

# Package ‘daewr’

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**Type** Package

**Title** Design and Analysis of Experiments with R

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**Maintainer** John Lawson <lawson@byu.edu>

**Description** Contains Data frames and functions used in the book “Design and Analysis of Experiments with R”.

**License** GPL-2

**Imports** lattice, FrF2, BsMD, graphics, grDevices, stats, stringi

**LazyLoad** yes

**LazyData** yes

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---

daewr-package      *Data frames and functions for Design and Analysis of Experiments  
with R*

---

### 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)**

John Lawson

Maintainer: John Lawson <lawson@byu.edu>

**References**

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

**Examples**

```
Fcrit(.05,2,15)
Fpower1(alpha=.05,nlev=3,nreps=4,Delta=3,sigma=sqrt(2.1))
BIBsize(6,3)
```

---

Altscreen

*Alternate 16 run screening designs*

---

**Description**

Recalls Jones and Montgomery's 16 run screening designs from data frames

**Usage**

```
Altscreen(nfac, randomize=FALSE)
```

**Arguments**

nfac           input- an integer  
randomize       input - logical

**Value**

a data frame containing the alternate screening design

**Author(s)**

John Lawson

**References**

Jones, B. and Montgomery, D. C. (2010) "Alternatives to resolution IV screening designs in 16 runs", Int. J. Experimental Design and Process Optimization, Vol 1, No. 4, 2010.

**Examples**

```
Altscreen(6)  
Altscreen(6, randomize=TRUE)
```

---

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

**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 variance 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

**Examples**

```
data(Apo)
```

---

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)
```

**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

**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

**Examples**

```
data(Bdish)
```

---

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	<i>mouse liver enzyme experiment</i>
-----	--------------------------------------

---

**Description**

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

**Usage**

```
data(bha)
```

**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	<i>Balanced incomplete blocksize</i>
---------	--------------------------------------

---

**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

**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)
}

#cat("lambda=",lambda)
}
cat("Possible BIB design with b=",b," and r=",r," lambda=",lambda,"\n")
}

```

---

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

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 34 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 3

Subject a factor with levels 1 2 3

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

**Examples**

```
data(bioeqv)
```

---

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<sup>4</sup> in chapter 3 of Design and Analysis of Experiments with 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

**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 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

**Examples**

```
data(BPmonitor)
```

---

bread	<i>Bread rise experiment data from Chapter 2</i>
-------	--

---

**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)
```

---

bstep	<i>This function performs Tukey's single degree of freedom test for interaction in an unreplicated two-factor design</i>
-------	--

---

**Description**

This function removes the term with the highest p-value from a model already created by ihstep or fhstep.

**Usage**

```
bstep(y, des, prvm)
```

**Arguments**

y	input - this is a data frame containing a single numeric column of response data.
des	input - this is a data frame containing the numeric columns of the candidate independent variables.
prvm	input - this is a vector of text names of the terms in the model. This is created as the value resulting from running ihstep or fhstep.

**Value**

returned vector of terms entered in the model at this step.

**Author(s)**

John Lawson

**Examples**

```
library(daewr)
des <- DefScreen( m = 8 )
pd<-c(5.35,4.4,12.91,3.79,4.15,14.05,11.4,4.29,3.56,11.4,10.09,5.9,9.54,4.53,3.919,8.1,5.35)
trm<-ihstep(pd,des)
trm<-fhstep(pd,des,trm)
trm<-fhstep(pd,des,trm)
trm<-fhstep(pd,des,trm)
trm<-bstep(pd,des,trm)
trm<-bstep(pd,des,trm)
```

---

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 6 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

**Source**

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

**Examples**

```
data(cake)
```

---

```
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

**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 3 4 5 6 7 8 9 10 11 12

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)
```

---

COdata

*CO emmissions 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)
```

---

colormap

*This function makes a colormap of correlations in a design matrix*


---

### Description

This function makes a colormap of the correlations of a design matrix stored in the data frame design

### Usage

```
colormap(design, mod)
```

### Arguments

design	input - a data frame containing columns of the numeric factor levels
mod	input - a number indicating the model for the colormap 1 = linear model containing only the terms in the dataframe 2 = linear model plus two factor interactions 3 = linear model plus 2 and 3 factor interactions 4 = linear model plus 2, 3, and 4 factor interactions

### Author(s)

John Lawson

### Examples

```
# color map of 2^(4-1) design
library(FrF2)
design <- FrF2(8, 4, randomize = FALSE)
colormap(design, mod=3)

# Makes color map for saturated 2^(7-4) design in Figure 6.14 p. 197
library(FrF2)
design <-FrF2( 8, 7)
colormap(design, mod=2)

# Makes colormap of an Alternate Screening Design
library(daewr)
ascr<-AltScreen(7)
colormap(ascr, mod=2)

# Makes colormap of a Model Robust Design
library(daewr)
MR16 <- ModelRobust('MR16m7g5', randomize = FALSE)
colormap(MR16, mod=2)

## The function is currently defined as
function(design,mod) {
##### Inputs #####
```

```

# design - a data frame containing columns of the numeric factor levels
# mod - the model for the color plot of correlations
#   1 = Linear model containing only the terms in the data frame
#   2 = Linear model plus two factor interactions
#   3 = Linear model plus 2 and 3 factor interactions
#   4 = Linear model plus 2, 3 and 4 factor interactions
#####
y<-runif(nrow(design),0,1)
if(mod==1) {test <- model.matrix(lm(y~.),data=design)}
if(mod==2) {test <- model.matrix(lm(y~.^2,data=design)}
if(mod==3) {test <- model.matrix(lm(y~.^3,data=design)}
if(mod==4) {test <- model.matrix(lm(y~.^4,data=design)}
names<-colnames(test)
names<-gsub(':',',',names)
names<-gsub('1',',',names)
colnames(test)<-names
cmas<-abs(cor(test[,ncol(test):2]))
cmas<-cmas[c((ncol(cmas)):1), ]
rgb.palette <- colorRampPalette(c("white", "black"), space = "rgb")
levelplot(cmas, main="Map of absolute correlations", xlab="", ylab="", col.regions=rgb.palette(120),
          cuts=100, at=seq(0,1,0.01),scales=list(x=list(rot=90))) }

