

Package ‘DanielBiostatistics10th’

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Type Package

Title Functions for Wayne W. Daniel's Biostatistics, Tenth Edition

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Description Functions to accompany Wayne W. Daniel's
Biostatistics: A Foundation for Analysis in the Health
Sciences, Tenth Edition.

License GPL-2

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DanielBiostatistics10th-package

Functions for Wayne W. Daniel's Biostatistics (Tenth Edition)

Description

Functions and examples to accompany Wayne W. Daniel's *Biostatistics: A Foundation for Analysis in the Health Sciences*, Tenth Edition, Wiley, ISBN: 978-1-119-62550-6.

<https://www.wiley.com/en-us/Biostatistics:+A+Foundation+for+Analysis+in+the+Health+Sciences,+10th+Edition-p-9781119625506>

Data sets from 10th edition <https://bcs.wiley.com/he-bcs/Books?action=resource&bcsId=7849&itemId=1118302796&resourceId=30373>.

Resources from 11th edition <https://bcs.wiley.com/he-bcs/Books?action=index&bcsId=11491&itemId=1119496578>, with errata of data.

binom2pois

Binomial Approaching Poisson

Description

Binomial Approaching Poisson

Usage

```
binom2pois(x, lambda, size = c(10L, 100L))
```

Arguments

x	integer scalar, observed number of responses
lambda	positive numeric scalar, parameter λ of Poisson distribution
size	integer vector, parameter n of binomial distribution

Details

[binom2pois](#) shows how binomial density approaches Poisson density when $n \rightarrow \infty$ and $p \rightarrow 0$, while holding a constant product $np = \lambda$.

Value

[binom2pois](#) returns a [binom2pois](#) object, for which a [print](#) method, an [autolayer](#) and an [autoplot](#) method are defined.

See Also

[dbinom](#) [dpois](#)

Examples

```
binom2pois(x = 4L, lambda = 6, size = seq.int(10L, 50L, by = 10L))
```

BooleanRisk-class *Boolean Risk-&-Disease Table*

Description

Boolean Risk-&-Disease Table

Slots

.Data two-by-two [integer matrix](#), contingency table of a Boolean test-&-disease decisions with layout

	Disease (+)	Disease (-)
Risk Factor (+)	x_{++}	x_{+-}
Risk Factor (-)	x_{-+}	x_{--}

BooleanTest-class *Boolean Test-&-Disease Table*

Description

Boolean Test-&-Disease Table

Slots

.Data two-by-two [integer matrix](#), contingency table of a Boolean test-&-disease decisions with layout

	Disease (+)	Disease (-)
Test (+)	x_{++}	x_{+-}
Test (-)	x_{-+}	x_{--}

Chapter01

Chapter 1

Description

Functions and examples for Chapter 1, *Introduction to Biostatistics*.

Usage

```
sampleRow(x, size, replace = FALSE, prob = NULL)
```

Arguments

x	a data.frame
size	positive integer scalar, number of rows to be selected
replace	logical scalar, whether sampling should be with replacement (default FALSE)
prob	numeric vector of probability weights for each row of input x being sampled. Default NULL indicates simple random sampling

Value

[sampleRow](#) returns a [data.frame](#), a simple random sample from the input.

References

Wayne W. Daniel, *Biostatistics: A Foundation for Analysis in the Health Sciences*, Tenth Edition. Wiley, ISBN: 978-1-119-62550-6.

See Also

[sample.int](#)

Examples

```
library(DanielBiostatistics10th)
# To run a line of code, use shortcut
# Command + Enter: Mac and RStudio Cloud
# Control + Enter: Windows, Mac and RStudio Cloud
# To clear the console
# Control + L: Mac and RStudio Cloud

# Page 8, Example 1.4.1
class(EXA_C01_S04_01) # `EXA_C01_S04_01` is a 'data.frame' (a specific class defined in R)
dim(EXA_C01_S04_01) # dimension, number-row and number-column
head(EXA_C01_S04_01, n = 8L) # first `n` rows of a 'data.frame'
names(EXA_C01_S04_01) # column names of a 'data.frame'
EXA_C01_S04_01$AGE # use `$` to obtain one column from a 'data.frame'
sampleRow(EXA_C01_S04_01, size = 10L, replace = FALSE) # to answer Example 1.4.1

# Page 11, Example 1.4.2
EXA_C01_S04_01[seq.int(from = 4L, to = 166L, by = 18L), ]
```

Description

Functions and examples for Chapter 2, *Descriptive Statistics*.

Usage

```
print_stats(x, na.rm = TRUE)
```

```
print_freqs(x, breaks, include.lowest = TRUE, right = TRUE)
```

Arguments

<code>x</code>	numeric vector, the observations. In print_freqs function, this argument can also be a factor
<code>na.rm</code>	logical scalar, whether to remove the missing observations (default TRUE)
<code>breaks</code>	numeric vector, see cut.default
<code>include.lowest</code>	logical scalar, default TRUE. See cut.default
<code>right</code>	logical scalar, see cut.default

Details

[print_freqs](#) prints the (relative) frequencies and cumulative (relative) frequencies, from a numeric input vector, specified interval breaks as well as open/close status of the ends of the intervals.

[print_stats](#) prints the simple statistics of the input observations, such as sample size, mean, median, (smallest) mode, variance, standard deviation, coefficient of variation (if all observations are non-negative), quartiles, inter-quartile range (IQR), range, skewness and kurtosis. A histogram is also printed.

Value

[print_freqs](#) returns a [freqs](#) object, for which a [show](#) method, an [autolayer](#) and an [autoplot](#) method are defined.

[print_stats](#) does not have a returned value.

References

Wayne W. Daniel, *Biostatistics: A Foundation for Analysis in the Health Sciences*, Tenth Edition. Wiley, ISBN: 978-1-119-62550-6.

See Also

[cut.default](#) [table](#) [cumsum](#) [mean.default](#) [median.default](#) [Mode](#) [var](#) [sd](#) [quantile](#) [skewness](#) [kurtosis](#)

Examples

```

library(DanielBiostatistics10th)

# Page 20, Example 2.2.1
head(EXA_C01_S04_01)
class(EXA_C01_S04_01$AGE) # 'integer'
class(age <- as.numeric(EXA_C01_S04_01$AGE)) # 'numeric'
sort(age) # Page 21, Table 2.2.1 # 'ordered vector'

# Page 23, Example 2.3.1
(ageB = seq.int(from = 30, to = 90, by = 10))
(r231 = print_freqs(age, breaks = ageB, right = FALSE)) # Page 25, Table 2.3.2
# The open/close of interval ends is determined by textbook using 30-39, 40-49, etc.
autoplot(r231, title = 'Page 27, Figure 2.3.2')

# Page 38-42, Example 2.4.1 - Example 2.4.6
# Page 44-46, Example 2.5.1 - Example 2.5.3
print_stats(age) # or some other data input

# Page 49, Example 2.5.4 (omitted)

# Page 50, Example 2.5.5
head(EXA_C02_S05_05)
boxplot(EXA_C02_S05_05$GRF, main = c('GRF from Page 50, Example 2.5.5'))
print_stats(EXA_C02_S05_05$GRF)
print_freqs(EXA_C02_S05_05$GRF, breaks = seq.int(10, 45, by = 5))

```

Chapter03

Chapter 3

Description

Functions for Chapter 3, *Some Basic Probability Concepts*.

Usage

```

addProbs(A)

predictiveValues(
  A,
  sensitivity = A[1, 1]/sum(A[, 1]),
  specificity = A[2, 2]/sum(A[, 2]),
  prevalence = stop("must provide prevalence")
)

```

Arguments

A [integer matrix](#), two-dimensional contingency table. For [predictiveValues](#) function, this must be a [BooleanTest](#) object, or a two-by-two [integer matrix](#) of the same with layout as outlined in [BooleanTest](#)

sensitivity, specificity **numeric** scalars, sensitivity and specificity of a test. By default, these are calculated by the test-disease contingency table A

prevalence **numeric** scalar or vector, prevalence(s) of disease

Details

addProbs provides the joint, marginal and conditional probabilities of a contingency table.

predictiveValues provides the predictive values based on the sensitivity, specificity of a test, as well as the disease prevalence.

Value

addProbs returns an **addProbs** object, which is a **list** consisting of a **noquote matrix** of joint probabilities, and two **noquote matrix** of conditional probabilities. A **print** method is defined for **addProbs** object.

predictiveValues returns a **predictiveValues** object, which is a **list** of three **double** vector elements named 'Prevalence', 'PVP' and 'PVN'. A **print** method and an **autoplot** method are defined for **predictiveValues** object.

References

Wayne W. Daniel, *Biostatistics: A Foundation for Analysis in the Health Sciences*, Tenth Edition. Wiley, ISBN: 978-1-119-62550-6.

See Also

[addmargins](#) [rowSums](#) [colSums](#) [proportions](#)

Examples

```
library(DanielBiostatistics10th)

# Page 69-75, Example 3.4.1 - Example 3.4.8
(d341 = matrix(c(28L, 19L, 41L, 53L, 35L, 38L, 44L, 60L), ncol = 2L, dimnames = list(
  FamilyHx = c('none', 'Bipolar', 'Unipolar', 'UniBipolar'),
  Onset = c('Early', 'Late'))))
class(d341) # 'matrix', i.e., a two-dimensional 'array'
addProbs(d341)

# Page 81, Example 3.5.1
(d351 = matrix(c(436L, 14L, 5L, 495L), nrow = 2L, dimnames = list(
  Test = c('Positive', 'Negative'), Alzheimer = c('Yes', 'No'))))
predictiveValues(d351, prevalence = .113)
predictiveValues(d351, prevalence = c(.005, .98))
```

Description

Functions for Chapter 4, *Probability Distributions*.

Usage

```
binomBar(size, prob, xlim = size, title)
```

```
poisBar(lambda, xlim, title)
```

Arguments

size	non-negative integer scalar, number of trials for binomial distribution
prob	numeric scalar between 0 and 1, probability of success on each trial for binomial distribution
xlim	length-two numeric vector, horizontal limit of the figure
title	character scalar, title of the figure
lambda	positive numeric scalar, mean of Poisson distribution

Details

[binomBar](#) and [poisBar](#) generate bar plots of binomial and Poisson distributions.

Value

[binomBar](#) and [poisBar](#) returns a 'discreteDistBar' object, for which a [print](#) method, an [autolayer](#) and an [autoplot](#) method are defined.

References

Wayne W. Daniel, *Biostatistics: A Foundation for Analysis in the Health Sciences*, Tenth Edition. Wiley, ISBN: 978-1-119-62550-6.

