

Package ‘multmod’

October 20, 2009

Title Testing of multiple outcomes

LazyLoad yes

LazyData yes

Version 0.6

Author Christian B. Pipper <pipper@life.ku.dk>, Christian Ritz <ritz@life.ku.dk>

Maintainer Christian B. Pipper <pipper@life.ku.dk>

Description Testing of multiple outcomes using i.i.d. decompositions

License GPL (>= 2)

Depends R (>= 2.6.0), gtools, mvtnorm, sandwich, MASS

Date 2009-08-31

URL <http://www.r-project.org>

Repository CRAN

Date/Publication 2009-10-20 09:33:27

R topics documented:

amtest	2
mn6.9	4
Index	5

amtest

*Testing multiple outcomes***Description**

Testing multiple endpoints or outcomes measures by providing a corrected significance level to assess the uncorrected, marginal p-values against.

Usage

```
amtest(modelList, varName, vcov. = c("sandwich", "model-based"),
       sig.level = 0.05, display = TRUE, adjp=FALSE)
```

Arguments

<code>modelList</code>	list of model fits. The fits may be obtained using <code>lm</code> , <code>glm</code> , or <code>coxph</code> . The fits should all be based on exactly the same dataset (data frame) and missing values should be handled using <code>na.omit</code> . All models should contain the explanatory variable of interest.
<code>varName</code>	string specifying the name of the explanatory variable for which the effect is to be assessed.
<code>vcov.</code>	string specifying the type of estimated variance-covariance matrix to use for calculating the test statistics. Two options are available model-based estimated standard errors or robust estimated sandwich standard errors, which is the default.
<code>sig.level</code>	numeric specifying the nominal significance level or type I error rate (default is 0.05).
<code>display</code>	logical indicating whether or not the results should be shown at the command line.
<code>adjp</code>	logical indicating whether or not the correction should be reported in terms of adjusted p-values.

Details

The function calculates a corrected overall significance level that asymptotically reaches the desired type I error using i.i.d. decompositions of parameter estimates by means of score components. The details are provided by Pipper and Ritz (2009).

The implemented method is less conservative than the practically identical Bonferroni/Slepian corrections and it flexibly adapts to the correlation structure between outcomes. Different types of outcome measures such as continuous, binomial, and event times, as well as missing values in outcome measures and explanatory variables can be handled by the procedure.

Value

The corrected significance level as well as the Slepian correction $1 - (1 - \text{Nominal significance level})^{(1/\text{Number of tests})}$ and the marginal p-values are shown by default (can be switched off using the argument 'display').

These values are also returned invisibly in a list.

Author(s)

Christian B. Pipper and Christian Ritz

References

Pipper, C. B. and Ritz, C (2009) An asymptotic correction for controlling the overall type I error when testing multiple outcomes, *Submitted manuscript*

Examples

```
##Example from: McCullagh, P. and Nelder, J.A. (1989,p. 239). Generalized Linear Models.  
##Second Edition. Chapman & Hall/CRC.
```

```
model1 <- glm(y1~group, family=binomial,na.action=na.omit,data=mn6.9)
```

```
model2 <- glm(y2~group, family=binomial,na.action=na.omit,data=mn6.9)
```

```
model3 <- glm(y3~group, family=binomial,na.action=na.omit,data=mn6.9)
```

```
model4 <- glm(y4~group, family=binomial,na.action=na.omit,data=mn6.9)
```

```
amtest(list(model1,model2,model3,model4),"group2",vcov="model-based")
```

```
amtest(list(model1,model2,model3,model4),"group2")
```

```
##Multiple testing in case of perfectly correlated outcomes
```

```
amtest(list(model2,model2),"group2",vcov="model-based")
```

```
amtest(list(model2,model2),"group2")
```

```
## Multiple testing in a subgroup analysis and overall analysis
```

```
subgroup<-mn6.9
```

```
set.seed(2982)
```

```
subgroup[sample(1:2982,1500),]<-NA
```

```
model1<- glm(y2~group, family=binomial,na.action=na.omit,data=mn6.9)
```

```
model2<- glm(y2~group, family=binomial,na.action=na.omit,data=subgroup)
```

```
amtest(list(model1,model2),"group2",vcov="model-based")
```

```
amtest(list(model1,model2),"group2")
```

`mn6.9`*I.Q. and attitude towards science*

Description

Responses given by 2982 New Jersey high-school seniors on 4 questions concerning attitude towards science. Also recorded was whether students had a high or low I.Q.

Usage

```
data(mn6.9)
```

Format

A data frame with 2982 observations on the following 5 variables.

y1 Agree=1/disagree=0 to "The development of new ideas is the scientist's greatest source of satisfaction"

y2 Agree=1/disagree=0 to "Scientists and engineers should be eliminated from the military draft"

y3 Agree=1/disagree=0 to "The scientist will make his maximum contribution to society when he has freedom to work on problems that interest him"

y4 Agree=1/disagree=0 to "The monetary compensation of a Nobel Prize-winner in physics should be at least equal to that given to popular entertainers"

group I.Q. levels: 1=low, 2=high

Source

McCullagh, P. and Nelder, J.A. (1989, p. 239). *Generalized Linear Models*. Second Edition. Chapman & Hall/CRC.

Examples

```
##Marginal assessment of effect of I.Q. group on response probability
model1 <- glm(y1~group, family=binomial,data=mn6.9)
model2 <- glm(y2~group, family=binomial,data=mn6.9)
model3 <- glm(y3~group, family=binomial,data=mn6.9)
model4 <- glm(y4~group, family=binomial,data=mn6.9)
```

Index

*Topic **datasets**

mn6.9, 4

*Topic **htest**

amtest, 2

*Topic **models**

amtest, 2

amtest, 2

coxph, 2

glm, 2

lm, 2

mn6.9, 4

na.omit, 2