# Package ‘multiCA’

August 7, 2017

<table>
<thead>
<tr>
<th>Type</th>
<th>Package</th>
</tr>
</thead>
<tbody>
<tr>
<td>Title</td>
<td>Multinomial Cochran-Armitage Trend Test</td>
</tr>
<tr>
<td>Version</td>
<td>1.1</td>
</tr>
<tr>
<td>Date</td>
<td>2017-08-07</td>
</tr>
<tr>
<td>Author</td>
<td>Aniko Szabo</td>
</tr>
<tr>
<td>Depends</td>
<td>R(&gt;= 2.10)</td>
</tr>
<tr>
<td>Imports</td>
<td>bitops, multcomp</td>
</tr>
<tr>
<td>Maintainer</td>
<td>Aniko Szabo <a href="mailto:aszabo@mcw.edu">aszabo@mcw.edu</a></td>
</tr>
</tbody>
</table>

**Description**
Implementes a generalization of the Cochran-Armitage trend test to multinomial data. In addition to an overall test, multiple testing adjusted p-values for trend in individual outcomes and power calculation is available.

**License**
GPL (>= 2)

**RoxygenNote**
5.0.1

**Suggests**
testthat

**Repository**
CRAN

**Repository/R-Forge/Project**
corrbin

**Repository/R-Forge/Revision**
106

**Repository/R-Forge/DateTimeStamp**
2017-08-07 16:05:30

**Date/Publication**
2017-08-07 19:36:33 UTC

**NeedsCompilation**
no

## R topics documented:

- multiCA-package
  - cnonct
  - multiCA.test
  - power.multiCA.test
  - stroke

Index

1
multiCA-package

Multinomial Cochran-Armitage Trend Test

Description

Implements a generalization of the Cochran-Armitage trend test to multinomial data. In addition to an overall test, multiple testing adjusted p-values for trend in individual outcomes and power calculation is available.

Details

The DESCRIPTION file:

Package: multiCA
Type: Package
Title: Multinomial Cochran-Armitage Trend Test
Version: 1.1
Date: 2017-08-07
Author: Aniko Szabo
Depends: R(>= 2.10)
Imports: bitops, multcomp
Maintainer: Aniko Szabo <aszabo@mcw.edu>
Description: Implements a generalization of the Cochran-Armitage trend test to multinomial data. In addition to an overall test, multiple testing adjusted p-values for trend in individual outcomes and power calculation is available.
License: GPL (>=2)
RoxygenNote: 5.0.1
Suggests: testthat
Repository: R-Forge
Repository/R-Forge/Project: corrbin
Repository/R-Forge/Revision: 106
Repository/R-Forge/DateTimeStamp: 2017-08-07 16:05:30
Date/Publication: 2017-08-07 16:05:30

Index of help topics:

cnonct Non-centrality parameter for chi-square distribution
multiCA-package Multinomial Cochran-Armitage Trend Test
multiCA.test Multinomial Cochran-Armitage trend test
power.multiCA.test Power calculations for the multinomial Cochran-Armitage trend test
stroke Stroke types over time

The main functionality is implemented in the multiCA.test function. The power.multiCA.test function can be used for power and sample size calculation.
Author(s)

Aniko Szabo

Maintainer: Aniko Szabo <aszabo@mcw.edu>

References

Szabo, A. (2016) Test for trend with a multinomial outcome.

Description

Calculates the non-centrality parameter for a chi-square distribution for a given quantile. This is often needed for sample size calculation for chi-square based tests.

Usage

cnonct(x, p, df)

Arguments

x a numeric value at which the distribution was evaluated
p a numeric value giving the cumulative probability at x
df an integer giving the degrees of freedom of the chi-square variable

Details

The function is modeled after the SAS function CNONCT. If p is larger than the cumulative probability of the central chi-square distribution at x, then there is no solution and NA is returned.

Examples

(ncp <- cnonct(qchisq(0.95, df=10), 0.8, df=10))
## check
pchisq(qchisq(0.95, df=10), df=10, ncp=ncp) ## 0.8
multiCA.test  

Multinomial Cochran-Armitage trend test

Description

The `multiCA.test` function performs a multinomial generalization of the Cochran-Armitage trend test.

