	a vector of strings for the labels for the mediators in the plot; if not supplied then the variable names are used
<code>plot.order</code>	a vector of numbers specifying the order of the plot

Value

If `plot = TRUE`, a plot of returned. In addition, a list is returned:

<code>results</code>	A formatted table of results
<code>values</code>	A table of values used for plotting; will be empty if <code>plot=FALSE</code>

Author(s)

Layla Parast

References

VanderWeele TJ, Vansteelandt S. Mediation Analysis with Multiple Mediators. *Epidemiol Methods*. 2014;2(1):95-115. doi:10.1515/em-2012-0010

Examples

```
data(datamed)

causal.calc.all(datas = datamed, confounders = c("con1","con2"), outcome = "outcome",
mainpred = "exposure", mediator.names = c("med1","med2","med3"),conf=FALSE, plot=FALSE)

#computationally intensive
## Not run: causal.calc.all(datas = datamed, confounders = c("con1","con2"),
outcome = "outcome", mainpred = "exposure", mediator.names = c("med1","med2",
"med3"),plot=TRUE)
## End(Not run)
```

`causal.calc.plot.only` *Function to plot causal mediation results*

Description

Function to plot causal mediation results; must be run after `causal.calc.all` and use the output from `causal.calc.all` where `plot` was set to `TRUE` so that the values are returned. The purpose of this function is to allow the user to make some edits to the plot, without having to re-run all the mediation analyses.

Usage

```
causal.calc.plot.only(values, mediator.names, plot.labels = NULL, plot.order = NULL)
```

Arguments

<code>values</code>	values to be used for plotting which come from <code>causal.calc.all</code>
<code>mediator.names</code>	names of mediators, should be provided as a vector of strings that corresponds to the variable names in <code>datas</code> e.g. <code>c("income","health")</code>
<code>plot.labels</code>	a vector of strings for the labels for the mediators in the plot; if not supplied then the variable names are used
<code>plot.order</code>	a vector of numbers specifying the order of the plot

Value

returns a plot

Author(s)

Layla Parast

Examples

```
data(datamed)

output = causal.calc.all(datas = datamed, confounders = c("con1","con2"), outcome = "outcome",
mainpred = "exposure", mediator.names = c("med1","med2","med3"),plot=TRUE)

causal.calc.plot.only(output$values, mediator.names = c("med1","med2","med3"),
plot.labels = c("Employed", "Pollution exposure","Prior arrest"), plot.order = c(2,3,4,1))
```

datamed

Example data

Description

Example data; simulated

Usage

```
data("datamed")
```

Format

A data frame with 5000 observations on the following 7 variables.

outcome the binary outcome, a numeric vector

exposure the binary exposure, a numeric vector

med1 mediator 1, a numeric vector

med2 mediator 2, a numeric vector

med3 mediator 3, a numeric vector

con1 confounder 1, a numeric vector

con2 confounder 2, a numeric vector

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