# Package 'daewr'

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| <b>Description</b> This package contains data frames and functions used in the book "Design and Analysis of Experiments with R" |                  |  |  |  |  |  |
| License GPL-2   |                  |  |  |  |  |  |
| <b>Dependencies</b> FrF2,BsMD   |                  |  |  |  |  |  |
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| R topics documented:  |                  |  |  |  |  |  |
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# Description

This package contains the data sets and functions from the book Design and Analysis of Experiments with R published by CRC in 2013.

#### **Details**

Package: daewr Type: Package Version: 1.0 Date: 2012-05-10 License: GPL-2 LazyLoad: yes

# Author(s)

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## References

J. Lawson, Design and Analysis of Experiments with R, CRC 2013.

# **Examples**

```
Fcrit(.05,2,15)
Fpower(0.05,2,15,6.428)
BIBsize(6,3)
```

antifungal

Two-period crossover study of antifungal agent

# Description

Data from the Two-period crossover study of an antifungal agent in chapter 9 of Design and Analysis of Experiments with R

4 Apo

#### Usage

```
data(antifungal)
```

#### **Format**

A data frame with 34 observations on the following 5 variables.

```
Group a factor with levels 1 2
Subject a factor with levels 1 2 3 4 5 6 7 8 9 10 11 12 14 15 16 17 18
Period a factor with levels 1 2
Treat a factor with levels A B
pl a numeric vector
```

## **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

## **Examples**

```
data(antifungal)
```

Apo

apolipoprotein survey varaince component study

# Description

Data from the apolipoprotein survey variance component study of Chapter 5 in Design and Analysis of Experiments with R

# Usage

```
data(Apo)
```

#### **Format**

A data frame with 30 observations on the following 2 variables.

```
lab a factor with levels A B C D conc a numeric vector
```

# Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

```
data(Apo)
```

apple 5

apple

Confounded apple slice browning experiment

# Description

Data from the confounded apple slice browning experiment in chapter 7 of Design and Analysis of Experiments with R

# Usage

```
data(apple)
```

## **Format**

A data frame with 24 observations on the following 4 variables.

Block a factor with levels 1 2 3 4

A a factor with levels 0 1 2 3

B a factor with levels 0 1 2

rating a numeric vector containing the response

## Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

## **Examples**

```
data(apple)
```

arso

 $2^{(7-3)}$  arsenic removal experiment

# Description

Data from the  $2^{(7-3)}$  arsenic removal experiment in chapter 6 of Design and Analysis of Experiments with R

# Usage

```
data(arso)
```

6 augm

#### **Format**

```
A data frame with 8 observations on the following 8 variables.
```

```
A a factor with levels -1 1
```

B a factor with levels -1 1

C a factor with levels -1 1

D a factor with levels -1 1

E a factor with levels -1 1

F a factor with levels -1 1

G a factor with levels -1 1

y1 a numeric vector

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

## **Examples**

```
data(arso)
```

augm

 $2^{(7-3)}$  arsenic removal experiment augmented with mirror image

## **Description**

Data from the  $2^{(7-3)}$  arsenic removal experiment augmented with mirror image in chapter 6 of Design and Analysis of Experiments with R

## Usage

```
data(augm)
```

#### **Format**

A data frame with 8 observations on the following 8 variables.

```
A a factor with levels -1 1
```

B a factor with levels -1 1

C a factor with levels -1 1

fold a factor with levels original mirror

D a factor with levels -1 1

E a factor with levels -1 1

F a factor with levels -1 1

G a factor with levels -1 1

y a numeric vector

Bdish 7

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

## **Examples**

```
data(augm)
```

Bdish

Confounded Block Dishwashing Experiment

# Description

Data from the Confounded Block Dishwashing Experiment in chapter 7 of Design and Analysis of Experiments with R

# Usage

```
data(Bdish)
```

#### **Format**

A data frame with 16 observations on the following 5 variables.

Blocks a factor with levels 1 2 3 4

A a factor with levels -1 1

B a factor with levels -1 1

C a factor with levels -1 1

D a factor with levels -1 1

y a numeric vector containing the response

## Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

```
data(Bdish)
```

8 bha

Bff

Confounded block fractional mouse growth experiment

## **Description**

Data from the Confounded block fractional factorial mouse growth experiment in chapter 7 of Design and Analysis of Experiments with R

## Usage

```
data(Bff)
```

#### **Format**

A data frame with 16 observations on the following 5 variables.

```
Blocks a factor with levels 1 2 3 4 5 6 7 8
```

- A a factor with levels -1 1
- B a factor with levels -1 1
- C a factor with levels -1 1
- D a factor with levels -1 1
- E a factor with levels -1 1
- F a factor with levels -1 1
- G a factor with levels -1 1
- H a factor with levels -1 1

weight a numeric vector containing the response

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

## **Examples**

```
data(Bff)
```

bha

mouse liver enzyme experiment

# Description

Data from the mouse liver enzyme experiment in chapter 4 of Design and Analysis of Experiments with R

# Usage

```
data(bha)
```

BIBsize 9

#### **Format**

A data frame with 16 observations on the following 4 variables.

```
block a factor with levels 1 2
strain a factor with levels A/J 12901a NIH BALB/c
treat a factor with levels treated control
y a numeric vector
```

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

## **Examples**

data(bha)

BIBsize

Balanced incomplete blocksize

# Description

This function computes the number of blocks, treatment frequency and lambda for a potential BIB design

# Usage

```
BIBsize(t,k)
```

## **Arguments**

t input - number of levels of the treatment factor

k input - blocksize or number of experimental units per block

# Value

a list containing the b=number of blocks, r=number of treatment replicates and lambda for a potential BIB design with t levels of treatment factor and blocksize k.

#### Author(s)

John Lawson

10 bioequiv

## **Examples**

```
BIBsize(6,3)
## The function is currently defined as
BIBsize<-function(t,k)
{
 b<-t
  r<-0
  lambda<-0
  check<-0
  while (check==0) {
  while (r==0) {
     #cat("r=",r)
     testr<-(b*k)/t
     #cat("testr=",testr,"b=",b)
     if (testr==floor(testr)) {
       r<-testr
       } else {
       b<-b+1
       }
      #cat("b=",b, "r=",r)
      testl < -(r*(k-1))/(t-1)
      #cat("testl=",testl,"b=",b)
      if (testl==floor(testl)) {
       lambda<-testl
       check=1
       } else {
       r<-0
       b<-b+1
      #cat("b=",b, "r=",r)
        }
}
```

bioequiv

Extra-period crossover bioequivalence study

# Description

Data from the extra-period crossover bioequivalence study in chapter 9 of Design and Analysis of Experiments with R

## Usage

```
data(bioequiv)
```

## **Format**

A data frame with 108 observations on the following 5 variables.

```
Group a factor with levels 1 2
```

bioeqv 11

```
Subject a factor with levels 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 21 23 24 25 26 27 28 30 31 32 33 code34 35 36 120 122 129

Period a factor with levels 1 2 3

Treat a factor with levels A B
```

Carry a factor with levels none A B

y a numeric vector

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

## **Examples**

```
data(bioequiv)
```

bioeqv

Latin Square bioequivalence experiment

# Description

Data from the Latin Square bioequivalence experiment in chapter 4 of Design and Analysis of Experiments with R

## Usage

```
data(bioeqv)
```

#### **Format**

A data frame with 9 observations on the following 4 variables.

```
Period a factor with levels 1 2 code3

Subject a factor with levels 1 2 code3

Treat a factor with levels A B C

AUC a numeric vector
```

# Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

```
data(bioeqv)
```

BoxM

blood

Variance component study of calcium in blood serum

## **Description**

Data from the Variance component study of calcium in blood serum in chapter 5 of Design and Analysis of Experiments with R

## Usage

```
data(blood)
```

#### **Format**

A data frame with 27 observations on the following 3 variables.

```
sol a factor with levels 1 2 3 4 lab a factor with levels A B C calcium a numeric vector
```

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

## **Examples**

data(blood)

BoxM

Box and Meyer's unreplicated 2<sup>4</sup> from Chapter 3

## **Description**

Data from Box and Meyer's unreplicated  $2^4$  in chapter 3 of Design and Analysis of Experiments with  ${\bf R}$ 

# Usage

```
data(BoxM)
```

#### **Format**

A data frame with 16 observations on the following 4 variables.