```

---

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
- lns2 a numeric vector

## 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

**Examples**

```
data(cpipe)
```

---

culture	<i>paecilomyces variotii culture experiment</i>
---------	---

---

**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)
```

**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)
```

---

DefScreen

*Definitive Screening Designs*

---

**Description**

Recalls Jones and Nachtshiem's Definitive screening designs for 3-level factors and 3-level factors with added 2-level categorical factors.

**Usage**

```
DefScreen(m, c=0, randomize=FALSE)
```

**Arguments**

m	input- an integer, the m=number of 3-level factors
c	input- an integer, the m=number of 2-level categorical factors, default is zero if not supplied
randomize	input - logical

**Value**

a data frame containing the definitive screening design with 3-level factors first followed by 2-level factors.

**Author(s)**

John Lawson

**References**

Jones, B. and Nachtsheim, C. J. (2011) "A Class of Three Level Designs for Definitive Screening in the Presence of Second-Order Effects", Journal of Quality Technology, Vol 43, No. 1, 2011, pp 1-15. Jones, B. and Nachtsheim, C. J. (2013) "Definitive Screening Designs with Added Two-Level Categorical Factors", Journal of Quality Technology, Vol 44, No. 2, 2013, pp. 121-129.

**Examples**

```
DefScreen(m=8,c=2)
DefScreen(12)
DefScreen(m=4,c=4, randomize=TRUE)
```

---

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

**Examples**

```
data(drug)
```

---

EEw1s1

*D-efficient Estimation Equivalent Response Surface Designs*

---

**Description**

Recalls Jones and Goos JQT Estimation Equivalent Response Surface Designs for 1 whole plot factor and 1 sub-plot factor from a catalog

**Usage**

```
EEw1s1(des, randomize=FALSE)
```

**Arguments**

des	input- a character variable containing the name of a design in the catalog. If left blank, the function prints a table showing all the design names in the catalog
randomize	input- a logical

**Value**

design

**Author(s)**

John Lawson

**References**

Jones, B. and Goos, P.(2012) "An Algorithm for Finding D-Efficient Equivalent-Estimation Second-Order Split Plot Designs", Journal of Quality Technology, Vol 44, No. 4, pp281-303, 2012.

**Examples**

```
EEw1s1()  
EEw1s1('EE8R4WP')  
EEw1s1('EE8R4WP', randomize=TRUE)
```

---

 EEw1s2

*D-efficient Estimation Equivalent Response Surface Designs*


---

**Description**

Recalls Jones and Goos JQT Estimation Equivalent Response Surface Designs for 1 whole plot factor and 2 sub-plot factors from a catalog

**Usage**

```
EEw1s2(des, randomize=FALSE)
```

**Arguments**

des	input- a character variable containing the name of a design in the catalog. If left blank, the function prints a table showing all the design names in the catalog
randomize	input- a logical

**Value**

design

**Author(s)**

John Lawson

**References**

Jones, B. and Goos, P.(2012) "An Algorithm for Finding D-Efficient Equivalent-Estimation Second-Order Split Plot Designs", Journal of Quality Technology, Vol 44, No. 4, pp281-303, 2012.

**Examples**

```
EEw1s2( )
EEw1s2('EE12R4WP')
EEw1s2('EE12R4WP', randomize=TRUE)
EEw1s2('EE12R6WP')
EEw1s2('EE12R6WP', randomize=TRUE)
EEw1s2('EE14R7WP')
EEw1s2('EE14R7WP', randomize=TRUE)
EEw1s2('EE15R5WP')
EEw1s2('EE15R5WP', randomize=TRUE)
EEw1s2('EE16R4WP')
EEw1s2('EE16R4WP', randomize=TRUE)
EEw1s2('EE18R6WP')
EEw1s2('EE18R6WP', randomize=TRUE)
EEw1s2('EE20R4WP')
EEw1s2('EE20R4WP', randomize=TRUE)
EEw1s2('EE20R5WP')
```

```

EEw1s2('EE20R5WP', randomize=TRUE)
EEw1s2('EE21R7WP')
EEw1s2('EE21R7WP', randomize=TRUE)
EEw1s2('EE24R4WP')
EEw1s2('EE24R4WP', randomize=TRUE)
EEw1s2('EE24R6WP')
EEw1s2('EE24R6WP', randomize=TRUE)
EEw1s2('EE25R5WP')
EEw1s2('EE25R5WP', randomize=TRUE)
EEw1s2('EE28R7WP')
EEw1s2('EE28R7WP', randomize=TRUE)
EEw1s2('EE30R6WP')
EEw1s2('EE30R6WP', randomize=TRUE)
EEw1s2('EE30R5WP')
EEw1s2('EE30R5WP', randomize=TRUE)
EEw1s2('EE35R7WP')
EEw1s2('EE35R7WP', randomize=TRUE)
EEw1s2('EE36R6WP')
EEw1s2('EE36R6WP', randomize=TRUE)
EEw1s2('EE42R7WP')
EEw1s2('EE42R7WP', randomize=TRUE)

```

---

EEw1s3

*D-efficient Estimation Equivalent Response Surface Designs*


---

### Description

Recalls Jones and Goos JQT Estimation Equivalent Response Surface Designs for 1 whole plot factor and 3 sub-plot factors from a catalog

### Usage

```
EEw1s3(des, randomize=FALSE)
```

### Arguments

des	input- a character variable containing the name of a design in the catalog. If left blank, the function prints a table showing all the design names in the catalog
randomize	input- a logical

### Value

design

### Author(s)

John Lawson

**References**

Jones, B. and Goos, P.(2012) "An Algorithm for Finding D-Efficient Equivalent-Estimation Second-Order Split Plot Designs", Journal of Quality Technology, Vol 44, No. 4, pp281-303, 2012.

