

Package ‘ANSM5’

January 20, 2025

Title Functions and Data for the Book ``Applied Nonparametric Statistical Methods'', 5th Edition

Version 1.1.1

Description Functions and data to accompany the 5th edition of the book ``Applied Nonparametric Statistical Methods'' (4th edition: Sprent & Smeeton, 2024, ISBN:158488701X), the revisions from the 4th edition including a move from describing the output from a miscellany of statistical software packages to using R. While the output from many of the functions can also be obtained using a range of other R functions, this package provides functions in a unified setting and give output using both p-values and confidence intervals, exemplifying the book's approach of treating p-values as a guide to statistical importance and not an end product in their own right. Please note that in creating the ANSM5 package we do not claim to have produced software which is necessarily the most computationally efficient nor the most comprehensive.

Encoding UTF-8

RoxygenNote 7.2.1

Suggests testthat (>= 3.0.0)

Config/testthat.edition 3

Imports stats

Depends R (>= 2.10)

LazyData true

License GPL (>= 3)

NeedsCompilation no

Author Neil Spencer [aut, cre, cph] (<<https://orcid.org/0000-0002-6068-5887>>)

Maintainer Neil Spencer <neilhspencer@gmail.com>

Repository CRAN

Date/Publication 2024-08-31 22:30:02 UTC

Contents

ansari.bradley	3
app1	4

binom	5
blomqvist	6
bowker	7
breslow.day	8
bs	9
ch10	9
ch11	11
ch12	12
ch13	13
ch14	15
ch15	15
ch3	16
ch4	17
ch5	18
ch6	19
ch7	21
ch8	22
ch9	24
chisqtest.ANSM	25
cochran.q	26
cohen.kappa	27
conover	28
control.median	30
cox.stuart	31
cramer.von.mises	32
fishertest.ANSM	33
friedman	34
friedman.lsd	35
gehan.wilcoxon	36
hettmansperger.elmore	37
hodges.ajne	38
jonckheere.terpstra	39
kendall.concordance	40
kendall.tau	41
kruskal.wallis	42
kruskal.wallis.lsd	43
kruskal.wallis.vdW	44
kstest.ANSM	45
lik.ratio	46
lilliefors	47
linear.by.linear	48
logoddsratio.2x2	49
logrank	50
mantel.haenszel	51
med.test	52
mood	53
moses.extreme.reactions	54
noether	55

normal.scores.test	55
oddsratio.2x2diff	56
pearson	58
pearson.beta	59
peto.wilcoxon	60
pitman	61
print.ANSMstat	62
print.ANSMtest	63
rng.test	63
runs.2cat	64
runs.ncat	65
sgn.test	66
shapiro.test.ANSM	67
siegel.tukey	68
spearman	69
spearman.beta	70
theil.kendall	71
wilcoxon.mann.whitney	72
wilcoxon.signedrank	74
zelen	75

Index**76**

ansari.bradley *Perform Ansari-Bradley test*

Description

ansari.bradley() performs the Ansari-Bradley test and is used in chapter 6 of "Applied Non-parametric Statistical Methods" (5th edition)

Usage

```
ansari.bradley(  
  x,  
  y,  
  H0 = NULL,  
  alternative = c("two.sided", "less", "greater"),  
  max.exact.cases = 25,  
  do.asymp = FALSE,  
  do.exact = TRUE  
)
```

Arguments

x	Numeric vector
y	Numeric vector
H0	Null hypothesis value (defaults to NULL)
alternative	Type of alternative hypothesis (defaults to two.sided)
max.exact.cases	Maximum number of cases allowed for exact calculations (defaults to 25)
do.asymp	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
do.exact	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 6.12 from "Applied Nonparametric Statistical Methods" (5th edition)
ansari.bradley(ch6$typeA, ch6$typeB)