- A a numeric vector containing the coded (-1,1) levels of factor A
- B a numeric vector containing the coded (-1,1) levels of factor B
- C a numeric vector containing the coded (-1,1) levels of factor C
- D a numeric vector containing the coded (-1,1) levels of factor D
- y a numeric vector containing the response

BPmonitor 13

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

#### References

```
Box, G. E. P. "George's Column", Quality Engineering, Vol. 3, pp. 405-410.
```

# **Examples**

```
data(BoxM)
```

**BPmonitor** 

blood pressure monitor experiment

## **Description**

Data from the blood pressure monitor experiment experiment in Chapter 7 of Design and Analysis of Experiments with R

# Usage

```
data(BPmonitor)
```

## **Format**

A data frame with 12 observations on the following 3 variables.

```
Block a factor with levels 1 2 3 4 5 6

Treatment a factor with levels "P" "A" "B" "C" pressure a numeric vector
```

## Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

```
data(BPmonitor)
```

14 cake

bread

Bread rise experiment data from Chapter 2

## **Description**

Data from the bread rise experiment in chapter 2 of Design and Analysis of Experiments with R

## Usage

```
data(bread)
```

#### **Format**

A data frame with 12 observations on the following 3 variables.

```
loaf a numeric vector
time a numeric vector
height a numeric vector
```

#### Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

## **Examples**

```
data(bread)
```

cake

Split-Plot response surface for cake baking experiment

# Description

Data from the Split-Plot response surface for cake baking experiment in chapter 10 of Design and Analysis of Experiments with R

# Usage

```
data(cake)
```

#### **Format**

A data frame with 11 observations on the following 4 variables.

```
Ovenrun a factor with levels 1 2 3 4
```

x1 a numeric vector

x2 a numeric vector

y a numeric vector

x1sq a numeric vector

x2sq a numeric vector

castf 15

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

```
data(cake)
```

castf

cast fatigue experiment

# Description

Data from the cast fatigue experiment in chapter 6 of Design and Analysis of Experiments with R

# Usage

```
data(castf)
```

#### **Format**

A data frame with 12 observations on the following 12 variables.

```
c8 a factor with levels -1 1
```

c9 a factor with levels -1 1

c10 a factor with levels -1 1

c11 a factor with levels -1 1

G a factor with levels -1 1

F a factor with levels -1 1

E a factor with levels -1 1

D a factor with levels -1 1

C a factor with levels -1 1

B a factor with levels -1 1

A a factor with levels -1 1

y a numeric vector

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

```
data(castf)
```

16 chem

cement

CCD design for cement workability experiment

## **Description**

Data from the CCD design for cement workability experiment in chapter 10 of Design and Analysis of Experiments with R

## Usage

```
data(cement)
```

## **Format**

A data frame with 20 observations on the following 4 variables.

Block a factor with levels 1 2

x1 a numeric vector

x2 a numeric vector

x3 a numeric vector

y a numeric vector

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

## **Examples**

```
data(cement)
```

chem

Chemical process experiment data from Chapter 3

## **Description**

Data from the Chemical process experiment in chapter 3 of Design and Analysis of Experiments with R

# Usage

```
data(chem)
```

#### **Format**

A data frame with 16 observations on the following 4 variables.

- A a numeric vector containing the coded (-1,1) levels of factor A
- B a numeric vector containing the coded (-1,1) levels of factor B
- C a numeric vector containing the coded (-1,1) levels of factor C
- D a numeric vector containing the coded (-1,1) levels of factor D
- y a numeric vector containing the response

chipman 17

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

```
data(chem)
```

chipman

Williams' crossover design for sprinting experiment

# Description

Data from the Williams' crossover design for sprinting experiment in chapter 9 of Design and Analysis of Experiments with R

# Usage

```
data(chipman)
```

## **Format**

A data frame with 36 observations on the following 5 variables.

Square a factor with levels 1 2

Group a factor with levels 1 2 3

Subject a factor with levels 1 2 code3 4 5 code6 7 8 code9 10 11 code12

Period a factor with levels 1 2 3

Treat a factor with levels 1 2 3

Carry a factor with levels 0 1 2 3

Time a numeric vector

## Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

data(chipman)

18 cont

COdata

CO emmisions experiment data from Chapter 3

#### **Description**

Data from the CO emissions experiment in chapter 3 of Design and Analysis of Experiments with R

# Usage

```
data(COdata)
```

#### **Format**

A data frame with 18 observations on the following 3 variables.

```
Eth a factor with levels 0.1 0.2 0.3
Ratio a factor with levels 14 15 16
CO a numeric vector
```

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

## **Examples**

```
data(COdata)
```

cont

Control factor array and summary statistics for controller circuit design experiment

#### **Description**

Data from the control factor array and summary statistics for controller circuit design experiment in chapter 12 of Design and Analysis of Experiments with R

## Usage

```
data(cont)
```

#### **Format**

A data frame with 18 observations on the following 6 variables.

A a numeric vector

B a numeric vector

C a numeric vector

D a numeric vector

F a numeric vector

1ns2 a numeric vector

cpipe 19

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

## **Examples**

```
data(cont)
```

cpipe

Split-plot response surface for ceramic pipe experiment

# Description

Data from the Split-plot response surface for ceramic pipe experiment in chapter 10 of Design and Analysis of Experiments with R

# Usage

```
data(cpipe)
```

#### **Format**

A data frame with 48 observations on the following 6 variables.

```
WP a factor with levels 1 2 3 4 5 6 7 8 9 10 11 12
```

A a numeric vector

B a numeric vector

P a numeric vector

Q a numeric vector

y a numeric vector

## Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

```
data(cpipe)
```

20 dairy

culture

paecilomyces variotii culture experiment

#### **Description**

Data from the paecilomyces variotii culture experiment experiment in chapter 6 of Design and Analysis of Experiments with R

## Usage

```
data(culture)
```

#### **Format**

A data frame with 16 observations on the following 9 variables.

```
A a factor with levels -1 1
```

B a factor with levels -1 1

C a factor with levels -1 1

D a factor with levels -1 1

E a factor with levels -1 1

F a factor with levels -1 1

G a factor with levels -1 1

H a factor with levels -1 1

y1 a numeric vector

# Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

## **Examples**

```
data(culture)
```

dairy

Repeated measures study with dairy cow diets

#### **Description**

Data from the Repeated measures study with dairy cow diets in chapter 9 of Design and Analysis of Experiments with R (compact format)