See Also

[dbinom](#) [dpois](#)

Examples

```

library(DanielBiostatistics10th)

# Page 93-97, Example 4.2.1 - Example 4.2.7
d421 = rep(1:8, times = c(62L, 47L, 39L, 39L, 58L, 37L, 4L, 11L))
(fq421 = print_freqs(factor(d421))) # Page 94, Table 4.2.1 and 4.2.2; Page 96, Table 4.2.3
autoplot(fq421, type = 'density', title = 'Page 95, Figure 4.2.1')
autoplot(fq421, type = 'distribution', title = 'Page 96, Figure 4.2.2')

# ?dbinom # 'd' for binomial 'density'; calculate Prob(X = x)
# ?pbinom # 'p' for binomial 'probability'
# `lower.tail = TRUE` (default), calculate Prob(X <= x)
# `lower.tail = FALSE`, calculate Prob(X > x)

# Page 99, Example 4.3.1
dbinom(x = 3L, size = 5L, prob = .858)
# Page 103, Example 4.3.2
dbinom(x = 4L, size = 10L, prob = .14)
# Page 103, Example 4.3.3
(pL = pbinom(q = 5L, size = 25L, prob = .1, lower.tail = TRUE)) # (a) including!
(pU = pbinom(q = 5L, size = 25L, prob = .1, lower.tail = FALSE)) # (b) excluding!
pL + pU # R makes sure they add up to 1
binomBar(size = 25L, prob = .1)
# Page 105, Example 4.3.4
dbinom(x = 7L, size = 12L, prob = .55)
pbinom(q = 5L, size = 12L, prob = .55)
pbinom(q = 7L, size = 12L, prob = .55, lower.tail = FALSE)

# Page 110, Example 4.4.1
dpois(x = 3L, lambda = 12)
# Page 110, Example 4.4.2
ppois(2L, lambda = 12, lower.tail = FALSE)
poisBar(lambda = 12, xlim = 30L)
# Page 110, Example 4.4.3
ppois(1L, lambda = 2)
# Page 111, Example 4.4.4
dpois(3L, lambda = 2)
# Page 112, Example 4.4.5
ppois(5L, lambda = 2, lower.tail = FALSE)

# Page 119. Example 4.6.1
pnorm(2)
# Page 120. Example 4.6.2
pnorm(2.55) - pnorm(-2.55)
1 - 2 * pnorm(-2.55) # alternative solution
# Page 121. Example 4.6.3
pnorm(1.53) - pnorm(-2.74)
# Page 121. Example 4.6.4
pnorm(2.71, lower.tail = FALSE)
# Page 122. Example 4.6.5
pnorm(2.45) - pnorm(.84)

```



```
# Page 122. Example 4.7.1
pnorm(q = 3, mean = 5.4, sd = 1.3)
pnorm(q = (3-5.4)/1.3) # manual solution
# Page 125. Example 4.7.2
pnorm(649, mean = 491, sd = 119) - pnorm(292, mean = 491, sd = 119)
# Page 122. Example 4.7.3
1e4L * pnorm(8.5, mean = 5.4, sd = 1.3, lower.tail = FALSE)
```

Chapter05to07

Chapter 5, 6 and 7

Description

Functions for Chapter 5, *Some Important Sampling Distributions*, Chapter 6, *Estimation* and Chapter 7, *Hypothesis Testing*.

Usage

```
aggregated_z(  
  xbar,  
  n,  
  sd,  
  null.value,  
  alternative = c("two.sided", "less", "greater"),  
  conf.level = 0.95,  
  ...  
)
```

```
aggregated_t(  
  xbar,  
  xsd,  
  n,  
  null.value,  
  var.equal = FALSE,  
  alternative = c("two.sided", "less", "greater"),  
  conf.level = 0.95,  
  ...  
)
```

```
prop_CLT(  
  x,  
  n,  
  bool_obs,  
  xbar = x/n,  
  null.value,  
  alternative = c("two.sided", "less", "greater"),  
  conf.level = 0.95,
```

```

    ...
  )

  aggregated_var(
    xsd,
    n,
    null.value,
    alternative = c("two.sided", "less", "greater"),
    conf.level = 0.95,
    ...
  )

```

Arguments

xbar	numeric scalar or length-two vector. Sample mean(s) for numeric variable(s) \bar{x} or (\bar{x}_1, \bar{x}_2) . Sample proportion(s) for binary (i.e., logical) variable(s) \hat{p} or (\hat{p}_1, \hat{p}_2) . In the case of two-sample tests, this could also be a numeric scalar indicating the difference in sample means $\bar{x}_1 - \bar{x}_2$ or sample proportions $\hat{p}_1 - \hat{p}_2$
n	integer scalar n or length-two vector. Sample size(s) n or (n_1, n_2)
sd	numeric scalar or length-two vector. population standard deviation(s) σ or (σ_1, σ_2)
null.value	(optional) numeric scalar or length-two vector. Null value(s) of the population mean(s) $(\mu_0, (\mu_{10}, \mu_{20}),$ or $\mu_{10} - \mu_{20})$ for aggregated_z and aggregated_t . Null value(s) of the population proportion(s) $(p_0, (p_{10}, p_{20}),$ or $p_{10} - p_{20})$ for prop_CLT . Null value(s) of the population variance(s) (ratio) $(\sigma_0^2, (\sigma_{10}^2, \sigma_{20}^2),$ or $\sigma_{10}^2/\sigma_{20}^2)$ for aggregated_var . If missing, only the confidence intervals will be computed.
alternative	character scalar, alternative hypothesis, either 'two.sided' (default), 'greater' or 'less'
conf.level	numeric scalar, confidence level, default 0.95
...	potential arguments, not in use currently
xsd	numeric scalar or length-two vector. Sample standard deviation(s) $\sigma_{\bar{x}}$ or $(\sigma_{\bar{x}_1}, \sigma_{\bar{x}_2})$
var.equal	logical scalar, whether to treat the two population variances as being equal (default FALSE) in aggregated_t
x	integer scalar or length-two vector, number of positive count(s) of binary (i.e., logical) variable(s)
bool_obs	logical vector of Boolean observations, used in one-sample z -test on proportion

Details

[aggregated_z](#) performs one- or two-sample z -test using the aggregated statistics of sample mean(s) and sample size(s) when `null.value` is provided. Otherwise, only the confidence interval based on z -distribution is computed.

[aggregated_t](#) performs one- or two-sample t -test using the aggregated statistics of sample mean(s), sample standard deviation(s) and sample size(s) when `null.value` is provided. Otherwise, only the confidence interval based on t -distribution is computed.

`prop_CLT` performs one- or two-sample z -test on proportion(s), using Central Limit Theorem when `null.value` is provided. Otherwise, only the confidence interval based on z -distribution is computed.

`aggregated_var` performs one-sample χ^2 -test on variance, or two-sample F -test on variances, using the aggregated statistics of sample standard deviation(s) and sample size(s) when `null.value` is provided. Otherwise, only the confidence interval based on χ^2 - or F -distribution is computed.

Value

`aggregated_z` returns an 'htest' object when `null.value` is provided, otherwise returns a length-two numeric vector.

`aggregated_t` returns an `htest` object when `null.value` is provided, otherwise returns a length-two numeric vector.

`prop_CLT` returns an `htest` object when `null.value` is provided, otherwise returns a length-two numeric vector.

`aggregated_var` returns an `htest` object when `null.value` is provided, otherwise returns a length-two numeric vector.

References

Wayne W. Daniel, *Biostatistics: A Foundation for Analysis in the Health Sciences*, Tenth Edition. Wiley, ISBN: 978-1-119-62550-6.

See Also

[t.test](#) [prop.test](#) [var.test](#)

Examples

```
library(DanielBiostatistics10th)

# Page 142, Example 5.3.2
aggregated_z(xbar = 190, sd = 12.7, n = 10L, null.value = 185.6, alternative = 'greater')
# Page 143, Example 5.3.3
pnorm(125, mean = 120, sd = 15/sqrt(50)) - pnorm(115, mean = 120, sd = 15/sqrt(50))
aggregated_z(125, sd = 15, n = 50L, null.value = 120, alternative = 'less')$p.value -
  aggregated_z(115, sd = 15, n = 50L, null.value = 120, alternative = 'less')$p.value

# Page 145, Example 5.4.1
aggregated_z(xbar = c(92, 105), sd = 20, n = 15L, null.value = 0, alternative = 'less')
# Page 148, Example 5.4.2
aggregated_z(xbar = 20, sd = c(15, 20), n = c(35L, 40L), null.value = c(45, 30),
  alternative = 'greater')

# Page 150, Example 5.5.1
prop_CLT(xbar = .4, n = 150L, null.value = .357, alternative = 'greater')
# Page 152, Example 5.5.2
prop_CLT(xbar = .45, n = 200L, null.value = .51, alternative = 'less')

# Page 155, Example 5.6.1
```

```

prop_CLT(xbar = .1, null.value = c(.28, .21), n = c(100L, 100L), alternative = 'greater')
# Page 155, Example 5.6.2
prop_CLT(xbar = .05, null.value = c(.34, .26), n = c(250L, 200L), alternative = 'less')

# Page 166, Example 6.2.1
aggregated_z(xbar = 22, n = 10L, sd = sqrt(45))
# Page 168, Example 6.2.2
aggregated_z(xbar = 84.3, n = 15L, sd = sqrt(144), conf.level = .99)
# Page 168, Example 6.2.3
aggregated_z(xbar = 17.2, n = 35L, sd = 8, conf.level = .9)
# Page 169, Example 6.2.4
head(EXA_C06_S02_04)
aggregated_z(xbar = mean(EXA_C06_S02_04$ACTIVITY), n = nrow(EXA_C06_S02_04), sd = sqrt(.36))

# Page 173, Example 6.3.1
aggregated_t(xbar = 250.8, xsd = 130.9, n = 19L)

# Page 177, Example 6.4.1
aggregated_z(xbar = c(4.5, 3.4), sd = sqrt(c(1, 1.5)), n = c(12L, 15L))
# Page 178, Example 6.4.2
aggregated_z(xbar = c(4.3, 13), sd = c(5.22, 8.97), n = c(328L, 64L), conf.level = .99)
# Page 180, Example 6.4.3
aggregated_t(xbar = c(4.7, 8.8), xsd = c(9.3, 11.5), n = c(18L, 10L), var.equal = TRUE)
# Page 181, Example 6.4.4
aggregated_t(xbar = c(4.7, 8.8), xsd = c(9.3, 11.5), n = c(18L, 10L))
# Welch slightly different from Cochran; textbook explained on Page 182

# Page 185, Example 6.5.1
prop_CLT(xbar = .18, n = 1220L)

# Page 187, Example 6.6.1
prop_CLT(x = c(31L, 53L), n = c(68L, 255L), conf.level = .99)

# Page 190, Example 6.7.1
n_671 = uniroot(f = function(n, sd, level = .95) {
  qnorm(1-(1-level)/2) * sd/sqrt(n) - 5 # half-width of CI <= 5 grams
}, interval = c(0, 2e2), sd = 20)
sprintf('Example 6.7.1 requires a sample size of %d.', ceiling(n_671$root))

# Page 192, Example 6.8.1
n_681 = uniroot(f = function(n, p, level = .95) {
  qnorm(1-(1-level)/2) * sqrt(p*(1-p)/n) - .05
}, interval = c(0, 1e3), p = .35)
sprintf('Example 6.8.1 requires a sample size of %d.', ceiling(n_681$root))

# Page 196, Example 6.9.1
d691 = c(9.7, 12.3, 11.2, 5.1, 24.8, 14.8, 17.7)
sqrt(aggregated_var(xsd = sd(d691), n = length(d691)))

# Page 200, Example 6.10.1
aggregated_var(xsd = c(8.1, 5.9), n = c(16L, 4L))

# Page 222, Example 7.2.1