**Examples**

```
EEw1s3()
EEw1s3('EE16R4WP')
EEw1s3('EE16R4WP', randomize=TRUE)
```

---

 EEw2s1

---

*D-efficient Estimation Equivalent Response Surface Designs*


---

**Description**

Recalls Jones and Goos JQT Estimation Equivalent Response Surface Designs for 2 whole plot factors and 1 sub-plot factor from a catalog

**Usage**

```
EEw2s1(des, randomize=FALSE)
```

**Arguments**

des	input- a character variable containing the name of a design in the catalog. If left blank, the function prints a table showing all the design names in the catalog
randomize	input- a logical

**Value**

design

**Author(s)**

John Lawson

**References**

Jones, B. and Goos, P.(2012) "An Algorithm for Finding D-Efficient Equivalent-Estimation Second-Order Split Plot Designs", Journal of Quality Technology, Vol 44, No. 4, pp281-303, 2012.

**Examples**

```
EEw2s1()
EEw2s1('EE21R7WP')
EEw1s1('EE21R7WP', randomize=TRUE)
```

**Description**

Recalls Jones and Goos JQT Estimation Equivalent Response Surface Designs for 2 whole plot factors and 1 sub-plot factor from a catalog

**Usage**

```
EEw2s2(des, randomize=FALSE)
```

**Arguments**

des	input- a character variable containing the name of a design in the catalog. If left blank, the function prints a table showing all the design names in the catalog
randomize	input- a logical

**Value**

design

**Author(s)**

John Lawson

**References**

Jones, B. and Goos, P.(2012) "An Algorithm for Finding D-Efficient Equivalent-Estimation Second-Order Split Plot Designs", Journal of Quality Technology, Vol 44, No. 4, pp281-303, 2012.

**Examples**

```
EEw2s2()  
EEw2s2('EE21R7WP')  
EEw1s2('EE21R7WP', randomize=TRUE)
```

---

EEw2s3

*D-efficient Estimation Equivalent Response Surface Designs*

---

### Description

Recalls Jones and Goos JQT Estimation Equivalent Response Surface Designs for 2 whole plot factors and 1 sub-plot factor from a catalog

### Usage

```
EEw2s3(des, randomize=FALSE)
```

### Arguments

des	input- a character variable containing the name of a design in the catalog. If left blank, the function prints a table showing all the design names in the catalog
randomize	input- a logical

### Value

design

### Author(s)

John Lawson

### References

Jones, B. and Goos, P.(2012) "An Algorithm for Finding D-Efficient Equivalent-Estimation Second-Order Split Plot Designs", Journal of Quality Technology, Vol 44, No. 4, pp281-303, 2012.

### Examples

```
EEw2s3()  
EEw2s3('EE24R8WP')  
EEw1s3('EE24R8WP', randomize=TRUE)
```

**Description**

Recalls Jones and Goos JQT Estimation Equivalent Response Surface Designs for 3 whole plot factors and 1-2 sub-plot factors from a catalog

**Usage**

```
EEw3(des, randomize=FALSE)
```

**Arguments**

des	input- a character variable containing the name of a design in the catalog. If left blank, the function prints a table showing all the design names in the catalog
randomize	input- a logical

**Value**

design

**Author(s)**

John Lawson

**References**

Jones, B. and Goos, P.(2012) "An Algorithm for Finding D-Efficient Equivalent-Estimation Second-Order Split Plot Designs", Journal of Quality Technology, Vol 44, No. 4, pp281-303, 2012.

**Examples**

```
EEw3()  
EEw3('EE22R11WP')  
EEw3('EE22R11WP', randomize=TRUE)  
EEw3('EE48R12WP')  
EEw3('EE48R12WP', randomize=TRUE)
```

---

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	<i>Control array and variance of response for silicon layer growth experiment</i>
---------	---

---

**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)
```

**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

### **Source**

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

### **Examples**

```
data(eptaxyb)
```

---

Fcrit	<i>F-Distribution critical values</i>
-------	---------------------------------------

---

**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

**Examples**

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

---

fhstep	<i>This function performs Tukey's single degree of freedom test for interaction in an unreplicated two-factor design</i>
--------	--

---

**Description**

This function performs a single step of a hierarchical forward stepwise regression by entering additional term(s) to a model already created by ihstep or fhstep. If an interaction or quadratic term is entered first, the parent main effects are also entered into the model.

**Usage**

```
fhstep(y, des, prvm)
```

**Arguments**

y	input - this is a data frame containing a single numeric column of response data.
des	input - this is a data frame containing the numeric columns of the candidate independent variables.
prvm	input - this is a vector of text names of the terms in the model. This is created as the value resulting from running ihstep or fhstep.

**Value**

returned vector of terms entered in the model at this step.

**Author(s)**

John Lawson

**Examples**