# Usage

```
data(dairy)
```

drug 21

#### **Format**

A data frame with 120 observations on the following 5 variables.

```
Diet a factor with levels "Barley" "Mixed" "Lupins" pr1 a numeric vector pr2 a numeric vector pr3 a numeric vector pr4 a numeric vector
```

#### Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

```
data(dairy)
```

drug

Data from rat behavior experiment in Chapter 4

# **Description**

Data from rat behavior experiment in Chapter 4 of Design and Analysis of Experiments with R

## Usage

```
data(drug)
```

#### **Format**

A data frame with 50 observations on the following 3 variables.

```
rat a factor with levels 1 2 3 4 5 6 7 8 9 10 dose a factor with levels 0.0 0.5 1.0 1.5 2.0 rate a numeric vector
```

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

```
data(drug)
```

22 eptaxs2

eptaxr

Single array and raw response for silicon layer growth experiment

## **Description**

Data from the single array and raw response for silicon layer growth experiment in chapter 12 of Design and Analysis of Experiments with R

# Usage

```
data(eptaxr)
```

#### **Format**

A data frame with 64 observations on the following 9 variables.

- A a numeric vector
- B a numeric vector
- C a numeric vector
- D a numeric vector
- E a numeric vector
- F a numeric vector
- G a numeric vector
- H a numeric vector
- y a numeric vector

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

data(eptaxr)

eptaxs2

Control array and variance of response for silicon layer growth experiment

# Description

Data from the control array and variance of response for silicon layer growth experiment in chapter 12 of Design and Analysis of Experiments with R

# Usage

```
data(eptaxs2)
```

eptaxyb 23

#### **Format**

A data frame with 16 observations on the following 9 variables.

A a numeric vector

B a numeric vector

C a numeric vector

D a numeric vector

E a numeric vector

F a numeric vector

G a numeric vector

H a numeric vector

s2 a numeric vector

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

## **Examples**

data(eptaxs2)

eptaxyb

Control array and mean response for silicon layer growth experiment

# Description

Data from the control array and mean response for silicon layer growth experiment in chapter 12 of Design and Analysis of Experiments with R

# Usage

data(eptaxyb)

#### **Format**

A data frame with 16 observations on the following 9 variables.

A a numeric vector

B a numeric vector

C a numeric vector

D a numeric vector

E a numeric vector

F a numeric vector

G a numeric vector

H a numeric vector

ybar a numeric vector

24 Fcrit

#### Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

```
data(eptaxyb)
```

Fcrit

F-Distribution critical values

# Description

Gets F-distribution critical values

# Usage

```
Fcrit(alpha, nu1, nu2)
```

# **Arguments**

alpha input- right tail area beyond critical value

nu1 input - numerator degrees of freedom for F-distribution

nu2 input - denominator degrees of freedom for F-distribution

# Value

returned critical value

# Author(s)

John Lawson

```
Fcrit(.05,2,15)
## The function is currently defined as
function(alpha,nu1,nu2) qf(1-alpha,nu1,nu2)
```

Fpower 25

| Fpower | F-Distribution Power Calculation |  |
|--------|----------------------------------|--|
|--------|----------------------------------|--|

#### **Description**

Calculates the power for the non-central F-distribution

## Usage

```
Fpower(alpha, nu1, nu2, nc)
```

## **Arguments**

alpha input - critical value alpha

nu1 input - degrees of freedom for numeratornu2 input - degrees of freedom for denominator

nc input - noncentrality parameter

# Value

probability of exceeding fcrit(alpha, nu1,nu2) with the non-central F-distribution with nu1 and nu2 degrees of freedom and noncentrality parameter nc

## Author(s)

John Lawson

# **Examples**

```
Fpower(0.05,2,15,6.428)
## The function is currently defined as
function(alpha,nu1,nu2,nc) 1-pf(Fcrit(alpha,nu1,nu2),nu1,nu2,nc)
```

fullnormal

Nornal Plot of Effects

## Description

This function makes a full normal plot of effects labeling those significant and by default adds the reference line (refline).

## Usage

```
fullnormal<-function(effects,labs,alpha=.05,refline="TRUE")</pre>
```

26 fullnormal

#### **Arguments**

effects input - This is the numeric vector of effects or regression coefficients to be plot-

ted; it should not include the intercept.

labs input - This is the character vector of effect labels usually obtained from the lm

function.

alpha input - The significance level of the test. The default is .05 if alpha is not speci-

fied.

refline input - logical, the default value is TRUE which indicates a reference line will

be added to the plot.

#### Author(s)

John Lawson

```
data(plasma)
sol<-lm(y~A*B*C*D*E,data=plasma)</pre>
# get whole plot effects and split plot effects
effects<-coef(sol)
effects<-effects[c(2:32)]
Wpeffects<-effects[c(1:4, 6:11, 16:19, 26)]</pre>
Speffects<-effects[c(5,12:15,20:25,27:31)]</pre>
#make separate normal plots
fullnormal(Wpeffects, names(Wpeffects), alpha=.10)
fullnormal(Speffects, names(Speffects), alpha=.05)
# make normal plot of all effects
fullnormal(effects, names(effects), alpha=.01, refline=FALSE)
## The function is currently defined as
fullnormal<-function(effects,labs,alpha=.05,refline="TRUE") {</pre>
crit<-LenthPlot(effects,alpha=alpha,plt=FALSE)["ME"]</pre>
names<-names(effects)</pre>
names<-gsub(':','',names)</pre>
names<-gsub('1','',names)</pre>
le<-length(effects)</pre>
 for (i in 1:le) {
     logc<-(abs(effects[i])<=crit)</pre>
     if (logc) {names[i]<-" "}</pre>
qqnorm(effects, ylab="Estimated Effects", xlab="Normal Scores")
x<-qqnorm(effects,plot=FALSE)</pre>
zscr<-(x$x)
# Splits effects into positive and negative for labeling
effp<-effects[zscr>0]
zp<-zscr[zscr>0]
namep<-names[zscr>0]
effn<-effects[zscr<0]
zn<-zscr[zscr<0]</pre>
namen<-names[zscr<0]</pre>
text(zp,effp,namep,pos=1)
text(zn,effn,namen,pos=3)
# calculate pse statistic
```

gagerr 27

```
ahe<-abs(effects)
s0<-1.5*median(ahe)
selhe<-ahe<(2.5*s0)
pse=1.5*median(ahe[selhe])
if (refline) {
# add reference line to plot
abline(0,pse)
}
}</pre>
```

gagerr

Gauge R&R Study

# Description

Data from the Gauge R&R Study in chapter 5 of Design and Analysis of Experiments with R

# Usage

```
data(gagerr)
```

## **Format**

A data frame with 60 observations on the following 3 variables.

```
part a factor with levels 1 2 3 4 5 6 7 8 9 10 oper a factor with levels 1 2 3 y a numeric vector
```

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

```
data(gagerr)
```

28 gapstat

gapstat

Gap Statistic

#### **Description**

This function calculates the gap statistic used in the automated Daniel method of detecting an outlier in an 8, 16 or 32 run  $2^{k-p}$  design.

## Usage

```
gapstat<-function(beta,pse)</pre>
```

## **Arguments**

beta input - This is the the vector of effects or regression coefficients calculated from

the design where an outlier is suspected. This vector of coefficients should not include the intercept. This function is called by the function Gaptest at the first

and second pass through the data.

pse input - This Lenth's pse statistic calculated on the vector beta befor the call to

this function.