```

```

aggregated_z(xbar = 27, sd = sqrt(20), n = 10L, null.value = 30)
# Page 226, Example 7.2.2
aggregated_z(xbar = 27, sd = sqrt(20), n = 10L, null.value = 30, alternative = 'less')
# Page 228, Example 7.2.3
head(EXA_C07_S02_03)
t.test(EXA_C07_S02_03$DAYS, mu = 15)
# Page 231, Example 7.2.4
aggregated_z(xbar = 146, sd = 27, n = 157L, null.value = 140, alternative = 'greater')
# Page 232, Example 7.2.5
d725 = c(33.38, 32.15, 34.34, 33.95, 33.46, 34.13, 33.99, 34.10, 33.85,
        34.23, 34.45, 34.19, 33.97, 32.73, 34.05)
t.test(d725, mu = 34.5)

# Page 237, Example 7.3.1
aggregated_z(xbar = c(4.5, 3.4), sd = sqrt(c(1, 1.5)), n = c(12L, 15L), null.value = 0)
# Page 239, Example 7.3.2
head(EXA_C07_S03_02)
with(EXA_C07_S03_02, t.test(x = CONTROL, y = SCI, alternative = 'less', var.equal = TRUE))
# Page 240, Example 7.3.3
aggregated_t(xbar = c(19.16, 9.53), xsd = c(5.29, 2.69), n = c(15L, 30L), null.value = 0)
# Page 242, Example 7.3.4
aggregated_z(xbar = c(59.01, 46.61), sd = c(44.89, 34.85), n = c(53L, 54L), null.value = 0,
            alternative = 'greater')

# Page 251, Example 7.4.1
head(EXA_C07_S04_01)
with(EXA_C07_S04_01, t.test(x = POSTOP, y = PREOP, alternative = 'greater', paired = TRUE))

# Page 258, Example 7.5.1
prop_CLT(x = 24L, n = 301L, null.value = .063, alternative = 'greater')

# Page 261, Example 7.6.1
prop_CLT(x = c(24L, 11L), n = c(44L, 29L), null.value = 0, alternative = 'greater')

# Page 264, Example 7.7.1
head(EXA_C07_S07_01)
aggregated_var(xsd = sd(EXA_C07_S07_01$mass), n = 16L, null.value = 600)

# Page 268, Example 7.8.1
aggregated_var(xsd = c(30.62, 11.37), n = 6L, null.value = 1, alternative = 'greater')
# Page 270, Example 7.8.2
with(EXA_C07_S03_02, var.test(x = CONTROL, y = SCI))

```

Description

Functions for Chapter 7, *Hypothesis Testing*.

Usage

```
power_z(
  x,
  null.value,
  sd,
  n,
  std.err = sd/sqrt(n),
  alternative = c("two.sided", "less", "greater"),
  sig.level = 0.05
)
```

Arguments

<code>x</code>	numeric vector, mean parameter(s) μ_1 in the alternative hypothesis
<code>null.value</code>	numeric scalar, mean parameter μ_0 in the null hypothesis
<code>sd</code>	numeric scalar, population standard deviation σ
<code>n</code>	integer scalar, sample size n
<code>std.err</code>	numeric scalar, standardized error. For one-sample z -test, this is σ/\sqrt{n} . Be aware of the name clash with <code>stderr</code>
<code>alternative</code>	character scalar, alternative hypothesis, either 'two.sided' (default), 'greater' or 'less'
<code>sig.level</code>	numeric scalar, significance level (i.e., Type-I-error rate), default .05

Details

`power_z` calculates the powers at each element of the alternative parameters μ_1 , for one-sample z -test

- $H_0 : \mu = \mu_0$ vs. $H_A : \mu \neq \mu_0$, if `alternative = 'two.sided'`
- $H_0 : \mu \leq \mu_0$ vs. $H_A : \mu > \mu_0$, if `alternative = 'greater'`
- $H_0 : \mu \geq \mu_0$ vs. $H_A : \mu < \mu_0$, if `alternative = 'less'`

Value

`power_z` returns a **power** object, for which a `show` method, an `autolayer` and an `autoplot` method are defined for `power_z` object.

References

Wayne W. Daniel, *Biostatistics: A Foundation for Analysis in the Health Sciences*, Tenth Edition. Wiley, ISBN: 978-1-119-62550-6.

See Also

[power.t.test](#)

Examples

```

library(DanielBiostatistics10th)

# Page 272, Example 7.9.1
(p791 = power_z(seq.int(from = 16, to = 19, by = .5), null.value = 17.5, sd = 3.6, n = 100L))
# Page 275, Table 7.9.1
autoplot(p791, title = 'Page 275, Figure 7.9.2')

# Page 276, Example 7.9.2
(p792 = power_z(seq.int(from = 50, to = 70, by = 5), null.value = 65, sd = 15, n = 20L,
  sig.level = .01, alternative = 'less'))
autoplot(p792, title = 'Page 277, Figure 7.9.4')
autoplot(p792, all.alternative = TRUE, title = '1-sided vs. 2-sided test')

# Page 278, Example 7.10.1
(n_d7101 <- uniroot(f = function(x) {
  power_z(55, null.value = 65, sd = 15, n = x, sig.level = .01, alternative = 'less') - .95
}, interval = c(0, 50))$root)
(C_d7101 = qnorm(p = .01, mean = 65, sd = 15/sqrt(ceiling(n_d7101)), lower.tail = TRUE))

```

Chapter09

Chapter 9

Description

Functions for Chapter 9, *Simple Linear Regression and Correlation*.

Usage

```
predict_lm(object, newx, level = 0.95, ...)
```

Arguments

object	lm object, with one and only one numeric predictor
newx	(optional) numeric scalar or vector, new x -value(s) for which the fitted response(s) are to be reported
level	numeric scalar, tolerance/confidence level, default .95
...	potential arguments, not in use currently

Value

`predict_lm` returns a `predict_lm` object, for which a `print` method, an `autolayer` and an `autoplot` method are defined.

References

Wayne W. Daniel, *Biostatistics: A Foundation for Analysis in the Health Sciences*, Tenth Edition. Wiley, ISBN: 978-1-119-62550-6.

See Also[predict.lm](#)**Examples**

```

library(DanielBiostatistics10th)

# Page 417, Example 9.3.1
head(EXA_C09_S03_01)
names(EXA_C09_S03_01)[2:3] = c('Waist', 'AT')
plot(AT ~ Waist, data = EXA_C09_S03_01, xlab = 'Waist circumference (cm), X',
      ylab = 'Deep abdominal AT area (cm2), Y', main = 'Page 419, Figure 9.3.1')

# Page 436, Example 9.4.2
summary(m931 <- lm(AT ~ Waist, data = EXA_C09_S03_01))
cor(EXA_C09_S03_01[2:3]); cor.test(~ AT + Waist, data = EXA_C09_S03_01)
confint(m931) # confidence interval of regression coefficients
anova(m931)

# Page 440, Example 9.4.3
plot(m931, which = 1, main = 'Page 440, Figure 9.4.8')

# Page 441, Section 9.5
autoplot(predict_lm(m931), xlab = 'Waist circumference (cm), X',
           ylab = 'Deep abdominal AT area (cm2), Y',
           title = 'Page 422, Figure 9.3.3; Page 442, Figure 9.5.1')

# Page 447, Example 9.7.1
head(EXA_C09_S07_01)
summary(mod_971 <- lm(CV ~ HEIGHT, data = EXA_C09_S07_01))
autoplot(predict_lm(mod_971), xlab = 'Height (cm)', ylab = 'Cv (units)',
          title = 'Page 449, Figure 9.7.2')

# Page 452, Example 9.7.2
cor(EXA_C09_S07_01); cor.test(~ CV + HEIGHT, data = EXA_C09_S07_01)
# Page 451, Figure 9.7.4, Figure 9.7.5

# Page 453, When the Hypothesized rho Is a Nonzero Value
# R does not have a function to do this

```

Description

Functions for Chapter 11, *Regression Analysis: Some Additional Techniques*.

Usage

```
predict_glm_binomial(object, newx, level = 0.95, ...)
```

Arguments

object	glm object with binomial link function, i.e., a logistic regression model, as well as one and only one numeric predictor
newx	(optional) numeric scalar or vector, new x -value(s) for which the fitted response(s) are to be reported
level	numeric scalar, tolerance/confidence level, default .95
...	potential arguments, not in use currently

Value

`predict_glm_binomial` returns a `predict_glm_binomial` object, for which a `print` method, an `auto-layer` and an `autoplot` method are defined.

References

Wayne W. Daniel, *Biostatistics: A Foundation for Analysis in the Health Sciences*, Tenth Edition. Wiley, ISBN: 978-1-119-62550-6.

See Also

[predict.glm](#)

Examples

```
library(DanielBiostatistics10th)
library(car)
library(DescTools)

# Page 540, Example 11.1.1
head(EXA_C11_S01_01)
head(log(EXA_C11_S01_01$conc, base = 10))
head(EXA_C11_S01_01$logConc)

# Page 542, Example 11.1.2
head(EXA_C11_S01_02)
cor.test(~ sbp + weight, data = EXA_C11_S01_02)
cor.test(~ sbp + bmi, data = EXA_C11_S01_02)

# Page 545, Example 11.2.1
head(EXA_C11_S02_01)
d1121 = within(EXA_C11_S02_01, expr = {
  SMOKE = as.logical(SMOKE)
})
xlab1121 = 'Length of gestation (weeks)'; ylab1121 = 'Birth weight (grams)'
car::scatterplot(GRAMS ~ WEEKS | SMOKE, data = d1121, regLine = FALSE, smooth = FALSE,
  xlab = xlab1121, ylab = ylab1121, main = 'Page 547, Figure 11.2.1')
```

```

# Page 547, Figure 11.2.2: main model (without interaction)
summary(m1121_main <- lm(GRAMS ~ WEEKS + SMOKE, data = d1121))
confint(m1121_main)
car::scatterplot(GRAMS ~ WEEKS | SMOKE, data = d1121, regLine = FALSE, smooth = FALSE,
                 xlab = xlab1121, ylab = ylab1121, main = 'Page 548, Figure 11.2.3')
(cf_main = m1121_main$coefficients)
abline(a = cf_main[1L], b = cf_main[2L], col = 'blue') # regression line for non-smoking mothers
abline(a = cf_main[1L] + cf_main[3L], b = cf_main[2L], col = 'magenta')

# Page 551, Example 11.2.3
d1123 = within(EXA_C11_S02_03, expr = {
  METHOD = factor(METHOD, levels = c('C', 'A', 'B')) # textbook designated 'C' as reference level
})
summary(mod_1123 <- lm(EFFECT ~ AGE * METHOD, data = d1123)) # Page 555, Figure 11.2.5
confint(mod_1123)
car::scatterplot(EFFECT ~ AGE | METHOD, data = d1123, smooth = FALSE,
                 xlab = 'Age', ylab = 'Treatment effectiveness', main = 'Page 555, Figure 11.2.6')

# (optional) Page 561, Example 11.3.1
head(EXA_C11_S03_01)
names(EXA_C11_S03_01) = c('JOBPER', 'ASRV', 'ENTH', 'AMB', 'COMM', 'PROB', 'INIT')
summary(mod_1131_raw <- lm(JOBPER ~ ASRV + ENTH + AMB + COMM + PROB + INIT, data = EXA_C11_S03_01))
# summary(mod_1131 <- MASS::stepAIC(mod_1131_raw, direction = 'backward'))
# the stepwise selection criterion used in MINITAB is not necessarily AIC

# Page 572, Example 11.4.1
addmargins(d1141 <- array(c(92L, 21L, 15L, 20L), dim = c(2L, 2L), dimnames = list(
  OCAD = c('Present', 'Absent'), Sex = c('Male', 'Female')))) # Page 572, Table 11.4.2
(d1141a = within(as.data.frame(as.table(d1141)), expr = {
  OCAD = (OCAD == 'Present')
  Sex = factor(Sex, levels = c('Female', 'Male'))
}))
(m1141 = glm(OCAD ~ Sex, family = binomial(link = 'logit'), weights = Freq, data = d1141a))
summary(m1141) # Page 573, Figure 11.4.1
exp(m1141$coefficients[2L]) # exp(beta_M)
exp(confint(m1141)) # confidence interval of exp(beta)
predict(m1141, newdata = data.frame(Sex = setNames(nm = c('Male', 'Female'))), type = 'response')

# Page 573, Example 11.4.2
head(EXA_C11_S04_02)
summary(mod_1142 <- glm(ATT ~ AGE, family = binomial, data = EXA_C11_S04_02))
# .. Page 575, Figure 11.4.2
exp(mod_1142$coefficients[2L])
exp(confint(mod_1142))
car::Anova(mod_1142) # Optional
autoplot(predict_glm_binomial(mod_1142, newx = c(50, 65, 80)), title = 'Page 576, Figure 11.4.3')

# (optional) Page 576, Example 11.4.3
head(REV_C11_24)
summary(glm(ONSET ~ HIAA + TRYPT, family = binomial(link = 'logit'), data = REV_C11_24))
# Page 577, Figure 11.4.4
# Predictor TRYPT should be removed from model due to p-value \approx 1
summary(glm(ONSET ~ HIAA, family = binomial(link = 'logit'), data = REV_C11_24))