```
library(daewr)
des <- DefScreen( m = 8 )
pd<-c(5.35, 4.4, 12.91, 3.79, 4.15, 14.05, 11.4, 4.29, 3.56, 11.4, 10.09, 5.9, 9.54, 4.53, 3.919, 8.1, 5.35)
trm<-ihstep(pd, des)
trm<-fhstep(pd, des, trm)
trm<-fhstep(pd, des, trm)
trm<-fhstep(pd, des, trm)
```

---

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 numerator
nu2	input - degrees of freedom for denominator
nc	input - noncentrality parameter

**Value**

probability of exceeding  $f_{crit}(\alpha, \nu_1, \nu_2)$  with the non-central F-distribution with  $\nu_1$  and  $\nu_2$  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)
```

---

Fpower1

*F-Distribution Power Calculation*

---

**Description**

Calculates the power for one-way ANOVA

**Usage**

```
Fpower1(alpha,nlev,nreps,Delta,sigma)
```

**Arguments**

alpha	input - significance level of the F-test.
nlev	input - the number of levels of the factor
nreps	input - the number of replicates in each level of the factor.
Delta	input - the size of a practical difference in two cell means.
sigma	input - the standard deviation of the experimental error.

**Value**

probability of exceeding  $f_{crit}(\alpha, \nu_1, \nu_2)$  with the non-central F-distribution with  $\nu_1$  and  $\nu_2$  degrees of freedom and noncentrality parameter  $nc$

**Author(s)**

John Lawson

**Examples**

```

Fpower1(alpha=.05,nlev=3,nreps=4,Delta=3,sigma=sqrt(2.1))

rmin <-2 #smallest number of replicates considered
rmax <-6 # largest number of replicates considered
alpha <- rep(0.05, rmax - rmin +1)
sigma <-rep(sqrt(2.1), rmax - rmin +1)
nreps <-c(rmin:rmax)
nlev <- rep(3,rmax - rmin +1)
nreps <- rmin:rmax
Delta <- rep(3,rmax - rmin +1)
power <- Fpower1(alpha,nlev,nreps,Delta,sigma)
data.frame(r=nreps,Power=power)

## The function is currently defined as
Fpower1<-function(alpha=NULL, nlev=NULL,nreps=NULL, Delta=NULL, sigma=NULL)
{
##### Power Calculation for one way ANOVA #####
# Argument list
# alpha the significance level of the test
# nlev the number of levels of the factor
# nreps the number of replicates in each level of the factor
# Delta the size of a practical difference in two cell means
# sigma the standard deviation of the experimental error
#####
if (is.null(alpha)|is.null(nlev)|is.null(nreps)|is.null(Delta)|is.null(sigma))
  stop("you must supply alpha, nlev, nreps, Delta and sigma")
css<-(Delta^2)/2
nc<- (nreps*css)/(sigma^2)
df1<-nlev-1
df2<-(nreps-1)*nlev
power <- 1-pf(Fcrit(alpha,df1,df2),df1,df2,nc)
return(power)
}

```

---

Fpower2

*F-Distribution Power Calculation*


---

**Description**

Calculates the power for a two-way ANOVA

**Usage**

```
Fpower2(alpha,nlev,nreps,Delta,sigma)
```

**Arguments**

alpha	input - significance level of the F-test.
nlev	input - vector of length two containing the number of levels of the factors.
nreps	input - the the number of replicates in each combination of factor levels.
Delta	input - the size of a practical difference in two marginal factor level means.
sigma	input - the standard deviation of the experimental error.

**Value**

probability of exceeding  $f_{crit}(\alpha, \nu_1, \nu_2)$  with the non-central F-distribution with  $\nu_1$  and  $\nu_2$  degrees of freedom and noncentrality parameter  $nc$

**Author(s)**

John Lawson

**Examples**

```
power <- Fpower2(.05, nlev = c(4,4), nreps=2, Delta= 1, sigma=.32)

rmin <- 2 # smallest number of replicates
rmax <- 4 # largest number of replicates
alpha <- .05
sigma <- .32
Delta <- 1.0
nlev <- c(4,4)
nreps <- c(rmin:rmax)
result <- Fpower2(alpha, nlev, nreps, Delta, sigma)
options(digits = 5)
result

## The function is currently defined as
Fpower2<-function(alpha=NULL, nlev=NULL,nreps=NULL, Delta=NULL, sigma=NULL)
{
##### Power Calculation for two way ANOVA #####
# Argument list
# alpha the significance level of the test.
# nlev vector containing the number of levels of the factors.
# nreps the number of replicates in each combination of factor levels.
# Delta the size of a practical difference in two marginal factor level means.
# sigma the standard deviation of the experimental error.
#####
if (is.null(alpha)|is.null(nlev)|is.null(nreps)|is.null(Delta)|is.null(sigma))
  stop("you must supply alpha, nlev, nreps, Delta and sigma")
if(length(nlev)<2)
  stop ("nlev must be a two component vecto containing levels of the 1st and 2nd factors")
a <- nlev[1]
b <- nlev[2]
cssb <- (Delta^2)/2
```

```

ncb <- a*(nreps*cssb)/(sigma^2)
cssa<-(Delta^2)/2
nca<- b*(nreps*cssa)/(sigma^2)
dfa<- a-1
dfb<- b-1
df2<-(nreps-1)*b*a
powera <- 1-pf(Fcrit(alpha,dfa,df2),dfa,df2,nca)
powerb <- 1-pf(Fcrit(alpha,dfb,df2),dfa,df2,nca)
result <-cbind(nreps,df2,powera,powerb)
}

```

---

fullnormal	<i>This function makes a full normal plot of the elements of the vector called effects</i>
------------	--

---

### Description

This function makes a full normal plot of the elements of the vector called effects