#### Value

The result is the gap statistic  $gap = ((\beta_s - \beta_l)/pse)/(z_s - z_l)$ .

#### Author(s)

John Lawson

#### References

Lawson, J. and Gatlin, J. "Finding bad values in factorials - revisited" *Quality Engineering* Vol 18, pages 491-501., 2006.

```
Beta<-c(.06,.25,-.01,.5,0,-.02,0,.14,.03,-.01,.02,.04,.02,.01,.02)
names(Beta)<-c("A","B","C","D","AB","AC","AD","BC","BD","CD","ABC","ABD","ACD","BCD","ABCD")
gapstat(Beta,.06)
## The function is currently defined as
gapstat<-function(beta,pse) {</pre>
# computes the standardized gap score
p<-length(beta)</pre>
psehe<-pse
# gets positive coefficients
sel<-beta >= 0
betap<-beta[sel]
# sorts positive elements
betap<-sort(betap)</pre>
# gets Beta_s
betas<-betap[1]
#gets negative coefficients
```

```
sel<-beta < 0
betan<-beta[sel]
nn<-length(betan)
# sorts negative coefficients
betan<-sort(betan)
#gets Beta_L
betal<-betan[nn]
# gets Z_L and Z_S
zl<-qnorm((nn-.375)/(p+.25))
zs<-qnorm((nn+1-.375)/(p+.25))
# calculates gap statistic
gap<-((betas-betal)/psehe)/(zs-zl)
return(gap)
}</pre>
```

Gaptest

Automated Daniel Method

## **Description**

This function performs and automated version of Daniels method for detecting an outlier in a  $2^{(k-p)}$  design by recognizing a gap in the full normal plot of effects. This function calls the functions gapstat and LGB.

## Usage

```
Gaptest<-function(DesY)</pre>
```

# Arguments

DesY

input - This is a data frame containing a  $2^{(k-p)}$  design augmented with a single column y of responses. This function only works for 8, 16 or 32 run designs.

#### Author(s)

John Lawson

## References

Lawson, J. and Gatlin, J. "Finding bad values in factorials - revisited" *Quality Engineering* Vol 18, pages 491-501., 2006.

```
# Lawson Gatlin Example
library(FrF2)
lawg<-FrF2(16,11,generators=c("ABC","BCD","ACD","ABD","ABCD","AB","AC"),randomize=FALSE)
y<-c(31,48,43,32,54,52,60,34,47,43,30,26,67,49,80,41)
lawg<-cbind(lawg,y)
Gaptest(lawg)
# eight run example
```

```
library(FrF2)
des<-FrF2(8,6,generators=c("AB","AC","BC"),randomize=FALSE)</pre>
y < -c(1.299, 1.601, 1.359, 1.461, 1.338, 1.486, 1.330, 1.470)
Des8<-cbind(des,y)
Gaptest(Des8)
#tes<-lm(y~(.)^3,data=Des8)</pre>
#summary(tes)
#c<-tes$coef
#cn<-names(c)</pre>
#ccn<-gsub("[^A-Z]","",cn)</pre>
## The function is currently defined as
Gaptest<-function(DesY) {</pre>
# function to compute gap statistic
ncheck<-dim(DesY)</pre>
ncheck<-ncheck[1]
 tcnd=TRUE
 if (ncheck==8) {tcnd=FALSE}
 if (ncheck==16) {tcnd=FALSE}
 if (ncheck==32) {tcnd=FALSE}
 if (tcnd) {stop("This function only works for 8, 16, or 32 run designs","\n")
    } else {
if (ncheck==8) ncheck=16
# 50th and 99th percentiles of the gap statistic ##
critg16<-c(1.7884,5.1009)
critg32<-c(1.7297,5.8758)
### First Pass through the data ####
###### Step 1 ######
#fit model to saturated design
modf < -lm(y^{(.)}^4, x=TRUE, data=DesY)
#extract the regression coefficients
nbeta<-dim(DesY)</pre>
nbeta<-nbeta[1]
he<-modf$coef
# This extracts the coefficients that are not NA
selcol<-which(!is.na(he))</pre>
he<-he[selcol]
he<-he[-1]
#number of coefficients
p<-length(he)</pre>
#number of runs
n<-p+1
# This trims unnecessary characters from coefficient names
cn1<-names(he)
ccn1<-gsub("[^A-Z]","",cn1)</pre>
names(he)<-ccn1
##### End of Step 1 #######
```

```
##### Steps 2 and 3 ######
#calculate the pse statistic
ahe<-abs(he)
s0<-1.5*median(ahe)
selhe < -ahe < (2.5*s0)
pse=1.5*median(ahe[selhe])
#library(BsMD)
#pse<-LenthPlot(modf,plt=FALSE)</pre>
#pse<-pse[2]</pre>
#calculate the gap statistic
gap<-gapstat(he,pse)</pre>
# checks to see if gap statistic exceeds 50th percentile
if (ncheck==16) {test=(gap>critg16[1])
} else {test=(gap>critg32[1])}
##### End Step 2 and 3 #####
if (test) {
##### Step 4 #####
#extract the model X matrix
X < -modf x
# This selects columns of the X matrix that correspond to non-missing
# coefficients
X<-X[,selcol]
X < -X[,-1]
#gets signs of regression coefficients
se<-as.matrix(sign(he),nrow=1)</pre>
# find signigicant effects using LGB
sigef<-LGB(he,rpt=FALSE,plt=FALSE)</pre>
# make signs of significant effects zero
 for (i in 1:length(he)) {
     if (sigef[i]=="yes")
                             {se[i]=0 }
                          }
#gets sum of products of signed effects and rows of X matrix
sp<-X
#finds index of largest sum of products as index of potential outlier
asp<-abs(sp)
oo<-max.col(t(asp))</pre>
### End Step 4 ####
##### Step 5 #####
# calculates the bias
 # first get absolute regression coefficients
ae<-abs(he)
 # next sort absolute effects
sae<-sort(ae)</pre>
 #get the number of effects in smallest half
nsmall<-round(length(he)/2)</pre>
# sum the smallest half absolute effects to get bias
bias<-2*sum(sae[1:nsmall])</pre>
##### Step 6 #####
# gets corrected response vector
y<-DesY$y
ycorr<-DesY$y
```

```
ycorr[oo]<-ycorr[oo]+(-1*sign(sp[oo]))*bias</pre>
 # makes vector of indicators for outlier
detect<-c(rep("no",n))</pre>
detect[oo]<-"yes"
cat("Initial Outlier Report","\n")
cat("Standardized-Gap = ",gap, "Significant at 50th percentile","\n")
### End of first pass throught the data ######
### Second Pass throught the data #########
### Step 1 ####
# augment DesY with corrected data
DesYc<-cbind(DesY[,1:(dim(DesY)[2]-1)],ycorr)</pre>
# fit saturated model to corrected data
modf<-lm(ycorr~(.)^4,x=TRUE,data=DesYc)</pre>
#extract the regression coefficients
che<-modf$coef
# This extracts the coefficients that are not NA
che<-che[!is.na(che)]</pre>
che<-che[-1]
#number of coefficients
p<-length(che)
#number of runs
n<-p+1
# This trims unnecessary characters from coefficient names
cn<-names(che)</pre>
ccn<-gsub("[^A-Z]","",cn)</pre>
names(che)<-ccn
### End of Step 1 ####
##### Steps 2 and 3 ######
#calculate the pse statistic
ache<-abs(che)
s0<-1.5*median(ache)
selche < -ache < (2.5*s0)
psec=1.5*median(ache[selche])
#psec<-LenthPlot(modf,plt=FALSE)</pre>
#psec<-psec[2]</pre>
#calculate the gap statistic
gap<-gapstat(he,psec)</pre>
# checks to see if gap statistic exceeds 99th percentile
if (ncheck==16) test2=(gap>critg16[2]) else test2=(gap>critg32[2])
##### End Step 2 and 3 #####
if (test2) {
cat("Final Outlier Report","\n")
cat("Standardized-Gap = ",gap, "Significant at 99th percentile","\n")
cat("
        ","\n")
cat("
        Corrrected Data Report ","\n")
cat("Response Corrected Response Detect Outlier","\n")
cat(paste(format(DesY$y, width=8), format(DesYc$ycorr, width=13),
            ", format(detect, width=10),"\n"),sep="")
# use LGB to test significance of effects calculated from corrected data
tce<-LGB(che)
```