```

```
# (optional) Page 578, Example 11.4.4
DescTools::PseudoR2(mod_1142, which = 'CoxSnell')
DescTools::PseudoR2(mod_1142, which = 'Nagelkerke')

# (optional) Page 579, Example 11.4.5 (same as Example 11.4.4)
```

Chapter12

Chapter 12

Description

Functions for Chapter 12, *The Chi-Square Distribution and The Analysis of Frequencies*.

Usage

```
relativeRisk(A)

oddsRatio(A)

print_OE(0, prob)
```

Arguments

A	a BooleanRisk object, or a two-by-two integer matrix of the same with layout as outlined in BooleanRisk
0	integer vector, observed counts
prob	numeric vector, anticipated probability. If missing (default), an uniform distribution across all categories are used.

Value

[relativeRisk](#) returns a 'logRelativeRisk' object, for which a [vcov](#) method and a [print](#) method are defined.

[oddsRatio](#) returns a 'logOddsRatio' object, for which a [vcov](#) method and a [print](#) method are defined.

[print_OE](#) prints a table with observed and expected frequencies, as well as the category-wise χ^2 statistics. A [double](#) vector of the category-wise χ^2 statistics is returned invisibly.

References

Wayne W. Daniel, *Biostatistics: A Foundation for Analysis in the Health Sciences*, Tenth Edition. Wiley, ISBN: 978-1-119-62550-6.

Examples

```

library(DanielBiostatistics10th)

# Page 605, Example 12.3.1
d1231_b = c(-Inf, seq.int(from = 125, to = 275, by = 25), Inf)
(d1231 = setNames( # Page 605, Table 12.3.1
  c(1L, 3L, 8L, 18L, 6L, 4L, 4L, 3L),
  nm = levels(cut(double(), breaks = d1231_b, right = FALSE, include.lowest = TRUE))))
chi1231 = print_OE(d1231, prob = diff.default(pnorm(q = d1231_b, mean = 198.67, sd = 41.31)))
pchisq(sum(chi1231), df = length(d1231) - 3L, lower.tail = FALSE)
# -3L: three restrictions (explained on Page 608)
# (1) making sum(xo) == sum(xe)
# (2) estimating mean
# (3) estimating sd

# Page 609, Example 12.3.2
# 100 doctors, 25 patients per doctor
d1232 = c(5L, 6L, 8L, 10L, 10L, 15L, 17L, 10L, 10L, 9L, 0L)
o1232 = setNames(c(sum(d1232[1:2]), d1232[-(1:2)]), nm = c('0-1', 2:9, '10 or more'))
(p1232 = sum((0:10) * d1232) / (25 * 100)) # binomial `prob`
chi1232 = print_OE(o1232, prob = c(
  pbinom(1L, size = 25L, prob = p1232),
  dbinom(2:9, size = 25L, prob = p1232),
  pbinom(9, size = 25L, prob = p1232, lower.tail = FALSE)))
pchisq(sum(chi1232), df = length(o1232) - 2L, lower.tail = FALSE)
# -2L: two restrictions (explained on Page 611)
# (1) making sum(o) == sum(e)
# (2) estimating p1232

# Page 611, Example 12.3.3
d1233 = c(5L, 14L, 15L, 23L, 16L, 9L, 3L, 3L, 1L, 1L, 0L)
o_1233 = setNames(c(d1233[1:8], sum(d1233[-(1:8)])), nm = c(0:7, '8 or more'))
p_1233 = c(dpois(0:7, lambda = 3), # lambda = 3 is provided by the textbook
  ppois(7L, lambda = 3, lower.tail = FALSE))
chi1233 = print_OE(o_1233, prob = p_1233)
pchisq(sum(chi1233), df = length(o_1233) - 1L, lower.tail = FALSE)
# -1L: one restrictions
# (1) making sum(xo) == sum(xe)
chisq.test(o_1233, p = p_1233) # equivalent # warning on any(E < 5)

# Page 614, Example 12.3.4
d1234 = c('Dec 05' = 62L, 'Jan 06' = 84L, 'Feb 06' = 17L, 'Mar 06' = 16L, 'Apr 06' = 21L)
chi1234 = print_OE(d1234)
pchisq(sum(chi1234), df = length(d1234) - 1L, lower.tail = FALSE)
chisq.test(d1234) # equivalent

# Page 616, Example 12.3.5
d1235 = c(dominant = 43L, heterozygous = 125L, recessive = 32L)
chi1235 = print_OE(d1235, prob = c(1, 2, 1))
pchisq(sum(chi1235), df = length(d1235) - 1L, lower.tail = FALSE)
chisq.test(d1235, p = c(1, 2, 1), rescale.p = TRUE) # equivalent

```

```

# Page 621, Example 12.4.1
addmargins(d1241 <- array(c(260L, 15L, 7L, 299L, 41L, 14L), dim = c(3L, 2L), dimnames = list(
  Race = c('White', 'Black', 'Other'),
  FolicAcid = c('TRUE', 'FALSE'))))
chisq.test(d1241) # ?stats::chisq.test

# Page 626, Example 12.4.2
addmargins(d1242 <- array(c(131L, 14L, 52L, 36L), dim = c(2L, 2L), dimnames = list(
  Type = c('Faller', 'Non-Faller'),
  LifestyleChange = c('TRUE', 'FALSE'))))
chisq.test(d1242, correct = FALSE)
chisq.test(d1242, correct = TRUE) # Page 627, Yates's Correction

# Page 631, Example 12.5.1
addmargins(d1251 <- array(c(21L, 19L, 75L, 77L), dim = c(2L, 2L), dimnames = list(
  Group = c('Narcoleptic', 'Healthy'),
  Migraine = c('TRUE', 'FALSE'))))
(chisq_1251 = chisq.test(d1251, correct = FALSE))
if (FALSE) {
  # (optional) using test on two proportions
  # only equivalent for 2*2 contingency table
  (clt_1251 = prop_CLT(x = c(21L, 19L), n = 96L, null.value = 0))
  all.equal.numeric(unnamed(clt_1251$statistic^2), unnamed(chisq_1251$statistic))
}

# Page 638, Example 12.6.1
addmargins(d1262 <- array(c(2L, 8L, 7L, 4L), dim = c(2L, 2L), dimnames = list(
  Group = c('PI_Naive', 'PA_Experienced'),
  Regimen2yr = c('TRUE', 'FALSE'))))
fisher.test(d1262)

# Page 644, Example 12.7.1
addmargins(d1271 <- array(c(22L, 18L, 216L, 199L), dim = c(2L, 2L),
  dimnames = list(Exercising = c('Extreme', 'No'), PretermLabor = c('TRUE', 'FALSE'))))
relativeRisk(d1271)
# textbook confidence interval (.65, 1.86) wrong (too many rounding in intermediate steps)

# Page 647, Example 12.7.2
addmargins(d1272 <- array(c(64L, 68L, 342L, 3496L), dim = c(2L, 2L), dimnames = list(
  SmkPregnancy = c('TRUE', 'FALSE'),
  Obesity = c('TRUE', 'FALSE'))))
oddsRatio(d1272)

# Page 650, Example 12.7.3
# Page 652, Example 12.7.4
(d1273 <- array(c(21L, 16L, 11L, 6L, 50L, 18L, 14L, 6L), dim = c(2L, 2L, 2L), dimnames = list(
  HTN = c('Present', 'Absent'), OCAD = c('Cases', 'Controls'),
  Age = c('<=55', '>55'))))
addmargins(d1273, margin = 1:2) # Page 651, Table 12.7.6
mantelhaen.test(d1273)

```

EXA_C01_S04_01 *Data of Example 1.4.1*

Description

EXA_C01_S04_01

Usage

EXA_C01_S04_01

Format

A [data.frame](#) with 189 rows and 2 columns

EXA_C02_S05_05 *Data of Example 2.5.5*

Description

EXA_C02_S05_05

Usage

EXA_C02_S05_05

Format

A [data.frame](#) with 20 rows and 1 columns

EXA_C06_S02_04 *Data of Example 6.2.4*

Description

EXA_C06_S02_04

Usage

EXA_C06_S02_04

Format

A [data.frame](#) with 35 rows and 1 columns

EXA_C07_S02_03 *Data of Example 7.2.3*

Description

EXA_C07_S02_03

Usage

EXA_C07_S02_03

Format

A [data.frame](#) with 17 rows and 2 columns

EXA_C07_S03_02 *Data of Example 7.3.2*

Description

EXA_C07_S03_02

Usage

EXA_C07_S03_02

Format

A [data.frame](#) with 10 rows and 2 columns

EXA_C07_S04_01 *Data of Example 7.4.1*

Description

EXA_C07_S04_01

Usage

EXA_C07_S04_01

Format

A [data.frame](#) with 12 rows and 2 columns

EXA_C07_S07_01 *Data of Example 7.7.1*

Description

EXA_C07_S07_01

Usage

EXA_C07_S07_01

Format

A [data.frame](#) with 17 rows and 1 columns

EXA_C08_S02_01 *Data of Example 8.2.1*

Description

EXA_C08_S02_01

Usage

EXA_C08_S02_01

Format

A [data.frame](#) with 144 rows and 2 columns

EXA_C08_S03_01 *Data of Example 8.3.1*

Description

EXA_C08_S03_01

Usage

EXA_C08_S03_01

Format

A [data.frame](#) with 15 rows and 3 columns

EXA_C08_S04_01 *Data of Example 8.4.1*

Description

EXA_C08_S04_01

Usage

EXA_C08_S04_01

Format

A [data.frame](#) with 72 rows and 3 columns

EXA_C08_S04_02 *Data of Example 8.4.2*

Description

EXA_C08_S04_02

Usage

EXA_C08_S04_02

Format

A [data.frame](#) with 25 rows and 6 columns

EXA_C08_S05_02 *Data of Example 8.5.2*

Description

EXA_C08_S05_02

Usage

EXA_C08_S05_02

Format

A [data.frame](#) with 80 rows and 3 columns

EXA_C09_S03_01 *Data of Example 9.3.1*

Description

EXA_C09_S03_01

Usage

EXA_C09_S03_01

Format

A [data.frame](#) with 109 rows and 3 columns

EXA_C09_S07_01 *Data of Example 9.7.1*

Description

EXA_C09_S07_01

Usage

EXA_C09_S07_01

Format

A [data.frame](#) with 155 rows and 2 columns

EXA_C10_S03_01 *Data of Example 10.3.1*

Description

EXA_C10_S03_01

Usage

EXA_C10_S03_01

Format

A [data.frame](#) with 71 rows and 3 columns

EXA_C10_S06_01	<i>Data of Example 10.6.1</i>
----------------	-------------------------------