### Usage

```
fullnormal(effects, labs, alpha = 0.05, reffline = "TRUE")
```

### Arguments

effects	input - vector of effects to be plotted
labs	input - vector of labels of the effects to be plotted
alpha	input - alpha level for labeling of significant effects using Lenth statistic
reffline	input - logical variable that indicates whether a reference line is added to the plot (default is "TRUE")

### Author(s)

John Lawson

### Examples

```

# Example Separate Normal plots of whole and split plot effects from an unreplicated split-plot
data(plasma)
sol<-lm(y~A*B*C*D*E,data=plasma)
summary(sol)
# 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)]
Speffects<-effects[c(5,12:15,20:25,27:31)]

#make separate normal plots

```

```

library(BsMD)
fullnormal(Wpeffects,names(Wpeffects),alpha=.10)
fullnormal(Speffects,names(Speffects),alpha=.05)

## The function is currently defined as
function (effects, labs, alpha = 0.05, refline = "TRUE")
{
  crit <- LenthPlot(effects, alpha = alpha, plt = FALSE)["ME"]
  names <- names(effects)
  names <- gsub(":", "", names)
  names <- gsub("1", "", names)
  le <- length(effects)
  for (i in 1:le) {
    logc <- (abs(effects[i]) <= crit)
    if (logc) {
      names[i] <- " "
    }
  }
  qqnorm(effects, ylab = "Estimated Effects", xlab = "Normal Scores")
  x <- qqnorm(effects, plot = FALSE)
  zscr <- (x$x)
  effp <- effects[zscr > 0]
  zp <- zscr[zscr > 0]
  namep <- names[zscr > 0]
  effn <- effects[zscr < 0]
  zn <- zscr[zscr < 0]
  namen <- names[zscr < 0]
  text(zp, effp, namep, pos = 1)
  text(zn, effn, namen, pos = 3)
  ahe <- abs(effects)
  s0 <- 1.5 * median(ahe)
  selhe <- ahe < (2.5 * s0)
  pse = 1.5 * median(ahe[selhe])
  if (refline) {
    abline(0, pse)
  }
}

```

**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

**Examples**

```
data(gagerr)
```

---

gapstat	<i>This function computes the gap statistic which is used to test for an outlier using Daniels method</i>
---------	---

---

**Description**

This function computes the gap statistic which is used to test for an outlier using Daniels method

**Usage**

```
gapstat(beta, pse)
```

**Arguments**

beta	input - vector of coefficients from saturated model fit to the data
pse	input - Lenth's PSE statistic calculated from the elements of beta

**Value**

returned gap statistic

**Author(s)**

John Lawson

**Examples**

```

## The function is currently defined as
function (beta, pse)
{
  p <- length(beta)
  psehe <- pse
  sel <- beta >= 0
  betap <- beta[sel]
  betap <- sort(betap)
  betas <- betap[1]
  sel <- beta < 0
  betan <- beta[sel]
  nn <- length(betan)
  betan <- sort(betan)
  betal <- betan[nn]
  z1 <- qnorm((nn - 0.375)/(p + 0.25))
  zs <- qnorm((nn + 1 - 0.375)/(p + 0.25))
  gap <- ((betas - betal)/psehe)/(zs - z1)
  return(gap)
}

```

---

Gaptest

*This function uses Daniel's Method to find an outlier in an unrepliated  $2^{(k-p)}$  design.*

---

**Description**

This function uses Daniel's Method to find an outlier in an unrepliated  $2^{(k-p)}$  design.

**Usage**

Gaptest(DesY)

**Arguments**

DesY           input this is a data frame containing an unrepliated  $2^{(k-p)}$  design. The last variable in the data frame should be the numeric response.

**Author(s)**

John Lawson

**References**

Box, G.E.P. (1991) "George's column: Finding bad values in factorial designs", Quality Engineering, 3, 249-254.

**Examples**

```

# Example from Box(1991)
data(BoxM)
Gaptest(BoxM)

## The function is currently defined as
function (DesY)
{
  ncheck <- dim(DesY)
  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
    critg16 <- c(1.7884, 5.1009)
    critg32 <- c(1.7297, 5.8758)
    modf <- lm(y ~ (.)^4, x = TRUE, data = DesY)
    nbeta <- dim(DesY)
    nbeta <- nbeta[1]
    he <- modf$coef
    selcol <- which(!is.na(he))
    he <- he[selcol]
    he <- he[-1]
    p <- length(he)
    n <- p + 1
    cn1 <- names(he)
    ccn1 <- gsub("[^A-Z]", "", cn1)
    names(he) <- ccn1
    ahe <- abs(he)
    s0 <- 1.5 * median(ahe)
    selhe <- ahe < (2.5 * s0)
    pse = 1.5 * median(ahe[selhe])
    gap <- gapstat(he, pse)
    if (ncheck == 16) {
      test = (gap > critg16[1])
    }
    else {
      test = (gap > critg32[1])
    }
  }
}