Usage

```r
multiCA.test(x, ...)  

## Default S3 method:
multiCA.test(x, scores = 1:ncol(x), outcomes = 1:nrow(x),
   "Westfall"), ...)  

## S3 method for class 'formula'
multiCA.test(formula, data, subset, na.action, weights, ...)
```

Arguments

- `x`  
a two-dimensional matrix of event counts with the outcomes as rows and ordered groups as columns.
- `scores`  
non-decreasing numeric vector of the same length as the number of ordered groups. Defaults to linearly increasing values.
- `outcomes`  
integer or character vector defining the set of outcomes (by row index or row name) over which the trend should be tested. Defaults to all outcomes.
- `p.adjust.method`  
character string defining the correction method for individual outcome p-values. Defaults to "closed.set" when length(outcomes)<=3, and "Holm-Shaffer" otherwise.
- `formula`  
a formula of the form `outcome ~ group` where `outcome` is a factor representing the categorical outcome and `group` is the grouping variable over which the trend is tested.
- `data`  
an optional matrix or data frame containing the variables in the formula `formula`. By default the variables are taken from `environment(formula)`. 
- `subset`  
an optional vector specifying a subset of observations to be used.
- `na.action`  
a function which indicates what should happen when the data contain NAs. Defaults to `getOption("na.action")`. 
- `weights`  
an integer-valued variable representing the number of times each outcome-group combination was observed.
- `...`  
other arguments
Value

- a list with two components
  - overall: an object of class "htest" with the results of the overall test
  - individual: a vector with adjusted p-values for individual outcomes

Author(s)

Aniko Szabo

References

Szabo, A. (2016) Test for trend with a multinomial outcome.

Examples

data(stroke)
## using formula interface
multiCA.test(Type ~ Year, weights=Freq, data=stroke)

## using Westfall's multiple testing adjustment
multiCA.test(Type ~ Year, weights=Freq, data=stroke, p.adjust.method="Westfall")

## using matrix interface and testing only the first 3 outcomes
strk.mat <- xtabs(Freq ~ Type + Year, data=stroke)
multiCA.test(strk.mat, outcomes=1:3)

Description

Given the probabilities of outcomes, compute the power of the overall multinomial Cochran-Armitage trend test or determine the sample size to obtain a target power.

Usage

power.multiCA.test(N = NULL, power = NULL, pmatrix = NULL, p.ave = NULL,
  p.start = NULL, p.end = NULL, slopes = NULL, scores = 1:G,
  n.prop = rep(1, G), G = length(p.ave), sig.level = 0.05)
power.multiCA.test

Arguments

N integer, the total sample size of the study. If NULL then power needs to be specified.

power target power. If NULL then N needs to be specified.

pmatrix numeric matrix of hypothesized outcome probabilities in each group, with '#' the outcomes as rows and ordered groups as columns. The columns should add up to 1.

p.ave numeric vector of average probability of each outcome over the groups weighted by n.prop.

p.start, p.end numeric vectors of the probability of each outcome for the first / last ordered group

slopes numeric vector of the hypothesized slope of each outcome when regressed against the column scores with weights n.prop

scores non-decreasing numeric vector of the same length as the number of ordered groups giving the trend test scores. Defaults to linearly increasing values.

n.prop numeric vector describing relative sample sizes of the ordered groups. Will be normalized to sum to 1. Defaults to equal sample sizes.

G integer, number of ordered groups

sig.level significance level

Details

The distribution of the outcomes can be specified in two ways: either the full matrix of outcome probabilities pmatrix can be specified, or exactly two of the parameters p.ave, slopes, p.start, and p.end must be specified, while

Value

object of class "power.htest"

Examples

power.multiCA.test(power=0.8, p.start=c(0.1,0.2,0.3,0.4), p.end=c(0.4, 0.3, 0.2, 0.1), G=5, n.prop=c(3,2,1,2,3))

## Power of stroke study with 100 subjects per year and observed trends
data(stroke)
strk.mat <- xtabs(Freq ~ Type + Year, data=stroke)
power.multiCA.test(N=900, pmatrix=prop.table(strk.mat, margin=2))
**Description**

Nakajima et al. (2014) collected information on stroke patients over a 9-year period. For each patient, the type of stroke was classified into one of 5 categories by etiology.

**Usage**

```r
data("stroke")
```

**Format**

A data frame with 45 observations on the following 3 variables.

- **Type** a factor with levels `Small vessel occlusion`, `Large artery atherosclerosis`, `Cardioembolism`, `Other determined aetiology`, and `Undetermined aetiology` giving the etiology of the stroke.
- **Year** a numeric vector with the year of the observation.
- **Freq** a numeric vector with the number of patients with a stroke of the given etiology that year.

**Source**


**Examples**

```r
data(stroke)
xtabs(Freq ~ Type + Year, data=stroke)

strk.props <- prop.table(xtabs(Freq ~ Year+Type, data=stroke), margin=1)
matplot(strk.props, type="l")
```
Index

*Topic **datasets**
  stroke, 7
*Topic **htest**
  multiCA-package, 2
*Topic **nonparametric**
  multiCA-package, 2
  multiCA.test, 4
*Topic **package**
  multiCA-package, 2

cronct, 3
multiCA (multiCA-package), 2
multiCA-package, 2
multiCA.test, 4

power.multiCA.test, 5
stroke, 7