gear 33

```
} else {
cat("Final Outlier Report","\n")
cat("No significant outlier detected in second pass","\n")
# use LGB to test significance of effects calculated from corrected data
LGB(he)
cat(" ","\n")
}
### End of second pass through the data #####
}
}
# end of function Gaptest
}
```

gear

Unreplicated split-plot fractional-factorial experiment on geometric distortion of drive gears

## **Description**

Data from the unreplicated split-plot fractional-factorial experiment on geometric distortion of drive gears in chapter 8 of Design and Analysis of Experiments with R

## Usage

```
data(gear)
```

## **Format**

A data frame with 16 observations on the following 6 variables.

```
A a factor with levels -1 1
```

B a factor with levels -1 1

C a factor with levels -1 1

P a factor with levels -1 1

Q a factor with levels -1 1

y a numeric vector containing the response

## Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

```
data(gear)
```

inject inject

hardwood

low grade hardwood conjoint study

## **Description**

Data from the low grade hardwood conjoint study in chapter 6 of Design and Analysis of Experiments with R

## Usage

```
data(hardwood)
```

#### **Format**

A data frame with 12 observations on the following 5 variables.

```
Design a factor with levels "RC" "AC" "OCI" "OCII"

Price a numeric variable

Density a factor with levels "Clear" "Heavy" "Medium"

Guarantee a factor with levels "1y" "Un"

Rating a numeric vector
```

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

```
data(hardwood)
```

inject

Single array for injection molding experiment

# Description

Data from the single array for injection molding experiment in chapter 12 of Design and Analysis of Experiments with R

## Usage

```
data(inject)
```

LGB 35

#### **Format**

A data frame with 20 observations on the following 8 variables.

A a numeric vector

B a numeric vector

C a numeric vector

D a numeric vector

E a numeric vector

F a numeric vector

G a numeric vector

shrinkage a numeric vector

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

## **Examples**

data(inject)

Lawson, Grimshaw, Burt Test

# Description

LGB

This function uses the LGB method (based on the half normal plot) to test significance of effects in unreplicated orthogonal 2-level designs.

# Usage

LGB<-function(Beta,alpha=.05,rpt=TRUE,plt=TRUE,pltl=TRUE)

## **Arguments**

| Beta  | input - This is the the vector of effects or regression coefficients to be tested; it should not include the intercept. This function only works for vetors of length 7,8,11,15,16,26,31,32,35,63,or 127.  |
|-------|--|
| alpha | input - The significance level of the test. The only options available are .1, .05, .025, or .01, and the default is .05 if alpha is not specified.  |
| rpt   | input - logical, the default value is TRUE which indicates the report printout is desired.   |
| plt   | input - logical, the default value is TRUE which indicates the half-normal plot with reference line and significance limit is desired.   |
| pltl  | input - logical, the default value is TRUE which indicates the significance limit will be shown on the half-normal plot. Make pltl=FALSE, if you want to remove the significance limit line from the plot. |

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#### Value

The result is a vector of indicators "yes" or "no" for the significance of the coefficients in Beta at the alpha significance level.

#### Author(s)

John Lawson

#### References

Lawson, J., Grimshaw, S. and Burt, J. "A quantitative method for identifying active contrasts in unreplicated factorial experiments based on the half-normal plot" *Computational Statistics and Data Analysis* Vol 26, pages 425-436., 1998.

```
Beta<-c(.06,.25,-.01,.5,0,-.02,0,.14,.03,-.01,.02,.04,.02,.01,.02)
names(Beta)<-c("A","B","C","D","AB","AC","AD","BD","CD","ABC","ABD","ACD","BCD","BCD","ABCD")
LGB(Beta,alpha=.05)
## The function is currently defined as
LGB<-function(Beta,alpha=.05,rpt=TRUE,plt=TRUE,pltl=TRUE) {
# function to compute the LGB statistic
siglev<-c(.1,.05,.025,.01)
df<-c(7,8,11,15,16, 17, 26, 31, 32, 35, 63, 127)
crittab<-matrix(c(1.265,1.196,1.161,1.122,1.110, 1.106, 1.072, 1.063, 1.060, 1.059, 1.037, 1.023,
1.534, 1.385, 1.291, 1.201, 1.186, 1.178, 1.115, 1.099, 1.093, 1.091, 1.056, 1.034,
1.889, 1.606, 1.449, 1.297, 1.274, 1.260, 1.165, 1.140, 1.130, 1.127, 1.074, 1.043,
2.506, 2.026, 1.74, 1.447, 1.421, 1.377, 1.232, 1.197, 1.185, 1.178, 1.096, 1.058),ncol=4,byrow=FALSE)
# get the critical value
colind<-which(siglev==alpha,arr.ind=TRUE)</pre>
if (length(colind)==0) {stop("this function works only when alpha= .1, .05, .025 or .01") }
rowind<-which(df==length(Beta),arr.ind=TRUE)</pre>
if (length(rowind)==0) {stop("this function works only for coefficent vectors of length
7,8,11,15,16,26,31,32,35,63,or 127")}
critL<-crittab[rowind,colind]</pre>
# calculate Beta1, Beta2 and the Rn statistic
acj<-abs(Beta)</pre>
ranks<-rank(acj,ties.method="first")</pre>
s0<-1.5*median(acj)
p<-(ranks-.5)/length(Beta)</pre>
z<-qnorm((p+1)/2)
moda < -lm(acj \sim -1 + z)
beta1<-moda$coef
sel < -acj < 2.5 *s0
modi<-lm(acj[sel]~-1+z[sel])</pre>
beta2<-modi$coef
Rn<-beta1/beta2
# finds prediction limits for values in sorted absolute Beta
pred<-beta2*z
n<-length(acj[sel])</pre>
df < -n-1
```

mod 37

```
sig<-sqrt(sum(modi$residuals^2)/df)</pre>
se.pred<-sig*(1+1/n+(z^2)/sum(z[sel]^2))^.5
pred.lim<-pred+qt(.975,df)*se.pred</pre>
# gets significance indicators
sigi<-c(rep("no",length(Beta)))</pre>
sel2<-acj>pred.lim
sigi[sel2]<-"yes"
if (plt) {
plot(z,acj,xlab="Half Normal Scores", ylab="Absoulute Effects")
lines(sort(z),sort(pred),lty=1)
 for (i in 1:length(Beta)) {
   if (sigi[i]=="yes") text(z[i],acj[i],names(Beta)[i],pos=1)
 if (pltl) {
lines(sort(z),sort(pred.lim),lty=3)\\
          }
          }
if (rpt) {
cat("Effect Report","\n")
cat(" ","\n")
cat("Label Half Effect Sig(.05)","\n")
cat(paste(format(names(Beta), width=8), format(Beta, width=8),
       ",format(sigi, width=10),"\n"),sep="")
cat(" ","\n")
cat("Lawson, Grimshaw & Burt Rn Statistic = ",Rn,"\n")
cat("95th percentile of Rn = ",critL,"\n")
return(sigi)
                     }
```

mod

Mod function

### **Description**

Gets mod of a to base b

### Usage

mod(a,b)