```

```

}
if (test) {
  X <- modf$x
  X <- X[, selcol]
  X <- X[, -1]
  se <- as.matrix(sign(he), nrow = 1)
  sigef <- LGB(he, rpt = FALSE, plt = FALSE)
  for (i in 1:length(he)) {
    if (sigef[i] == "yes") {
      se[i] = 0
    }
  }
  sp <- X %**% se
  asp <- abs(sp)
  oo <- max.col(t(asp))
  ae <- abs(he)
  sae <- sort(ae)
  nsmall <- round(length(he)/2)
  bias <- 2 * sum(sae[1:nsmall])
  y <- DesY$y
  ycorr <- DesY$y
  ycorr[oo] <- ycorr[oo] + (-1 * sign(sp[oo])) * bias
  detect <- c(rep("no", n))
  detect[oo] <- "yes"
  cat("Initial Outlier Report", "\n")
  cat("Standardized-Gap = ", gap, "Significant at 50th percentile",
      "\n")
  DesYc <- cbind(DesY[, 1:(dim(DesY)[2] - 1)], ycorr)
  modf <- lm(ycorr ~ (.)^4, x = TRUE, data = DesYc)
  che <- modf$coef
  che <- che[!is.na(che)]
  che <- che[-1]
  p <- length(che)
  n <- p + 1
  cn <- names(che)
  ccn <- gsub("[^A-Z]", "", cn)
  names(che) <- ccn
  ache <- abs(che)
  s0 <- 1.5 * median(ache)
  selche <- ache < (2.5 * s0)
  psec = 1.5 * median(ache[selche])
  gap <- gapstat(he, psec)
  if (ncheck == 16)
    test2 = (gap > critg16[2])
  else test2 = (gap > critg32[2])
  if (test2) {
    cat("Final Outlier Report", "\n")
    cat("Standardized-Gap = ", gap, "Significant at 99th percentile",
        "\n")
    cat(" ", "\n")
    cat(" Corrected 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 = "")
        tce <- LGB(che)
    }
    else {
        cat("Final Outlier Report", "\n")
        cat("No significant outlier detected in second pass",
            "\n")
        LGB(he)
        cat("      ", "\n")
    }
}
}
}

```

---

gear	<i>Unreplicated split-plot fractional-factorial experiment on geometric distortion of drive gears</i>
------	---

---

### 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

### Examples

```
data(gear)
```

---

halfnorm	<i>This function makes a half normal plot of the elements of the vector called effects</i>
----------	--

---

## Description

This function makes a half normal plot of the elements of the vector called effects

## Usage

```
halfnorm(effects, labs, alpha = 0.05, refline = "TRUE")
```

## Arguments

effects	input - vector of effects to be plotted
labs	input - vector of labels of the effects to be plotted
alpha	input - alpha level for labeling of significant effects using Lenth statistic
refline	input - logical variable that indicates whether a reference line is added to the plot (default is "TRUE")

## Author(s)

John Lawson

## Examples

```
# Example Separate Normal plots of whole and split plot effects from an unreplicated split-plot
data(plasma)
sol<-lm(y~A*B*C*D*E,data=plasma)
# 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)]
Speffects<-effects[c(5,12:15,20:25,27:31)]

#make separate half normal plots
library(BsMD)
halfnorm(Wpeffects,names(Wpeffects),alpha=.10)
halfnorm(Speffects,names(Speffects),alpha=.05)
```

---

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)
```

---

ihstep

*This function performs Tukey's single degree of freedom test for interaction in an unreplicated two-factor design*

---

**Description**

This function performs the first step of a hierarchical forward stepwise regression. If an interaction or quadratic term is entered first, the parent main effects are also entered into the model.

**Usage**

```
ihstep(y, des)
```

**Arguments**

y input - this is a data frame containing a single numeric column of response data.  
des input - this is a data frame containing the numeric columns of the candidate independent variables.

**Value**

returned vector of terms entered in the model at this step.

**Author(s)**

John Lawson

**Examples**

```
library(daewr)
des <- DefScreen( m = 8 )
pd<-c(5.35,4.4,12.91,3.79,4.15,14.05,11.4,4.29,3.56,11.4,10.09,5.9,9.54,4.53,3.919,8.1,5.35)
trm<-ihstep(pd,des)
```

---

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)
```

**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)
```

---

LGB

*This function uses the LGB Method to detect significant effects in unreplicated fractional factorials.*

---

**Description**

This function uses the LGB Method to detect significant effects in unreplicated fractional factorials.

**Usage**

```
LGB(Beta, alpha = 0.05, rpt = TRUE, plt = TRUE, plt1 = TRUE)
```

**Arguments**

Beta	input - this is the numeric vector of effects or coefficients to be tested
alpha	input - This is the significance level of the test
rpt	input - this is a logical variable that controls whether the report is written (default is TRUE)
plt	input - this is a logical variable that controls whether a half-normal plot is made (default is TRUE)
plt1	input - this is a logical variable that controls whether the significance limit line is drawn on the half-normal plot (default is TRUE)

**Author(s)**

John Lawson

**References**

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

**Examples**

```

data(chem)
modf<-lm(y~A*B*C*D,data=chem)
LGB(coef(modf)[-1],rpt=FALSE)

## The function is currently defined as
LGB <- function(Beta, alpha=.05,rpt=TRUE, plt=TRUE, pltl=TRUE) {
  sigLGB<-LGBc(Beta,alpha,rpt,plt,pltl)
}

```

LGBc

---

*This function does the calculations for the LGB Method to detect significant effects in unrepliated fractional factorials.*

---

**Description**

This function uses the LGB Method to detect significant effects in unrepliated fractional factorials.

**Usage**

```
LGBc(Beta, alpha = 0.05, rpt = TRUE, plt = TRUE, pltl = TRUE)
```

**Arguments**

Beta	input - this is the numeric vector of effects or coefficients to be tested
alpha	input - This is the significance level of the test
rpt	input - this is a logical variable that controls whether the report is written (default is TRUE)
plt	input - this is a logical variable that controls whether a half-normal plot is made (default is TRUE)
pltl	input - this is a logical variable that controls whether the significance limit line is drawn on the half-normal plot (default is TRUE)

**Author(s)**

John Lawson

**References**

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

## Examples

```

data(chem)
modf<-lm(y~A*B*C*D,data=chem)
sig<-LGBc(coef(modf)[-1],rpt=FALSE)