Description

EXA_C10_S06_01

Usage

EXA_C10_S06_01

FormatA [data.frame](#) with 29 rows and 3 columns

EXA_C11_S01_01	<i>Data of Example 11.1.1</i>
----------------	-------------------------------

Description

EXA_C11_S01_01

Usage

EXA_C11_S01_01

FormatA [data.frame](#) with 25 rows and 3 columns

EXA_C11_S01_02	<i>Data of Example 11.1.2</i>
----------------	-------------------------------

Description

EXA_C11_S01_02

Usage

EXA_C11_S01_02

FormatA [data.frame](#) with 15 rows and 4 columns

EXA_C11_S02_01 *Data of Example 11.2.1*

Description

EXA_C11_S02_01

Usage

EXA_C11_S02_01

Format

A [data.frame](#) with 100 rows and 4 columns

EXA_C11_S02_03 *Data of Example 11.2.3*

Description

EXA_C11_S02_03

Usage

EXA_C11_S02_03

Format

A [data.frame](#) with 36 rows and 3 columns

EXA_C11_S03_01 *Data of Example 11.3.1*

Description

EXA_C11_S03_01

Usage

EXA_C11_S03_01

Format

A [data.frame](#) with 30 rows and 7 columns

EXA_C11_S04_02	<i>Data of Example 11.4.2</i>
----------------	-------------------------------

Description

EXA_C11_S04_02

Usage

EXA_C11_S04_02

FormatA [data.frame](#) with 184 rows and 2 columns

EXA_C11_S05_01	<i>Data of Example 11.5.1</i>
----------------	-------------------------------

Description

EXA_C11_S05_01

Usage

EXA_C11_S05_01

FormatA [data.frame](#) with 45 rows and 4 columns

EXA_C12_S02_03	<i>Data of Example 12.2.3</i>
----------------	-------------------------------

