Package ‘GFD’

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Type Package
Title Tests for General Factorial Designs
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Depends R (>= 3.3.0)
Description Implemented are the Wald-type statistic,
a permuted version thereof as well as the ANOVA-type statistic
for general factorial designs, even with non-normal error terms
and/or heteroscedastic variances, for crossed designs with an
arbitrary number of factors and nested designs with up to three factors.
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(>= 1.5-6), plotrix (>= 3.5-12), methods
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R topics documented:

  calculateGUI .................................................. 2
  curdies .......................................................... 2
  GFD .............................................................. 3
  pizza ............................................................. 4
  startup .......................................................... 5

Index .......... 6
calculateGUI  

*A graphical user interface for the package GFD*

### Description

This function provides a graphical user interface for calculating statistical tests in general factorial designs.

### Usage

```r
calculateGUI()
```

### Details

The function produces a GUI for the calculation of the test statistics and for plotting. Data can be loaded via the "load data" button. The formula, number of permutations (default: 10,000) and the significance level alpha (default: 0.05) need to be specified. If the plot option is chosen, an additional window opens containing information on the plots.

curdies  

*Curdies river data set*

### Description

A dataset containing the number of flatworms (dugesia) sampled in two seasons at different sites in the Curdies River in Western Victoria.

### Usage

```r
data(curdies)
```

### Format

A data frame with 36 rows and 3 variables:

- **season** a factor with levels "SUMMER" and "WINTER"
- **site** a factor with levels 1 to 6, nested within "season"
- **dugesia** number of flatworms counted on a particular stone (in no./dm^2)

### Source

Description

The GFD function calculates the Wald-type statistic (WTS), the ANOVA-type statistic (ATS) as well as a permutation version of the WTS for general factorial designs.

Usage

GFD(formula, data = NULL, nperm = 10000, alpha = 0.05,
     nested.levels.unique = FALSE, CI.method = "t-quantile")

Arguments

- **formula**: A model formula object. The left hand side contains the response variable and the right hand side contains the factor variables of interest. An interaction term must be specified.
- **data**: A data.frame, list or environment containing the variables in formula. The default option is NULL.
- **nperm**: The number of permutations used for calculating the permuted Wald-type statistic. The default option is 10000.
- **alpha**: A number specifying the significance level; the default is 0.05.
- **nested.levels.unique**: A logical specifying whether the levels of the nested factor(s) are labeled uniquely or not. Default is FALSE, i.e., the levels of the nested factor are the same for each level of the main factor.
- **CI.method**: Method for calculating the confidence intervals. Default is ‘t-quantile’ for CIs based on the corresponding t-quantile. Additionally, the quantile of the permutation distribution can be used (‘perm’).

Details

The package provides the Wald-type statistic, a permuted version thereof as well as the ANOVA-type statistic for general factorial designs, even with non-normal error terms and/or heteroscedastic variances. It is implemented for both crossed and hierarchically nested designs and allows for an arbitrary number of factor combinations as well as different sample sizes in the crossed design. The GFD function returns three p-values: One for the ATS based on an F-quantile and two for the WTS, one based on the $\chi^2$ distribution and one based on the permutation procedure. Since the ATS is only an approximation and the WTS based on the $\chi^2$ distribution is known to be very liberal for small sample sizes, we recommend to use the WTPS in these situations.
**Value**

A GFD object containing the following components:

- **Descriptive**
  Some descriptive statistics of the data for all factor level combinations. Displayed are the number of individuals per factor level combination, the mean, variance and \(100\times(1-\alpha)\)\% confidence intervals.

- **WTS**
  The value of the WTS along with degrees of freedom of the central chi-square distribution and p-value, as well as the p-value of the permutation procedure.

- **ATS**
  The value of the ATS, degrees of freedom of the central F distribution and the corresponding p-value.

**References**


**Examples**

```r
GFD(weightgain ~ source * type, data = HSAUR::weightgain, nperm = 1000)

data(startup)
model <- GFD(Costs ~ company, data = startup, CI.method = "perm")
summary(model)
```

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**pizza**

*Pizza delivery times*

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**Description**

A dataset containing the delivery times for pizza (in minutes) under different conditions.

**Usage**

data(pizza)

**Format**

A data frame with 16 rows and 6 variables:

- **Crust** a factor with levels "thick" and "thin"
- **Coke** whether or not Coke was ordered with the pizza ("yes" or "no")
- **Bread** whether or not garlic bread was ordered with the pizza ("yes" or "no")
- **Driver** the sex of the driver, a factor with levels "M" and "F"
- **Hour** time of order in hours after midnight
- **Delivery** Delivery time in minutes
startup

Source

http://www.statsci.org/data/oz/pizza.html

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startup  

Startup Costs of five different companies

Description

A dataset containing the startup costs (in thousands of dollars) of five companies.

Usage

data(startup)

Format

A data frame with 60 rows and 2 variables:

- Costs  price, in thousands of dollars
- company  company, a factor with levels "pets", "pizza", "gifts", "shoes" and "bakery"

Source

http://college.cengage.com/mathematics/brase/understandable_statistics/7e/students/datasets/owan/frames/frame.html
Index

*Topic datasets
  curdies, 2
  pizza, 4
  startup, 5

  calculateGUI, 2
  curdies, 2

  formula, 3

  GFD, 3
  GUI (calculateGUI), 2

  pizza, 4
  startup, 5