## The function is currently defined as
function (Beta, alpha = 0.05, rpt = TRUE, plt = TRUE, pltl = TRUE)
{
  siglev <- c(0.1, 0.05, 0.025, 0.01)
  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, 1.11, 1.106,
    1.072, 1.063, 1.06, 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, 1.26,
    1.165, 1.14, 1.13, 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)
  colind <- which(siglev == alpha, arr.ind = TRUE)
  if (length(colind) == 0) {
    stop("this function works only when alpha= .1, .05, .025 or .01")
  }
  rowind <- which(df == length(Beta), arr.ind = TRUE)
  if (length(rowind) == 0) {
    stop("this function works only for coefficient vectors of
length 7,8,11,15,16,26,31,32,35,63,or 127")
  }
  critL <- crittab[rowind, colind]
  acj <- abs(Beta)
  ranks <- rank(acj, ties.method = "first")
  s0 <- 1.5 * median(acj)
  p <- (ranks - 0.5)/length(Beta)
  z <- qnorm((p + 1)/2)
  moda <- lm(acj ~ -1 + z)
  beta1 <- moda$coef
  sel <- acj < 2.5 * s0
  modi <- lm(acj[sel] ~ -1 + z[sel])
  beta2 <- modi$coef
  Rn <- beta1/beta2
  pred <- beta2 * z
  n <- length(acj[sel])
  df <- n - 1
  sig <- sqrt(sum(modi$residuals^2)/df)
  se.pred <- sig * (1 + 1/n + (z^2)/sum(z[sel]^2))^0.5
  pred.lim <- pred + qt(0.975, df) * se.pred
  sigi <- c(rep("no", length(Beta)))
  sel2 <- acj > pred.lim
  sigi[sel2] <- "yes"
  if (plt) {
    plot(z, acj, xlab = "Half Normal Scores", ylab = "Absolute 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 (plt1) {
        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 integer
b	input - an integer

### Value

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

### Author(s)

John Lawson

**Examples**

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

---

ModelRobust

*Model Robust Factorial Designs*

---

**Description**

Recalls Li and Nachtsheim's model robust factorial designs from a catalog of data frames

**Usage**

```
ModelRobust(des, randomize=FALSE)
```

**Arguments**

des	input- a character variable containing the name of a design in the catalog. If left blank, the function prints a table showing all the design names in the catalog
randomize	input- a logical

**Value**

design

**Author(s)**

John Lawson

**References**

Li, W. and Nachtsheim, C. J. (2000) "Model Robust factorial Designs", Technometrics, Vol 42, No. 4, pp345-352, 2000.

**Examples**

```
ModelRobust()
ModelRobust('MR8m4g3')
ModelRobust('MR8m4g3', randomize=TRUE)
```

---

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

**Examples**

```
data(MPV)
```

---

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)
```

---

OptPB

*Optimum Plackett-Burman Designs*

---

**Description**

Selects the columns from a Plackett-Burman Design Produced by FrF2 that will minimize model dependence for main effects and two factor interactions

**Usage**

```
OptPB(nruns, nfactors, randomize=FALSE)
```

**Arguments**

nruns	input- an integer representing the number of runs in the design
nfactors	input - in integer representing the number of factors in the design
randomize	input - logical

**Value**

design

**Author(s)**

John Lawson

**References**

Fairchild, K. (2011) "Screening Designs that Minimize Model Dependence", MS Project Department of Statistics Brigham Young University, Dec. 2011.

**Examples**

```
OptPB(12,8)
```

---

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

**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	<i>pesticide application experiment</i>
-----------	---

---

**Description**

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

**Usage**

```
data(pesticide)
```

**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	<i>Unreplicated split-plot 2<sup>5</sup> experiment on plasma treatment of paper</i>
--------	--

---

**Description**

Data from the unreplicated split-plot 2<sup>5</sup> 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

**Examples**

```
data(plasma)
```

---

polvdat	<i>Polvoron mixture experiment</i>
---------	------------------------------------

---

**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	<i>polymerization strength variability study</i>
---------	--

---

**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

**Source**

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

**Examples**

```
data(polymer)
```

---

prodstd	<i>Complete control factor array and noise factor array for connector experiment</i>
---------	--

---

**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)
```

---

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

teehtgt a factor with levels 1 2 3

cdistance a numeric vector

**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

**Examples**

```
data(residue)
```

---

rubber	<i>Rubber Elasticity data</i>
--------	-------------------------------

---

**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	<i>Split-plot experiment on sausage casing with RCB in whole plot</i>
---------	---

---

**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

**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

**Examples**

```
data(Smotor)
```

---

soup	<i>dry mix soup experiment</i>
------	--------------------------------

---

**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	<i>dry soup mix variance component study</i>
--------	--

---

**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

**Source**

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

**Examples**

```
data(soupmx)
```

---

splitPdes	<i>Split-plot cookie baking experiment</i>
-----------	--

---

**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

**Examples**

```
data(splitPdes)
```

---

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)
```

---

step.forward	<i>RSM forward regression keeping model hierarchy</i>
--------------	---

---

### Description

This function performs a hierarchical forward stepwise regression. If an interaction or quadratic term is entered in the model, the parent main effects are also entered into the model.

### Usage

```
step.forward(y,x,step)
```

### Arguments

y	input - this is a vector containing a single numeric column of response data.
x	input - this is a data frame containing the numeric columns of the candidate independent variables.
step	input - this is a single numeric value containing the number of steps requested.

### Value

returned data frame the first column is a factor variable containing the formula for the model fit at each step, the second numeric column is the R-square statistic for the model fit with each formula.

### Author(s)

Gerhard Krennrich

### Examples