### **Arguments**

a input- an integerb input - an integer

### Value

remainder of a/b or mod(a,b)

# Author(s)

John Lawson

38 MPV

### **Examples**

```
mod(5,3)
## The function is currently defined as
mod<-function(a,b)
{a-b*floor(a/b)}</pre>
```

MPV

mixture process variable experiment with mayonnaise

# Description

Data from the mixture process variable experiment with mayonnaise in chapter 11 of Design and Analysis of Experiments with R

### Usage

data(MPV)

### **Format**

A data frame with 35 observations on the following 4 variables.

x1 a numeric vector

x2 a numeric vector

x3 a numeric vector

z1 a numeric vector

z2 a numeric vector

y a numeric vector

# Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

```
data(MPV)
```

Naph 39

Naph

Yields of naphthalene black

### **Description**

Data from the Yields of naphthalene black of Chapter 5 in Design and Analysis of Experiments with R

### Usage

```
data(Naph)
```

### **Format**

A data frame with 30 observations on the following 2 variables.

```
sample a factor with levels 1 2 3 4 5 6 yield a numeric vector
```

### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

### **Examples**

```
data(Naph)
```

pastry

Blocked response surface design for pastry dough experiment

### **Description**

Data from the Blocked response surface design for pastry dough experiment in chapter 10 of Design and Analysis of Experiments with R

# Usage

```
data(pastry)
```

### **Format**

A data frame with 28 observations on the following 5 variables.

```
Block a factor with levels 1 2 3 4 5 6 7
```

x1 a numeric vector

x2 a numeric vector

x3 a numeric vector

y a numeric vector

40 pesticide

### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

```
data(pastry)
```

pest

Pesticide formulation experiment

# Description

Data from the Pesticide formulation experiment in chapter 11 of Design and Analysis of Experiments with R

# Usage

```
data(pest)
```

### **Format**

A data frame with 13 observations on the following 4 variables.

x1 a numeric vector

x2 a numeric vector

x3 a numeric vector

y a numeric vector

# Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

```
data(pest)
```

pesticide

pesticide application experiment

# Description

Data from the pesticide application experiment in chapter 5 of Design and Analysis of Experiments with R

# Usage

```
data(pesticide)
```

plasma 41

### **Format**

A data frame with 16 observations on the following 4 variables.

```
form a factor with levels A B tech a factor with levels 1 2 plot a factor with levels 1 2 residue a numeric vector
```

### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

```
data(pesticide)
```

plasma

Unreplicated split-plot 2^5 experiment on plasma treatment of paper

# **Description**

Data from the unreplicated split-plot  $2^5$  experiment on plasma treatment of paper in chapter 8 of Design and Analysis of Experiments with R

### Usage

```
data(plasma)
```

#### **Format**

A data frame with 32 observations on the following 6 variables.

```
A a factor with levels -1 1
```

B a factor with levels -1 1

C a factor with levels -1 1

D a factor with levels -1 1

E a factor with levels -1 1

y a numeric vector containing the response

# Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

```
data(plasma)
```

42 polymer

polvdat

Polvoron mixture experiment

### **Description**

Data from the Polvoron mixture experiment in chapter 11 of Design and Analysis of Experiments with R

### Usage

```
data(polvdat)
```

### **Format**

A data frame with 12 observations on the following 4 variables.

x1 a numeric vector

x2 a numeric vector

x3 a numeric vector

y a numeric vector

### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

### **Examples**

```
data(polvdat)
```

polymer

polymerization strength variability study

## **Description**

Data from the polymerization strength variability study in chapter 5 of Design and Analysis of Experiments with R

### Usage

```
data(polymer)
```

### **Format**

A data frame with 120 observations on the following 5 variables.

```
lot a factor with levels 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30
```

box a factor with levels 1 2

prep a factor with levels 1 2

test a factor with levels 1 2

strength a numeric vector

prodstd 43

### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

data(polymer)

prodstd

Complete control factor array and noise factor array for connector experiment

# Description

Data from the complete control factor array and noise factor array for connector experiment in chapter 12 of Design and Analysis of Experiments with R

# Usage

data(prodstd)

### **Format**

A data frame with 16 observations on the following 16 variables.

- A a numeric vector
- B a numeric vector
- C a numeric vector
- D a numeric vector
- E a numeric vector
- F a numeric vector
- G a numeric vector
- Pof a numeric vector

# Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

data(prodstd)

rcb

qsar

Library of substituted hydroxyphenylurea compounds

### **Description**

Data from the Library of substituted hydroxyphenylurea compounds in chapter 10 of Design and Analysis of Experiments with R (compact format)

### Usage

```
data(qsar)
```

### **Format**

A data frame with 36 observations on the following 4 variables.

Compound a numeric vector

HE a numeric vector

DMz a numeric vector

S0K a numeric vector

### Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

### **Examples**

```
data(qsar)
```

rcb

generalized RCB golf driving experiment

### **Description**

Data from the generalized RCB golf driving experiment in chapter 4 of Design and Analysis of Experiments with R

### Usage

```
data(rcb)
```

### **Format**

A data frame with 135 observations on the following 3 variables.

```
id a factor with levels 1 2 3 4 5 6 7 8 9
teehgt a factor with levels 1 2 3
cdistance a numeric vector
```

residue 45

### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

```
data(rcb)
```

residue

Herbicide degradation experiment

# Description

Data from the Herbicide degradation experiment in chapter 9 of Design and Analysis of Experiments with R

# Usage

```
data(residue)
```

### **Format**

A data frame with 16 observations on the following 3 variables.

```
soil a factor with levels "C" "P"
moisture a factor with levels "L" "H"
temp a factor with levels 10 30
X1 a numeric vector
X2 a numeric vector
X3 a numeric vector
X4 a numeric vector
X5 a numeric vector
```

### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

```
data(residue)
```

46 sausage

rubber

Rubber Elasticity data

# Description

Data from the Rubber Elasticity Study in chapter 5 of Design and Analysis of Experiments with R

### Usage

```
data(rubber)
```

#### **Format**

A data frame with 96 observations on the following 4 variables.

```
supplier a factor with levels A B C D
batch a factor with levels I II III IV
sample a factor with levels 1 2
elasticity a numeric vector
```

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

### **Examples**

```
data(rubber)
```

sausage

Split-plot experiment on sausage casing with RCB in whole plot

# **Description**

Data from the Split-plot experiment on sausage casing with RCB in whole plot in chapter 7 of Design and Analysis of Experiments with R

### Usage

```
data(sausage)
```

#### **Format**

A data frame with 32 observations on the following 5 variables.

```
Block a factor with levels 1 2
Gbatch a factor with levels 1 2 3 4
A a factor with levels -1 1
B a factor with levels -1 1
C a factor with levels -1 1
```

D a factor with levels -1 1

ys a numeric vector containing the response

Smotor 47

### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

### **Examples**

```
data(sausage)
```

Smotor

Single array for starting motor experiment

# Description

Data from the single array for starting motor experiment in chapter 12 of Design and Analysis of Experiments with R

# Usage

```
data(Smotor)
```

### **Format**

A data frame with 18 observations on the following 6 variables.