# Exercise 6.16 from "Applied Nonparametric Statistical Methods" (5th edition)
ansari.bradley(ch6$travel, ch6$politics)
```

Description

Data in Appendix 1 of "Applied Nonparametric Statistical Methods" (5th edition)

- McAlpha (used in example 4.5)
- McBeta (used in example 6.6)
- McGamma (used in exercise 4.1, example 6.6)
- McDelta (used in examples 10.4, 10.8, exercise 10.5)

Usage

app1

Format

app1:

A list with 4 data vectors

Source

"Applied Nonparametric Statistical Methods" (5th edition)

binom	<i>Perform Binomial test</i>
-------	------------------------------

Description

binom() performs the Binomial test and calculates the Binomial confidence interval and is used in chapters 4, 5 and 13 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
binom(  
  r,  
  n,  
  H0 = NULL,  
  alternative = c("two.sided", "less", "greater"),  
  CI.width = 0.95,  
  max.exact.cases = 1e+07,  
  do.asymp = FALSE,  
  do.exact = TRUE,  
  do.CI = TRUE  
)
```

Arguments

r	Number of successes
n	Number of trials
H0	Null hypothesis value (defaults to NULL)
alternative	Type of alternative hypothesis (defaults to two.sided)
CI.width	Confidence interval width (defaults to 0.95)
max.exact.cases	Maximum number of cases allowed for exact calculations (defaults to 10000000)
do.asymp	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
do.exact	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)
do.CI	Boolean indicating whether or not to perform confidence interval calculations (defaults to TRUE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 4.6 from "Applied Nonparametric Statistical Methods" (5th edition)
binom(3, 20)

# Exercise 5.8 from "Applied Nonparametric Statistical Methods" (5th edition)
binom(24, 40, 0.5)
```

blomqvist

Calculate Blomqvist coefficient

Description

`blomqvist()` calculates the Blomqvist coefficient and is used in chapter 10 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
blomqvist(
  x,
  y,
  alternative = c("two.sided", "less", "greater"),
  max.exact.cases = 1000,
  nsims.mc = 1e+05,
  seed = NULL,
  do.exact = TRUE,
  do.mc = FALSE
)
```

Arguments

<code>x</code>	Numeric vector of same length as <code>y</code>
<code>y</code>	Numeric vector of same length as <code>x</code>
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to 1000)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 100000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to <code>NULL</code>)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to <code>TRUE</code>)
<code>do.mc</code>	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to <code>FALSE</code>)

Value

An ANSMstat object with the results from applying the function

Examples

```
# Example 10.9 from "Applied Nonparametric Statistical Methods" (5th edition)
blomqvist(ch10$q1, ch10$q2, alternative = "greater")

# Exercise 10.7 from "Applied Nonparametric Statistical Methods" (5th edition)
blomqvist(ch10$ERA, ch10$SSS)
```

bowker

Perform Bowker's extension of McNemar's test

Description

`bowker()` performs the Bowker's extension of McNemar's test and is used in chapter 12 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
bowker(x, y = NULL, do.asymp = TRUE)
```

Arguments

- | | |
|----------|---|
| x | Factor of same length as y, or two-dimensional square table |
| y | Factor of same length as x (or NULL if x is table) (defaults to NULL) |
| do.asymp | Boolean indicating whether or not to perform asymptotic calculations (defaults to TRUE) |

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 12.12 from "Applied Nonparametric Statistical Methods" (5th edition)
bowker(ch12$side.effect.new, ch12$side.effect.old)

# Exercise 12.12 from "Applied Nonparametric Statistical Methods" (5th edition)
bowker(ch12$first.response, ch12$second.response)
```

breslow.day *Perform Breslow and Day test*

Description

`breslow.day()` performs the Breslow and Day test and is used in chapter 13 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
breslow.day(x, y, z, CI.width = 0.95, do.asymp = TRUE, do.CI = TRUE)
```

Arguments

x	Binary factor of same length as y, z
y	Binary factor of same length as x, z
z	Factor of same length as x, y
CI.width	Confidence interval width (defaults to 0.95)
do.asymp	Boolean indicating whether or not to perform asymptotic calculations (defaults to TRUE)
do.CI	Boolean indicating whether or not to perform confidence interval calculations (defaults to TRUE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 13.3 from "Applied Nonparametric Statistical Methods" (5th edition)
breslow.day(ch13$machine, ch13$output.status, ch13$material.source)
```

```
# Exercise 13.7 from "Applied Nonparametric Statistical Methods" (5th edition)
breslow.day(ch13$medicine, ch13$response, ch13$location)
```