Description

EXA_C12_S02_03

Usage

EXA_C12_S02_03

FormatA [data.frame](#) with 90 rows and 2 columns

EXA_C13_S03_02 *Data of Example 13.3.2*

Description

EXA_C13_S03_02

Usage

EXA_C13_S03_02

Format

A [data.frame](#) with 12 rows and 3 columns

EXA_C13_S05_01 *Data of Example 13.5.1*

Description

EXA_C13_S05_01

Usage

EXA_C13_S05_01

Format

A [data.frame](#) with 16 rows and 2 columns

EXA_C13_S06_01 *Data of Example 13.6.1*

Description

EXA_C13_S06_01

Usage

EXA_C13_S06_01

Format

A [data.frame](#) with 15 rows and 2 columns

EXA_C13_S07_01 *Data of Example 13.7.1*

Description

EXA_C13_S07_01

Usage

EXA_C13_S07_01

Format

A [data.frame](#) with 36 rows and 1 columns

EXA_C13_S08_02 *Data of Example 13.8.2*

Description

EXA_C13_S08_02

Usage

EXA_C13_S08_02

Format

A [data.frame](#) with 10 rows and 5 columns

EXA_C13_S09_01 *Data of Example 13.9.1*

Description

EXA_C13_S09_01

Usage

EXA_C13_S09_01

Format

A [data.frame](#) with 9 rows and 4 columns

EXA_C13_S09_02 *Data of Example 13.9.2*

Description

EXA_C13_S09_02

Usage

EXA_C13_S09_02

Format

A [data.frame](#) with 16 rows and 5 columns

EXA_C13_S10_01 *Data of Example 13.10.1*

Description

EXA_C13_S10_01

Usage

EXA_C13_S10_01

Format

A [data.frame](#) with 20 rows and 3 columns

EXA_C13_S10_02 *Data of Example 13.10.2*

Description

EXA_C13_S10_02

Usage

EXA_C13_S10_02

Format

A [data.frame](#) with 35 rows and 3 columns

EXA_C14_S03_01 *Data of Example 14.3.1*

Description

EXA_C14_S03_01

Usage

EXA_C14_S03_01

Format

A [data.frame](#) with 39 rows and 4 columns

EXA_C14_S05_01 *Data of Example 14.5.1*

Description

EXA_C14_S05_01

Usage

EXA_C14_S05_01

Format

A [data.frame](#) with 40 rows and 5 columns

EXR_C02_S03_01 *Data of Exercise 2.3.1*

Description

EXR_C02_S03_01

Usage

EXR_C02_S03_01

Format

A [data.frame](#) with 90 rows and 1 columns

EXR_C02_S03_02 *Data of Exercise 2.3.2*

Description

EXR_C02_S03_02

Usage

EXR_C02_S03_02

Format

A [data.frame](#) with 159 rows and 1 columns

EXR_C02_S03_03 *Data of Exercise 2.3.3*

Description

EXR_C02_S03_03

Usage

EXR_C02_S03_03

Format

A [data.frame](#) with 29 rows and 1 columns

EXR_C02_S03_04 *Data of Exercise 2.3.4*

Description

EXR_C02_S03_04

Usage

EXR_C02_S03_04

Format

A [data.frame](#) with 53 rows and 1 columns

EXR_C02_S03_05 *Data of Exercise 2.3.5*

Description

EXR_C02_S03_05

Usage

EXR_C02_S03_05

Format

A [data.frame](#) with 45 rows and 1 columns

EXR_C02_S03_06 *Data of Exercise 2.3.6*

Description

EXR_C02_S03_06

Usage

EXR_C02_S03_06

Format

A [data.frame](#) with 60 rows and 1 columns

EXR_C02_S03_07 *Data of Exercise 2.3.7*

Description

EXR_C02_S03_07

Usage

EXR_C02_S03_07

Format

A [data.frame](#) with 155 rows and 1 columns

EXR_C02_S03_08 *Data of Exercise 2.3.8*

Description

EXR_C02_S03_08

Usage

EXR_C02_S03_08

Format

A [data.frame](#) with 30 rows and 1 columns

EXR_C02_S03_09 *Data of Exercise 2.3.9*

Description

EXR_C02_S03_09

Usage

EXR_C02_S03_09

Format

A [data.frame](#) with 50 rows and 2 columns

EXR_C02_S03_11 *Data of Exercise 2.3.11*

Description

EXR_C02_S03_11

Usage

EXR_C02_S03_11

Format

A [data.frame](#) with 216 rows and 1 columns

EXR_C02_S03_12 *Data of Exercise 2.3.12*

Description

EXR_C02_S03_12

Usage

EXR_C02_S03_12

Format

A [data.frame](#) with 109 rows and 1 columns

EXR_C02_S05_03 *Data of Exercise 2.5.3*

Description

EXR_C02_S05_03

Usage

EXR_C02_S05_03

Format

A [data.frame](#) with 30 rows and 1 columns

EXR_C02_S05_06 *Data of Exercise 2.5.6*

Description

EXR_C02_S05_06

Usage

EXR_C02_S05_06

Format

A [data.frame](#) with 20 rows and 1 columns

EXR_C06_S02_05 *Data of Exercise 6.2.5*

Description

EXR_C06_S02_05

Usage

EXR_C06_S02_05

Format

A [data.frame](#) with 16 rows and 1 columns

EXR_C06_S04_10 *Data of Exercise 6.4.10*

Description

EXR_C06_S04_10

Usage

EXR_C06_S04_10

Format

A [data.frame](#) with 32 rows and 2 columns

EXR_C06_S09_07 *Data of Exercise 6.9.7*

Description

EXR_C06_S09_07

Usage

EXR_C06_S09_07

Format

A [data.frame](#) with 20 rows and 1 columns

EXR_C06_S10_07 *Data of Exercise 6.10.7*

Description

EXR_C06_S10_07

Usage

EXR_C06_S10_07

Format

A [data.frame](#) with 26 rows and 2 columns

EXR_C07_S02_13 *Data of Exercise 7.2.13*

Description

EXR_C07_S02_13

Usage

EXR_C07_S02_13

Format

A [data.frame](#) with 20 rows and 1 columns

EXR_C07_S02_15 *Data of Exercise 7.2.15*

Description

EXR_C07_S02_15

Usage

EXR_C07_S02_15

Format

A [data.frame](#) with 50 rows and 1 columns

EXR_C07_S02_16 *Data of Exercise 7.2.16*

Description

EXR_C07_S02_16

Usage

EXR_C07_S02_16

Format

A [data.frame](#) with 21 rows and 1 columns

EXR_C07_S03_03 *Data of Exercise 7.3.3*

Description

EXR_C07_S03_03

Usage

EXR_C07_S03_03

Format

A [data.frame](#) with 63 rows and 2 columns

EXR_C07_S03_04 *Data of Exercise 7.3.4*

Description

EXR_C07_S03_04

Usage

EXR_C07_S03_04

Format

A [data.frame](#) with 174 rows and 2 columns

EXR_C07_S03_05 *Data of Exercise 7.3.5*

Description

EXR_C07_S03_05

Usage

EXR_C07_S03_05

Format

A [data.frame](#) with 82 rows and 2 columns

EXR_C07_S03_10 *Data of Exercise 7.3.10*

Description

EXR_C07_S03_10

Usage

EXR_C07_S03_10

Format

A [data.frame](#) with 24 rows and 2 columns

EXR_C07_S03_11 *Data of Exercise 7.3.11*

Description

EXR_C07_S03_11

Usage

EXR_C07_S03_11

Format

A [data.frame](#) with 20 rows and 2 columns

EXR_C07_S03_12 *Data of Exercise 7.3.12*

Description

EXR_C07_S03_12

Usage

EXR_C07_S03_12

Format

A [data.frame](#) with 90 rows and 2 columns

EXR_C07_S04_01 *Data of Exercise 7.4.1*

Description

EXR_C07_S04_01

Usage

EXR_C07_S04_01

Format

A [data.frame](#) with 15 rows and 2 columns

EXR_C07_S04_02 *Data of Exercise 7.4.2*

Description

EXR_C07_S04_02

Usage

EXR_C07_S04_02

Format

A [data.frame](#) with 66 rows and 2 columns

EXR_C07_S04_03 *Data of Exercise 7.4.3*

Description

EXR_C07_S04_03

Usage

EXR_C07_S04_03

Format

A [data.frame](#) with 11 rows and 2 columns

EXR_C07_S04_04 *Data of Exercise 7.4.4*

Description

EXR_C07_S04_04

Usage

EXR_C07_S04_04

Format

A [data.frame](#) with 20 rows and 2 columns

EXR_C07_S08_07 *Data of Exercise 7.8.7*

Description

EXR_C07_S08_07

Usage

EXR_C07_S08_07

Format

A [data.frame](#) with 23 rows and 2 columns

EXR_C08_S02_01 *Data of Exercise 8.2.1*

Description

EXR_C08_S02_01

Usage

EXR_C08_S02_01

Format

A [data.frame](#) with 329 rows and 2 columns

EXR_C08_S02_02 *Data of Exercise 8.2.2*

Description

EXR_C08_S02_02

Usage

EXR_C08_S02_02

Format

A [data.frame](#) with 96 rows and 2 columns

EXR_C08_S02_03 *Data of Exercise 8.2.3*

Description

EXR_C08_S02_03

Usage

EXR_C08_S02_03

Format

A [data.frame](#) with 113 rows and 2 columns

EXR_C08_S02_04 *Data of Exercise 8.2.4*

Description

EXR_C08_S02_04

Usage

EXR_C08_S02_04

Format

A [data.frame](#) with 164 rows and 2 columns

EXR_C08_S02_05 *Data of Exercise 8.2.5*

Description

EXR_C08_S02_05

Usage

EXR_C08_S02_05

Format

A [data.frame](#) with 29 rows and 2 columns

EXR_C08_S02_06 *Data of Exercise 8.2.6*

Description

EXR_C08_S02_06

Usage

EXR_C08_S02_06

Format

A [data.frame](#) with 90 rows and 2 columns

EXR_C08_S02_07 *Data of Exercise 8.2.7*

Description

EXR_C08_S02_07

Usage

EXR_C08_S02_07

Format

A [data.frame](#) with 178 rows and 2 columns

EXR_C08_S03_01 *Data of Exercise 8.3.1*

Description

EXR_C08_S03_01

Usage

EXR_C08_S03_01

Format

A [data.frame](#) with 96 rows and 3 columns

EXR_C08_S03_02 *Data of Exercise 8.3.2*

Description

EXR_C08_S03_02

Usage

EXR_C08_S03_02

Format

A [data.frame](#) with 10 rows and 5 columns

EXR_C08_S03_03 *Data of Exercise 8.3.3*

Description

EXR_C08_S03_03

Usage

EXR_C08_S03_03

Format

A [data.frame](#) with 20 rows and 3 columns

EXR_C08_S03_04 *Data of Exercise 8.3.4*

Description

EXR_C08_S03_04

Usage

EXR_C08_S03_04

Format

A [data.frame](#) with 16 rows and 3 columns

EXR_C08_S03_05 *Data of Exercise 8.3.5*

Description

EXR_C08_S03_05

Usage

EXR_C08_S03_05

Format

A [data.frame](#) with 12 rows and 3 columns

EXR_C08_S04_01 *Data of Exercise 8.4.1*

Description

EXR_C08_S04_01

Usage

EXR_C08_S04_01

Format

A [data.frame](#) with 40 rows and 3 columns

EXR_C08_S04_02 *Data of Exercise 8.4.2*

Description

EXR_C08_S04_02

Usage

EXR_C08_S04_02

Format

A [data.frame](#) with 35 rows and 3 columns

EXR_C08_S04_03 *Data of Exercise 8.4.3*

Description

EXR_C08_S04_03

Usage

EXR_C08_S04_03

Format

A [data.frame](#) with 48 rows and 3 columns

EXR_C08_S04_06 *Data of Exercise 8.4.6*

Description

EXR_C08_S04_06

Usage

EXR_C08_S04_06

Format

A [data.frame](#) with 20 rows and 6 columns

EXR_C08_S05_01 *Data of Exercise 8.5.1*

Description

EXR_C08_S05_01

Usage

EXR_C08_S05_01

Format

A [data.frame](#) with 24 rows and 3 columns

EXR_C08_S05_02 *Data of Exercise 8.5.2*

Description

EXR_C08_S05_02

Usage

EXR_C08_S05_02

Format

A [data.frame](#) with 72 rows and 3 columns

EXR_C08_S05_03 *Data of Exercise 8.5.3*

Description

EXR_C08_S05_03

Usage

EXR_C08_S05_03

Format

A [data.frame](#) with 44 rows and 3 columns

EXR_C08_S05_04 *Data of Exercise 8.5.4*

Description

EXR_C08_S05_04

Usage

EXR_C08_S05_04

Format

A [data.frame](#) with 13 rows and 3 columns

EXR_C09_S03_02 *Data of Exercise 9.3.2*

Description

EXR_C09_S03_02

Usage

EXR_C09_S03_02

Format

A [data.frame](#) with 10 rows and 2 columns

EXR_C09_S03_03 *Data of Exercise 9.3.3*

Description

EXR_C09_S03_03

Usage

EXR_C09_S03_03

Format

A [data.frame](#) with 17 rows and 2 columns

EXR_C09_S03_04 *Data of Exercise 9.3.4*

Description

EXR_C09_S03_04

Usage

EXR_C09_S03_04

Format

A [data.frame](#) with 90 rows and 2 columns

EXR_C09_S03_06 *Data of Exercise 9.3.6*

Description

EXR_C09_S03_06

Usage

EXR_C09_S03_06

Format

A [data.frame](#) with 22 rows and 2 columns

EXR_C09_S03_07 *Data of Exercise 9.3.7*

Description

EXR_C09_S03_07

Usage

EXR_C09_S03_07

Format

A [data.frame](#) with 27 rows and 2 columns

EXR_C09_S07_01 *Data of Exercise 9.7.1*

Description

EXR_C09_S07_01

Usage

EXR_C09_S07_01

Format

A [data.frame](#) with 20 rows and 2 columns

EXR_C09_S07_02 *Data of Exercise 9.7.2*

Description

EXR_C09_S07_02

Usage

EXR_C09_S07_02

Format

A [data.frame](#) with 90 rows and 2 columns

EXR_C09_S07_04 *Data of Exercise 9.7.4*

Description

EXR_C09_S07_04

Usage

EXR_C09_S07_04

Format

A [data.frame](#) with 18 rows and 2 columns

EXR_C09_S07_05 *Data of Exercise 9.7.5*

Description

EXR_C09_S07_05

Usage

EXR_C09_S07_05

Format

A [data.frame](#) with 30 rows and 2 columns

EXR_C09_S07_06 *Data of Exercise 9.7.6*

Description

EXR_C09_S07_06

Usage

EXR_C09_S07_06

Format

A [data.frame](#) with 15 rows and 2 columns

EXR_C10_S03_01 *Data of Exercise 10.3.1*

Description

EXR_C10_S03_01

Usage

EXR_C10_S03_01

Format

A [data.frame](#) with 35 rows and 3 columns

EXR_C10_S03_02 *Data of Exercise 10.3.2*

Description

EXR_C10_S03_02

Usage

EXR_C10_S03_02

Format

A [data.frame](#) with 100 rows and 4 columns

EXR_C10_S03_03 *Data of Exercise 10.3.3*

Description

EXR_C10_S03_03

Usage

EXR_C10_S03_03

Format

A [data.frame](#) with 10 rows and 3 columns

EXR_C10_S03_04 *Data of Exercise 10.3.4*

Description

EXR_C10_S03_04

Usage

EXR_C10_S03_04

Format

A [data.frame](#) with 20 rows and 3 columns

EXR_C10_S03_05 *Data of Exercise 10.3.5*

Description

EXR_C10_S03_05

Usage

EXR_C10_S03_05

Format

A [data.frame](#) with 25 rows and 3 columns

EXR_C10_S03_06 *Data of Exercise 10.3.6*

Description

EXR_C10_S03_06

Usage

EXR_C10_S03_06

Format

A [data.frame](#) with 20 rows and 8 columns

EXR_C10_S06_01 *Data of Exercise 10.6.1*

Description

EXR_C10_S06_01

Usage

EXR_C10_S06_01

Format

A [data.frame](#) with 40 rows and 4 columns

EXR_C10_S06_02 *Data of Exercise 10.6.2*

Description

EXR_C10_S06_02

Usage

EXR_C10_S06_02

Format

A [data.frame](#) with 12 rows and 3 columns

EXR_C10_S06_03 *Data of Exercise 10.6.3*

Description

EXR_C10_S06_03

Usage

EXR_C10_S06_03

Format

A [data.frame](#) with 15 rows and 3 columns

EXR_C10_S06_04	<i>Data of Exercise 10.6.4</i>
----------------	--------------------------------

Description

EXR_C10_S06_04

Usage

EXR_C10_S06_04

FormatA [data.frame](#) with 15 rows and 6 columns

EXR_C11_S02_01	<i>Data of Exercise 11.2.1</i>
----------------	--------------------------------

Description

EXR_C11_S02_01

Usage

EXR_C11_S02_01

FormatA [data.frame](#) with 44 rows and 3 columns

EXR_C11_S02_02	<i>Data of Exercise 11.2.2</i>
----------------	--------------------------------