```
#Definitive Screening Design
library(daewr)
set.seed(1234)
x <- DefScreen(m=5,c=0)
colnames(x) <- paste("x",1:5,sep="")
x$y <- 3*x$x1 + 2*x$x2 + 2*x$x4*x$x5 + x$x3^2 + 2*x$x1^2 + rnorm(nrow(x),0,1)
(z <- step.forward(x$y,x[,-6], 4 ))
# Example p. 240 Design and Analysis of Experiments with R PB Design
library(BsMD)
data(PB12Des,package="BsMD")
colnames(PB12Des)<-c("c11","c10","c9","c8","G","F","E","D","C","B","A")
#Reorder the columns to match Table 6.11
castf<-PB12Des[c(11,10,9,8,7,6,5,4,3,2,1)]
y<-c(4.733,4.625,5.899,7.0,5.752,5.682,6.607,5.818,5.917,5.863,6.058,4.809)
castf<-cbind(castf,y)
castfr <- castf[ , c(1:7, 12)]
library(daewr)
des<-castfr[ ,c(1:7)]
```

```
y<-castfr[ , 8]
step.forward(y,des,2)
```

---

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)
```

### 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 experiment*


---

### 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

**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

**Source**

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

**Examples**

```
data(taste)
```

---

teach	<i>Teaching experiment data from Chapter 2</i>
-------	--

---

**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	<i>Tetracycline concentration in plasma</i>
-----	---

---

**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)
```

**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	<i>Control factor array and summary statistics for Ina tile experiment</i>
------	--

---

**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

lns2 a numeric vector

**Source**

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

**Examples**

```
data(tile)
```

---

Treb	<i>Box-Behnken design for trebuchet experiment</i>
------	--

---

**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	<i>This function performs Tukey's single degree of freedom test for interaction in an unreplicated two-factor design</i>
----------	--

---

**Description**

This function performs Tukey's single degree of freedom test for interaction in an unreplicated two-factor design

**Usage**

```
Tukey1df(data)
```

**Arguments**

`data` input - this is a data frame with three variables, the first variable is a numeric response and next two variables are factors. There should be  $ab$  lines in the data frame where  $a$  is the number of levels of the first factor, and  $b$  is the number of levels of the second factor.

**Author(s)**

John Lawson

**Examples**

```
library(daewr)
data(virus)
Tukey1df(virus)

## The function is currently defined as
function (data)
{
  y <- data[, 1]
  Afactor <- data[, 2]
  Bfactor <- data[, 3]
  tst1 <- is.factor(Afactor)
  tst2 <- is.factor(Bfactor)
  tst3 <- is.numeric(y)
  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 be coded as factors")
  }
  tst4 <- max(a, b) > 2
  tst5 <- length(y) == a * b
  if (tst4 & tst5) {
    ybb <- with(data, tapply(y, Bfactor, mean))
    yba <- with(data, tapply(y, Afactor, mean))
    sbb <- with(data, tapply(y, Bfactor, sum))
    sba <- with(data, tapply(y, Afactor, sum))
    ybardd <- mean(y)
    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)
    prody <- y * ybdj
    sumprod <- tapply(prody, Afactor, sum)
    leftsum <- sum(sumprod * yba)
    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(F, 1, ((a - 1) * (b - 1) - 1))
cat("Source      df      SS      MS      F      Pr>F",
    "\n")
cat("A          ", paste(format(a - 1, width = 6),
    " ", format(round(ssA, 4), justify = "right"), " ",
    format(round(ssA/(a - 1), 4), justify = "right"),
    "\n"), sep = "")
cat("B          ", paste(format(b - 1, width = 6),
    " ", format(round(ssB, 4), justify = "right"), " ",
    format(round(ssB/(b - 1), 4), justify = "right"),
    "\n"), sep = "")
cat("Error       ", paste(format((b - 1) * (a - 1),
    width = 6), " ", format(round(ssE, 4), justify = "right"),
    " ", format(round(ssE/((a - 1) * (b - 1)), 4), justify = "right"),
    "\n"), sep = "")
cat("NonAdditivity", paste(format(1, width = 6), " ",
    format(round(ssAB, 4), justify = "right"), " ",
    format(round(ssAB, 4), justify = "right"), " ",
    format(round(F, 2), justify = "right"), " ", format(round(Pval,
    4), justify = "right"), "\n"), sep = "")
cat("Residual    ", paste(format((b - 1) * (a - 1) -
    1, width = 6), " ", format(round(ssR, 4), justify = "right"),
    " ", format(round(ssR/((a - 1) * (b - 1) - 1), 4),
    justify = "right"), "\n"), sep = "")
}
else {
  stop("This function only works for unreplicated 2-factor
factorials with >2 levels for one of the factors")
}
}

```

---

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(conf1, c1, ms1, nu1, c2, ms2, nu2)
```

### Arguments

conf1            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

**Examples**

```
vci(.90,.05,.014852,2,.05,.026885,18)
## The function is currently defined as
vci<-function(conf1,c1,ms1,nu1,c2,ms2,nu2){
  delta<-c1*ms1-c2*ms2
  alpha<-1-conf1
  Falpha1<-qf(conf1,nu1,10000000)
  Falpha12<-qf(conf1,nu1,nu2)
  Fconf2<-qf(alpha,nu2,10000000)
  Fconf12<-qf(alpha,nu1,nu2)
  Falpha2<-qf(conf1,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
  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")
}
```

**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)
```

**Format**

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

y a numeric vector

A a factor containing the levels (22, 32) of factor A

B a factor containing the levels (0.5, 5.0) of factor B

C a factor containing the levels (0.5, 5.0) 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

**Examples**

```
data(web)
```

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