A a factor with levels 1 2

B a factor with levels 1 2 3

C a factor with levels 1 2 3

D a factor with levels 1 2 3

E a factor with levels 1 2

torque a numeric vector

### Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

```
data(Smotor)
```

48 soupmx

soup

dry mix soup experiment

### **Description**

Data from the dry mix soup experiment in chapter 6 of Design and Analysis of Experiments with R

### Usage

```
data(soup)
```

### **Format**

A data frame with 16 observations on the following 6 variables.

```
A a factor with levels -1 1
```

- B a factor with levels -1 1
- C a factor with levels -1 1
- D a factor with levels -1 1
- E a factor with levels -1 1
- y a numeric vector

### Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

### **Examples**

data(soup)

soupmx

dry soup mix variance component study

### **Description**

Data from the dry soup mix variance component study of Chapter 5 in Design and Analysis of Experiments with R

# Usage

```
data(soupmx)
```

### **Format**

A data frame with 12 observations on the following 2 variables.

```
batch a factor with levels 1 2 3 4 weight a numeric vector
```

splitPdes 49

### Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

```
data(soupmx)
```

splitPdes

Split-plot cookie baking experiment

# Description

Data from the Split-plot cookie baking experiment in chapter 8 of Design and Analysis of Experiments with R

# Usage

```
data(splitPdes)
```

### **Format**

A data frame with 24 observations on the following 5 variables.

```
short a factor with levels 100 80
trayT a factor with levels RoomT Hot
bakeT a factor with levels low mid high
batch a factor with levels 1 2
y a numeric vector
```

### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

```
data(splitPdes)
```

50 strung

SPMPV

Split-plot mixture process variable experiment with vinyl

# Description

Data from the Split-plot mixture process variable experiment with vinyl in chapter 10 of Design and Analysis of Experiments with R

### Usage

```
data(SPMPV)
```

### **Format**

A data frame with 28 observations on the following 7 variables.

```
wp a factor with levels 1 2 3 4 5 6 7
```

- z1 a numeric vector
- z2 a numeric vector
- x1 a numeric vector
- x2 a numeric vector
- x3 a numeric vector
- y a numeric vector

### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

data(SPMPV)

strung

Repeated measures study with dairy cow diets

### **Description**

Data from the Repeated measures study with dairy cow diets in chapter 9 of Design and Analysis of Experiments with R (strung out format)

### Usage

```
data(strung)
```

strungtile 51

### **Format**

```
A data frame with 120 observations on the following 5 variables.
```

```
Diet a factor with levels "Barley" "Mixed" "Lupins"
Cow a factor with levels 1 2 3 4 5 6 7 8 9 10
week a factor with levels 1 2 3 4
protein a numeric vector
```

### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

### **Examples**

```
data(strung)
```

| strungtile | Strung out control factor array and raw response data for Ina tile ex- |
|------------|--|
|            | periment   |

### **Description**

Data from the strung out control factor array and raw response data for Ina tile experiment in chapter 12 of Design and Analysis of Experiments with R

### Usage

```
data(strungtile)
```

# **Format**

A data frame with 16 observations on the following 16 variables.

- A a numeric vector
- B a numeric vector
- C a numeric vector
- D a numeric vector
- E a numeric vector
- F a numeric vector
- G a numeric vector
- H a numeric vector
- AH a numeric vector
- BH a numeric vector
- CH a numeric vector
- DH a numeric vector
- EH a numeric vector
- FH a numeric vector
- GH a numeric vector
- y a numeric vector

52 taste

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

### **Examples**

```
data(strungtile)
```

sugarbeet

Sugarbeet data from Chapter 2

### **Description**

Sugarbeet data from chapter 2 of Design and Analysis of Experiments with R

### Usage

```
data(sugarbeet)
```

#### **Format**

```
A data frame with 18 observations on the following 2 variables.
```

```
treat a factor with levels A B C D yield a numeric vector
```

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

### **Examples**

```
data(sugarbeet)
```

taste

taste test panel experiment

### **Description**

Data from the taste test panel experiment in Chapter 7 of Design and Analysis of Experiments with R

# Usage

```
data(taste)
```

### **Format**

```
A data frame with 24 observations on the following 3 variables.
```

```
panelist a factor with levels 1 2 3 4 5 6 7 8 9 10 11 12 recipe a factor with levels "A" "B" "C" "D" score a numeric vector
```

teach 53

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

### **Examples**

```
data(taste)
```

teach

Teaching experiment data from Chapter 2

### **Description**

Data from the teaching experiment in chapter 2 of Design and Analysis of Experiments with R

### Usage

```
data(teach)
```

#### **Format**

A data frame with 30 observations on the following 4 variables.

```
class a numeric vector
method a factor with levels 1 2 3
score a factor with levels 1 2 3 4 5
count a numeric vector
```

### Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

```
data(teach)
```

Tet

Tetracycline concentration in plasma

### **Description**

Data from the Tetracycline concentration in plasma study in chapter 10 of Design and Analysis of Experiments with R (compact format)

# Usage

```
data(Tet)
```

54 tile

#### **Format**

A data frame with 9 observations on the following 2 variables.

```
Time a numeric vector

Conc a numeric vector
```

### Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

```
data(Tet)
```

tile

Control factor array and summary statistics for Ina tile experiment

# Description

Data from the control factor array and summary statistics for Ina tile experiment in chapter 12 of Design and Analysis of Experiments with R

### Usage

```
data(tile)
```

### **Format**

A data frame with 8 observations on the following 11 variables.

A a numeric vector

B a numeric vector

C a numeric vector

D a numeric vector

E a numeric vector

F a numeric vector

G a numeric vector

y1 a numeric vector

y2 a numeric vector

ybar a numeric vector

1ns2 a numeric vector

### Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

```
data(tile)
```

Treb 55

Treb Box-Behnken design for trebuchet experiment

### **Description**

Data from the Box-Behnken design for trebuchet experiment in chapter 10 of Design and Analysis of Experiments with R

### Usage

```
data(Treb)
```

#### **Format**

A data frame with 15 observations on the following 4 variables.

x1 a numeric vector

x2 a numeric vector

x3 a numeric vector

y a numeric vector

### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

data(Treb)

Tukey1df

Tukey's single degree of freedom test for additivity

### **Description**

This function performs Tukey's single degree of freedom test for additivity in an unreplicated two-factor factorial.

### Usage

```
Tukey1df<-function(data)</pre>
```

### **Arguments**

data

input - This is a data frame containing an unreplicated two-factor factorial. The first column should be an numeric response, the second and third columns should be factors. At least one of the factors should have more than 2 levels.