Description

EXR_C11_S02_02

Usage

EXR_C11_S02_02

FormatA [data.frame](#) with 100 rows and 3 columns

EXR_C11_S02_03 *Data of Exercise 11.2.3*

Description

EXR_C11_S02_03

Usage

EXR_C11_S02_03

Format

A [data.frame](#) with 17 rows and 3 columns

EXR_C11_S02_04 *Data of Exercise 11.2.4*

Description

EXR_C11_S02_04

Usage

EXR_C11_S02_04

Format

A [data.frame](#) with 90 rows and 3 columns

EXR_C11_S03_01 *Data of Exercise 11.3.1*

Description

EXR_C11_S03_01

Usage

EXR_C11_S03_01

Format

A [data.frame](#) with 100 rows and 8 columns

EXR_C11_S03_02 *Data of Exercise 11.3.2*

Description

EXR_C11_S03_02

Usage

EXR_C11_S03_02

Format

A [data.frame](#) with 35 rows and 7 columns

EXR_C11_S03_03 *Data of Exercise 11.3.3*

Description

EXR_C11_S03_03

Usage

EXR_C11_S03_03

Format

A [data.frame](#) with 68 rows and 7 columns

EXR_C11_S04_01 *Data of Exercise 11.4.1*

Description

EXR_C11_S04_01

Usage

EXR_C11_S04_01

Format

A [data.frame](#) with 4 rows and 3 columns

EXR_C11_S04_02 *Data of Exercise 11.4.2*

Description

EXR_C11_S04_02

Usage

EXR_C11_S04_02

Format

A [data.frame](#) with 184 rows and 2 columns

EXR_C11_S05_01 *Data of Exercise 11.5.1*

Description

EXR_C11_S05_01

Usage

EXR_C11_S05_01

Format

A [data.frame](#) with 45 rows and 4 columns

EXR_C11_S05_02 *Data of Exercise 11.5.2*

Description

EXR_C11_S05_02

Usage

EXR_C11_S05_02

Format

A [data.frame](#) with 45 rows and 5 columns

EXR_C11_S05_04 *Data of Exercise 11.5.4*

Description

EXR_C11_S05_04

Usage

EXR_C11_S05_04

Format

A [data.frame](#) with 90 rows and 3 columns

EXR_C13_S05_01 *Data of Exercise 13.5.1*

Description

EXR_C13_S05_01

Usage

EXR_C13_S05_01

Format

A [data.frame](#) with 30 rows and 2 columns

EXR_C13_S05_02 *Data of Exercise 13.5.2*

Description

EXR_C13_S05_02

Usage

EXR_C13_S05_02

Format

A [data.frame](#) with 30 rows and 2 columns

EXR_C13_S06_01 *Data of Exercise 13.6.1*

Description

EXR_C13_S06_01

Usage

EXR_C13_S06_01

Format

A [data.frame](#) with 70 rows and 2 columns

EXR_C13_S06_02 *Data of Exercise 13.6.2*

Description

EXR_C13_S06_02

Usage

EXR_C13_S06_02

Format

A [data.frame](#) with 17 rows and 2 columns

EXR_C13_S06_03 *Data of Exercise 13.6.3*

Description

EXR_C13_S06_03

Usage

EXR_C13_S06_03

Format

A [data.frame](#) with 83 rows and 2 columns

EXR_C13_S07_02 *Data of Exercise 13.7.2*

Description

EXR_C13_S07_02

Usage

EXR_C13_S07_02

Format

A [data.frame](#) with 30 rows and 1 columns

EXR_C13_S08_01 *Data of Exercise 13.8.1*

Description

EXR_C13_S08_01

Usage

EXR_C13_S08_01

Format

A [data.frame](#) with 232 rows and 2 columns

EXR_C13_S08_02 *Data of Exercise 13.8.2*

Description

EXR_C13_S08_02

Usage

EXR_C13_S08_02

Format

A [data.frame](#) with 15 rows and 2 columns

EXR_C13_S08_03 *Data of Exercise 13.8.3*

Description

EXR_C13_S08_03

Usage

EXR_C13_S08_03

Format

A [data.frame](#) with 53 rows and 2 columns

EXR_C13_S08_04 *Data of Exercise 13.8.4*

Description

EXR_C13_S08_04

Usage

EXR_C13_S08_04

Format

A [data.frame](#) with 22 rows and 2 columns

EXR_C13_S08_05 *Data of Exercise 13.8.5*

Description

EXR_C13_S08_05

Usage

EXR_C13_S08_05

Format

A [data.frame](#) with 44 rows and 2 columns

EXR_C13_S08_06 *Data of Exercise 13.8.6*

Description

EXR_C13_S08_06

Usage

EXR_C13_S08_06

Format

A [data.frame](#) with 22 rows and 3 columns

EXR_C13_S09_01 *Data of Exercise 13.9.1*

Description

EXR_C13_S09_01

Usage

EXR_C13_S09_01

Format

A [data.frame](#) with 9 rows and 4 columns

EXR_C13_S09_02 *Data of Exercise 13.9.2*

Description

EXR_C13_S09_02

Usage

EXR_C13_S09_02

Format

A [data.frame](#) with 15 rows and 11 columns

EXR_C13_S09_03 *Data of Exercise 13.9.3*

Description

EXR_C13_S09_03

Usage

EXR_C13_S09_03

Format

A [data.frame](#) with 10 rows and 6 columns

EXR_C13_S10_01 *Data of Exercise 13.10.1*

Description

EXR_C13_S10_01

Usage

EXR_C13_S10_01

Format

A [data.frame](#) with 15 rows and 3 columns

EXR_C13_S10_02 *Data of Exercise 13.10.2*

Description

EXR_C13_S10_02

Usage

EXR_C13_S10_02

Format

A [data.frame](#) with 10 rows and 3 columns

EXR_C13_S10_03 *Data of Exercise 13.10.3*

Description

EXR_C13_S10_03

Usage

EXR_C13_S10_03

Format

A [data.frame](#) with 20 rows and 2 columns

EXR_C13_S10_04 *Data of Exercise 13.10.4*

Description

EXR_C13_S10_04

Usage

EXR_C13_S10_04

Format

A [data.frame](#) with 20 rows and 2 columns

EXR_C13_S10_05 *Data of Exercise 13.10.5*

Description

EXR_C13_S10_05

Usage

EXR_C13_S10_05

Format

A [data.frame](#) with 30 rows and 2 columns

EXR_C13_S10_06 *Data of Exercise 13.10.6*

Description

EXR_C13_S10_06

Usage

EXR_C13_S10_06

Format

A [data.frame](#) with 17 rows and 3 columns

EXR_C14_S03_01 *Data of Exercise 14.3.1*

Description

EXR_C14_S03_01

Usage

EXR_C14_S03_01

Format

A [data.frame](#) with 53 rows and 3 columns

EXR_C14_S03_02 *Data of Exercise 14.3.2*

Description

EXR_C14_S03_02

Usage

EXR_C14_S03_02

Format

A [data.frame](#) with 62 rows and 2 columns

EXR_C14_S04_03	<i>Data of Exercise 14.4.3</i>
----------------	--------------------------------

Description

EXR_C14_S04_03

Usage

EXR_C14_S04_03

FormatA [data.frame](#) with 50 rows and 4 columns

freqs-class	<i>S4 Class freqs</i>
-------------	---------------------------------------

DescriptionS4 Class [freqs](#)**Slots**.Data [integer](#) vector, frequency countsdata.name [character](#) integer, name of the data, only used in output

Gosset_Welch	<i>Two-Sample Student's t-statistic and Welch–Satterthwaite Equation</i>
--------------	---

DescriptionTo determine the degree of freedom, as well as the standard error, of two-sample t -statistic, with or without the equal-variance assumption.**Usage**

Gosset_Welch(s1, s2, v1 = s1^2, v2 = s2^2, n1, n2, var.equal = FALSE)

Arguments

<code>s1, s2</code>	(optional) double vectors, sample standard deviations of the two samples
<code>v1, v2</code>	double vectors, sample variances of the two samples, default $v_1 = s_1^2, v_2 = s_2^2$.
<code>n1, n2</code>	integer vectors, sample sizes of the two samples
<code>var.equal</code>	logical scalar, whether to treat the two variances as being equal when calculating the degree of freedom and the standard error of the mean-difference. If FALSE (default), Welch–Satterthwaite equation is used. If TRUE, the original two-sample <i>t</i> -test from William Sealy Gosset is used.

Value

`Gosset_Welch` returns a **numeric** scalar of the degree of freedom, with a **numeric** scalar attribute 'std.err' of the standard error of the mean-difference.

References

Student's *t*-test by William Sealy Gosset, [doi:10.1093/biomet/6.1.1](https://doi.org/10.1093/biomet/6.1.1).

Welch–Satterthwaite equation by Bernard Lewis Welch and F. E. Satterthwaite [doi:10.2307/3002019](https://doi.org/10.2307/3002019) and [doi:10.1093/biomet/34.12.28](https://doi.org/10.1093/biomet/34.12.28).

See Also

[t.test](#)

Examples

```
x = rnorm(32L, sd = 1.6); y = rnorm(57L, sd = 2.1)
vx = var(x); vy = var(y); nx = length(x); ny = length(y)
t.test(x, y, var.equal = FALSE)[c('parameter', 'stderr')]
Gosset_Welch(v1 = vx, v2 = vy, n1 = nx, n2 = ny, var.equal = FALSE)
t.test(x, y, var.equal = TRUE)[c('parameter', 'stderr')]
Gosset_Welch(v1 = vx, v2 = vy, n1 = nx, n2 = ny, var.equal = TRUE)
```

Description

LDS_C02_NCBIRTH800

Usage

LDS_C02_NCBIRTH800

Format

A [data.frame](#) with 800 rows and 14 columns

LDS_C06_BABYWGTS *Large Data BABYWGTS from Chapter 6*

Description

LDS_C06_BABYWGTS

Usage

LDS_C06_BABYWGTS

Format

A [data.frame](#) with 1200 rows and 2 columns

LDS_C06_BOYHGTS *Large Data BOYHGTS from Chapter 6*

Description

LDS_C06_BOYHGTS

Usage

LDS_C06_BOYHGTS

Format

A [data.frame](#) with 1000 rows and 2 columns

LDS_C06_CHOLEST *Large Data CHOLEST from Chapter 6*

Description

LDS_C06_CHOLEST

Usage

LDS_C06_CHOLEST

Format

A [data.frame](#) with 1000 rows and 2 columns

LDS_C07_HEADCIRC *Large Data HEADCIRC from Chapter 7*

Description

LDS_C07_HEADCIRC

Usage

LDS_C07_HEADCIRC

Format

A [data.frame](#) with 1000 rows and 3 columns

LDS_C07_HEMOGLOB *Large Data HEMOglob from Chapter 7*

Description

LDS_C07_HEMOGLOB

Usage

LDS_C07_HEMOGLOB

Format

A [data.frame](#) with 1000 rows and 2 columns

LDS_C07_MANDEXT *Large Data MANDEXT from Chapter 7*

Description

LDS_C07_MANDEXT

Usage

LDS_C07_MANDEXT

Format

A [data.frame](#) with 1000 rows and 2 columns

LDS_C07_PCKDATA	<i>Large Data PCKDATA from Chapter 7</i>
-----------------	--

Description

LDS_C07_PCKDATA

Usage

LDS_C07_PCKDATA

FormatA [data.frame](#) with 1005 rows and 3 columns

LDS_C07_PROTHROM	<i>Large Data PROTHROM from Chapter 7</i>
------------------	---

Description

LDS_C07_PROTHROM

Usage

LDS_C07_PROTHROM

FormatA [data.frame](#) with 1000 rows and 2 columns

LDS_C08_CSFDATA	<i>Large Data CSFDATA from Chapter 8</i>
-----------------	--

Description

LDS_C08_CSFDATA

Usage

LDS_C08_CSFDATA

FormatA [data.frame](#) with 300 rows and 6 columns

LDS_C08_LSADATA *Large Data LSADATA from Chapter 8*

Description

LDS_C08_LSADATA

Usage

LDS_C08_LSADATA

Format

A [data.frame](#) with 350 rows and 5 columns

LDS_C08_MEDSCORES *Large Data MEDSCORES from Chapter 8*

Description

LDS_C08_MEDSCORES

Usage

LDS_C08_MEDSCORES

Format

A [data.frame](#) with 582 rows and 4 columns

LDS_C08_RBCDATA *Large Data RBCDATA from Chapter 8*

Description

LDS_C08_RBCDATA

Usage

LDS_C08_RBCDATA

Format

A [data.frame](#) with 350 rows and 4 columns

LDS_C08_SACEDATA *Large Data SACEDATA from Chapter 8*

Description

LDS_C08_SACEDATA

Usage

LDS_C08_SACEDATA

Format

A [data.frame](#) with 400 rows and 5 columns

LDS_C08_SERUMCHO *Large Data SERUMCHO from Chapter 8*

Description

LDS_C08_SERUMCHO

Usage

LDS_C08_SERUMCHO

Format

A [data.frame](#) with 347 rows and 4 columns

LDS_C09_CALCIIUM *Large Data CALCIIUM from Chapter 9*

Description

LDS_C09_CALCIIUM

Usage

LDS_C09_CALCIIUM

Format

A [data.frame](#) with 100 rows and 13 columns

LDS_C09_CEREBRAL *Large Data CEREBRAL from Chapter 9*

Description

LDS_C09_CEREBRAL

Usage

LDS_C09_CEREBRAL

Format

A [data.frame](#) with 1050 rows and 3 columns

LDS_C09_HYPERTEN *Large Data HYPERTEN from Chapter 9*

Description

LDS_C09_HYPERTEN

Usage

LDS_C09_HYPERTEN

Format

A [data.frame](#) with 1050 rows and 3 columns

LDS_C10_LTEXER *Large Data LTEXER from Chapter 10*

Description

LDS_C10_LTEXER

Usage

LDS_C10_LTEXER

Format

A [data.frame](#) with 248 rows and 5 columns

LDS_C10_RESPDIS	<i>Large Data RESPDIS from Chapter 10</i>
-----------------	---

Description

LDS_C10_RESPDIS

Usage

LDS_C10_RESPDIS

FormatA [data.frame](#) with 1200 rows and 6 columns

LDS_C10_RISKFACT	<i>Large Data RISKFACT from Chapter 10</i>
------------------	--

Description

LDS_C10_RISKFACT

Usage

LDS_C10_RISKFACT

FormatA [data.frame](#) with 1000 rows and 6 columns

LDS_C10_STERLENGTH	<i>Large Data STERLENGTH from Chapter 10</i>
--------------------	--

Description

LDS_C10_STERLENGTH

Usage

LDS_C10_STERLENGTH

FormatA [data.frame](#) with 1162 rows and 4 columns

LDS_C11_AQUATICS *Large Data AQUATICS from Chapter 11*

Description

LDS_C11_AQUATICS

Usage

LDS_C11_AQUATICS

Format

A [data.frame](#) with 142 rows and 7 columns

LDS_C11_TEACHERS *Large Data TEACHERS from Chapter 11*

Description

LDS_C11_TEACHERS

Usage

LDS_C11_TEACHERS

Format

A [data.frame](#) with 212 rows and 7 columns

LDS_C11_WGTLOSS *Large Data WGTLOSS from Chapter 11*

Description

LDS_C11_WGTLOSS

Usage

LDS_C11_WGTLOSS

Format

A [data.frame](#) with 1185 rows and 3 columns

LDS_C12_SMOKING	<i>Large Data SMOKING from Chapter 12</i>
-----------------	---

Description

LDS_C12_SMOKING

Usage

LDS_C12_SMOKING

FormatA [data.frame](#) with 1200 rows and 6 columns

LDS_C13_KLETTER	<i>Large Data KLETTER from Chapter 13</i>
-----------------	---

Description

LDS_C13_KLETTER

Usage

LDS_C13_KLETTER

FormatA [data.frame](#) with 168 rows and 3 columns

power-class	<i>S4 class power</i>
-------------	-----------------------

DescriptionS4 class [power](#)**Slots**.Data [numeric](#) scalar or vector, power(s) calculated at alternative parameter(s) μ_1 x [numeric](#) scalar or vector, alternative parameter(s) μ_1 rr [RejectionRegion](#) object

RejectionRegion-class *S4 class [RejectionRegion](#)*

Description

S4 class [RejectionRegion](#)

Slots

.Data [numeric](#) scalar for one-sided test, or length-two [numeric](#) vector for two-sided test
 null.value [numeric](#) scalar, null value
 std.err [numeric](#) scalar, standard error of sampling distribution
 parameter [numeric](#) vector, additional parameters
 alternative [character](#) scalar, alternative hypothesis
 sig.level [numeric](#) scalar, significance level (Type I error probability)
 test [character](#) scalar, type of test. Currently only 'z' is supported

REV_C02_13

Review Exercise 13 of Chapter 2

Description

REV_C02_13

Usage

REV_C02_13

Format

A [data.frame](#) with 50 rows and 1 columns

REV_C02_15

Review Exercise 15 of Chapter 2

Description

REV_C02_15

Usage

REV_C02_15

Format

A [data.frame](#) with 28 rows and 1 columns

REV_C02_16

Review Exercise 16 of Chapter 2

Description

REV_C02_16

Usage

REV_C02_16

Format

A [data.frame](#) with 53 rows and 2 columns

REV_C02_19

Review Exercise 19 of Chapter 2

Description

REV_C02_19

Usage

REV_C02_19

Format

A [data.frame](#) with 22 rows and 1 columns

REV_C02_29

Review Exercise 29 of Chapter 2

Description

REV_C02_29

Usage

REV_C02_29

Format

A [data.frame](#) with 107 rows and 1 columns

REV_C06_22 *Review Exercise 22 of Chapter 6*

Description

REV_C06_22

Usage

REV_C06_22

Format

A [data.frame](#) with 27 rows and 2 columns

REV_C06_23 *Review Exercise 23 of Chapter 6*

Description

REV_C06_23

Usage

REV_C06_23

Format

A [data.frame](#) with 28 rows and 2 columns

REV_C06_28 *Review Exercise 28 of Chapter 6*

Description

REV_C06_28

Usage

REV_C06_28

Format

A [data.frame](#) with 110 rows and 2 columns

REV_C07_18 *Review Exercise 18 of Chapter 7*

Description

REV_C07_18

Usage

REV_C07_18

Format

A [data.frame](#) with 107 rows and 3 columns

REV_C07_19 *Review Exercise 19 of Chapter 7*

Description

REV_C07_19

Usage

REV_C07_19

Format

A [data.frame](#) with 107 rows and 3 columns

REV_C07_22 *Review Exercise 22 of Chapter 7*

Description

REV_C07_22

Usage

REV_C07_22

Format

A [data.frame](#) with 76 rows and 2 columns

REV_C07_24 *Review Exercise 24 of Chapter 7*

Description

REV_C07_24

Usage

REV_C07_24

Format

A [data.frame](#) with 37 rows and 2 columns

REV_C07_29 *Review Exercise 29 of Chapter 7*

Description

REV_C07_29

Usage

REV_C07_29

Format

A [data.frame](#) with 12 rows and 2 columns

REV_C07_40 *Review Exercise 40 of Chapter 7*

Description

REV_C07_40

Usage

REV_C07_40

Format

A [data.frame](#) with 8 rows and 3 columns

REV_C07_41 *Review Exercise 41 of Chapter 7*

Description

REV_C07_41

Usage

REV_C07_41

Format

A [data.frame](#) with 11 rows and 3 columns

REV_C07_42 *Review Exercise 42 of Chapter 7*

Description

REV_C07_42

Usage

REV_C07_42

Format

A [data.frame](#) with 10 rows and 6 columns

REV_C07_43 *Review Exercise 43 of Chapter 7*

Description

REV_C07_43

Usage

REV_C07_43

Format

A [data.frame](#) with 31 rows and 2 columns

REV_C07_44 *Review Exercise 44 of Chapter 7*

Description

REV_C07_44

Usage

REV_C07_44

Format

A [data.frame](#) with 98 rows and 4 columns

REV_C07_45 *Review Exercise 45 of Chapter 7*

Description

REV_C07_45

Usage

REV_C07_45

Format

A [data.frame](#) with 15 rows and 11 columns

REV_C07_46 *Review Exercise 46 of Chapter 7*

Description

REV_C07_46

Usage

REV_C07_46

Format

A [data.frame](#) with 17 rows and 2 columns

REV_C07_47

Review Exercise 47 of Chapter 7

Description

REV_C07_47

Usage

REV_C07_47

Format

A [data.frame](#) with 66 rows and 2 columns

REV_C07_48

Review Exercise 48 of Chapter 7

Description

REV_C07_48

Usage

REV_C07_48

Format

A [data.frame](#) with 51 rows and 2 columns

REV_C07_49

Review Exercise 49 of Chapter 7

Description

REV_C07_49

Usage

REV_C07_49

Format

A [data.frame](#) with 22 rows and 2 columns

REV_C07_50 *Review Exercise 50 of Chapter 7*

Description

REV_C07_50

Usage

REV_C07_50

Format

A [data.frame](#) with 28 rows and 4 columns

REV_C07_51 *Review Exercise 51 of Chapter 7*

Description

REV_C07_51

Usage

REV_C07_51

Format

A [data.frame](#) with 22 rows and 2 columns

REV_C07_52 *Review Exercise 52 of Chapter 7*

Description

REV_C07_52

Usage

REV_C07_52

Format

A [data.frame](#) with 24 rows and 2 columns

REV_C07_53

Review Exercise 53 of Chapter 7

Description

REV_C07_53

Usage

REV_C07_53

Format

A [data.frame](#) with 55 rows and 2 columns

REV_C07_54

Review Exercise 54 of Chapter 7

Description

REV_C07_54

Usage

REV_C07_54

Format

A [data.frame](#) with 17 rows and 2 columns

REV_C07_55

Review Exercise 55 of Chapter 7

Description

REV_C07_55

Usage

REV_C07_55

Format

A [data.frame](#) with 50 rows and 2 columns

REV_C08_13 *Review Exercise 13 of Chapter 8*

Description

REV_C08_13

Usage

REV_C08_13

Format

A [data.frame](#) with 75 rows and 2 columns

REV_C08_14 *Review Exercise 14 of Chapter 8*

Description

REV_C08_14

Usage

REV_C08_14

Format

A [data.frame](#) with 91 rows and 2 columns

REV_C08_15 *Review Exercise 15 of Chapter 8*

Description

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Description

REV_C10_19

Usage

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FormatA [data.frame](#) with 70 rows and 5 columns

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FormatA [data.frame](#) with 28 rows and 3 columns

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REV_C11_28 *Review Exercise 28 of Chapter 11*

Description

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REV_C11_28

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Description

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Description

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show,BooleanRisk-method

Show [BooleanRisk](#) Object

Description

Show [BooleanRisk](#) object

Usage

```
## S4 method for signature 'BooleanRisk'  
show(object)
```


Arguments

object a [BooleanRisk](#) object

Value

The [show](#) method for [BooleanRisk](#) object does not have a returned value.

show, BooleanTest-method

Show [BooleanTest](#) Object

Description

Show [BooleanTest](#) object

Usage

```
## S4 method for signature 'BooleanTest'  
show(object)
```

Arguments

object a [BooleanTest](#) object

Value

The [show](#) method for [BooleanTest](#) object does not have a returned value.

show, freqs-method

Show [freqs](#) Object

Description

Show [freqs](#) object

Usage

```
## S4 method for signature 'freqs'  
show(object)
```

Arguments

object an [freqs](#) object

Value

The [show](#) method for [freqs](#) object does not have a returned value.

show,power-method	<i>Show power Object</i>
-------------------	--------------------------

Description

Show [power](#) object

Usage

```
## S4 method for signature 'power'  
show(object)
```

Arguments

object a [power](#) object

Value

The [show](#) method for [power](#) object does not have a returned value.

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