### Author(s)

John Lawson

56 Tukey1df

#### References

Tukey, J. "One degree of freedom test for non-additivity" Biometrics Vol 5, pages 232-242., 1949.

```
Dilution<-rep(c(3,4,5),6)
 Sample < -rep(c(1,2,3,4,5,6), each=3)
       y < -c(1.87506, 1.38021, 0.60206, 1.74036, 1.36173, 0.90309, 1.79934, 1.25527, 0.95424, 2.02119, 1.39794, 1.000, 1.79936, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.36173, 1.361
       virus<-data.frame(y=y, Sample=factor(Sample),Dilution=factor(Dilution))</pre>
Tukey1df(virus)
 ## The function is currently defined as
 Tukey1df<-function(data) {</pre>
 y<-data[,1]
 Afactor<-data[,2]
Bfactor<-data[,3]
 tst1<-is.factor(Afactor)</pre>
 tst2<-is.factor(Bfactor)</pre>
 tst3<-is.numeric(y)</pre>
 if (tst1&tst2&tst3) {
a<-nlevels(Afactor)
b<-nlevels(Bfactor)
                } else {stop("The first column of the data frame is the numeric response, the 2nd and 3rd columns should
 tst4 < -max(a,b) > 2
 tst5 < -length(y) = = a * b
  if (tst4&tst5) {
 ybb<-with(data, tapply(y, Bfactor, mean))</pre>
 yba<-with(data, tapply(y, Afactor, mean))</pre>
 sbb<-with(data, tapply(y, Bfactor, sum))</pre>
 sba<-with(data, tapply(y, Afactor, sum))</pre>
 ybardd<-mean(y)</pre>
 CT < -(sum(y)^2)/(a*b)
 ssA<-sum(sba^2/b)-CT
 ssB<-sum(sbb^2/a)-CT
 ssE<-sum(y^2)-CT-ssA-ssB
 ybdj<-rep(ybb,6)</pre>
 prody<-y*ybdj
 sumprod<-tapply(prody,Afactor,sum)</pre>
 leftsum<-sum(sumprod*yba)</pre>
 ssAB < -(a*b*(leftsum-(ssA+ssB+a*b*ybardd^2)*ybardd)^2/(ssA*ssB))
 ssR<-ssE-ssAB
F < -ssAB/(ssR/((a-1)*(b-1)-1))
Pval < -1-pf(1,((a-1)*(b-1)-1),F)
                                                                                                                                                                                                                                                                                                                                                                                                                                      Pr>F","\n")
 cat("Source
                                                                                                                                                                   df
                                                                                                                                                                                                                                                                                                                                                                                        F
                                                                                                                                           ", paste(format(a-1, width=6),"", format(round(ssA,4), justify="right")," \\ ", format(suA,4), justify="right")," \\ ", format
 cat("A
                                                                                                                                           ",paste(format(b-1, width=6)," ", format(round(ssB,4),justify="right")," ",format(round
 cat("B
                                                                                                                                           ", paste(format((b-1)*(a-1), width=6),"", format(round(ssE,4), justify="right")," \\ ", format(round(ssE,4), justify="right")," \\ ", format(ssE,4), justify="right")," \\ ", f
cat("Error
cat("NonAdditivity",paste(format(1, width=6)," ", format(round(ssAB,4),justify="right")," ",format(round(ssAB,4),justify="right")," ",format(suAB,4),justify="right")," ",format(suAB,4),justify=
                                                                                                                                           ",paste(format((b-1)*(a-1)-1, width=6)," ", format(round(ssR,4),justify="right")," ",format(round(ssR,4),justify="right")," ",format(ssR,4),justify="right")," ",format(ssR,4),justify="right"
 cat("Residual
                        } else {stop("This function only works for unreplicated 2-factor factorials with >2 levels for one of the stop of the factorials with >2 levels for one of the f
```

vci 57

| vci | confidence limits for method of moments estimators of variance components |
|-----|---|
|     |   |

# Description

function for getting confidence intervals on variance components estimated by the method of moments

### Usage

```
vci(confl,c1,ms1,nu1,c2,ms2,nu2)
```

### **Arguments**

| confl | input- confidence level   |
|-------|---|
| c1    | input - linear combination coefficient of ms1 in the estimated variance component |
| ms1   | input - Anova mean square 1   |
| nu1   | input - Anova degrees of freedom for mean square 1                                |
| c2    | input - linear combination coefficient of ms2 in the estimated variance component |
| ms2   | input - Anova mean square 2   |
| nu2   | input - Anova degrees of freedom for mean square 2                                |

# Value

returned delta, Lower and Upper limits

### Author(s)

John Lawson

```
vci(.90,.05,.014852,2,.05,.026885,18)
## The function is currently defined as
vci<-function(confl,c1,ms1,nu1,c2,ms2,nu2){</pre>
  delta<-c1*ms1-c2*ms2
  alpha<-1-confl
  Falpha1<-qf(confl,nu1,10000000)
  Falpha12<-qf(confl,nu1,nu2)
  Fconf2<-qf(alpha,nu2,10000000)
  Fconf12<-qf(alpha,nu1,nu2)
  Falpha2<-qf(confl,nu2,10000000)
  Fconf1<-qf(alpha,nu1,10000000)
  Fconf12<-qf(alpha,nu1,nu2)
  G1<-1-(1/Falpha1)
  H2<-(1/Fconf2)-1
  G12<-((Falpha12-1)**2-G1**2*Falpha12**2-H2**2)/Falpha12
  VL<-G1**2*c1**2*ms1**2+H2**2*c2**2*ms2**2+G12*c1*c2*ms1*ms2
```

58 volt

```
H1<-(1/Fconf1)-1
G2<-1-(1/Falpha2)
H12<-((1-Fconf12)**2-H1**2*Fconf12**2-G2**2)/Fconf12
VU<-H1**2*c1**2*ms1**2+G2**2*c2**2*ms2**2
L<-delta-sqrt(VL)
U<-delta+sqrt(VU)
cat("delta=",delta," Lower Limit=",L," Upper Limit=",U,"\n")
}
```

virus

Assay of Viral Contamination experiment data from Chapter 3

# Description

Data from the Assay of Viral Contamination experiment in chapter 3 of Design and Analysis of Experiments with R

### Usage

```
data(virus)
```

#### **Format**

A data frame with 18 observations on the following 3 variables.

```
y a numeric vector

Sample a factor with levels 1 2 3 4 5 6

Dilution a factor with levels 3 4 5
```

### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

### **Examples**

```
data(virus)
```

volt

Volt meter experiment data from Chapter 3

### **Description**

Data from the Volt meter experiment in chapter 3 of Design and Analysis of Experiments with R

# Usage

```
data(volt)
```

web 59

#### **Format**

A data frame with 16 observations on the following 3 variables.

y a numeric vector

XA a numeric vector containing the coded levels (-1,1) of factor A

XB a numeric vector containing the coded levels (-1,1) of factor B

XC a numeric vector containing the coded levels (-1,1) of factor C

#### **Source**

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

# **Examples**

```
data(volt)
```

web

Web page design experiment data from Chapter 3

## **Description**

Data from the web page design experiment in chapter 3 of Design and Analysis of Experiments with R

### Usage

```
data(web)
```

# Format

A data frame with 36 observations on the following 6 variables.

A a factor with levels 1 2

B a factor with levels 1 2

C a factor with levels 1 2

D a factor with levels 1 2

visitors a numeric vector

signup a numeric vector

## Source

Design and Analysis of Experiments with R, by John Lawson, CRC/Chapman Hall

```
data(web)
```

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