bs	<i>Create bootstrap confidence interval</i>
----	---

Description

bs() creates a bootstrap confidence interval and is used in chapter 14 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
bs(x, y = NULL, CI.width = 0.95, nsims.bs = 10000, seed = NULL)
```

Arguments

x	Numeric vector
y	Numeric vector or NULL (defaults to NULL)
CI.width	Confidence interval width (defaults to 0.95)
nsims.bs	Number of bootstrap samples to be taken (defaults to 10000)
seed	Random number seed to be used for Monte Carlo simulations (defaults to NULL)

Value

A list object object with the results from applying the function

Examples

```
# Example 14.5 from "Applied Nonparametric Statistical Methods" (5th edition)
bs(ch14$example14.2, nsims.bs = 2000, CI.width = 0.95, seed = 1)
bs(ch14$example14.2, nsims.bs = 2000, CI.width = 0.99, seed = 1)
```

ch10	<i>Data used in Chapter 10</i>
------	--------------------------------

Description

Data used in Chapter 10 of "Applied Nonparametric Statistical Methods" (5th edition)

- q1 (used in section 10.1.2, examples 10.2, 10.5, 10.9)
- q2 (used in section 10.1.2, examples 10.2, 10.5, 10.9)
- death.year (used in examples 10.4, 10.8)
- diving.rank (used in example 10.10)
- competitors (used in example 10.10)

- judges (used in example 10.10)
- dentistA (used in example 10.11)
- dentistB (used in example 10.11)
- questionnaire (used in example 10.12, exercise 10.13)
- demonstration (used in example 10.12, exercise 10.13)
- gender (used in exercise 10.13)
- items (used in example 10.12)
- ERA (used in exercises 10.1, 10.3, 10.6, 10.7)
- ESMS (used in exercises 10.1, 10.3, 10.6)
- SSS (used in exercise 10.7)
- British (used in example 10.8, exercise 10.10)
- American (used in example 10.8, exercise 10.10)
- Canadian (used in example 10.9, exercise 10.10)
- Australian (used in example 10.9, exercise 10.10)
- design (used in exercise 10.10)
- country (used in exercise 10.10)
- marks (used in exercise 10.11)
- script (used in exercise 10.11)
- examiner (used in exercise 10.11)
- observerA (used in exercise 10.12)
- observerB (used in exercise 10.12)

Usage

`ch10`

Format

`ch10:`

A list with 26 data vectors

Source

"Applied Nonparametric Statistical Methods" (5th edition)

ch11

Data used in Chapter 11

Description

Data used in Chapter 11 of "Applied Nonparametric Statistical Methods" (5th edition)

- parentlimit (used in examples 11.2, 11.3, 11.4, 11.6)
- reportedtime (used in examples 11.2, 11.3, 11.4, 11.6)
- age (used in example 11.5)
- length (used in example 11.5)
- parentlimit.2 (used in example 11.7)
- reportedtime.2 (used in example 11.7)
- days.stored (used in exercise 11.3)
- rotten (used in exercise 11.3)
- ERA (used in exercise 11.6)
- ESMS (used in exercise 11.6)
- depth (used in exercise 11.8)
- ammonia (used in exercise 11.8)
- food.weight.A (used in exercise 11.9)
- weight.gain.A (used in exercise 11.9)
- food.weight.B (used in exercise 11.9)
- weight.gain.B (used in exercise 11.9)
- SW.England (used in exercise 11.10)
- N.Scotland (used in exercise 11.10)

Usage

ch11

Format

ch11:

A list with 18 data vectors

Source

"Applied Nonparametric Statistical Methods" (5th edition)

Description

Data used in Chapter 12 of "Applied Nonparametric Statistical Methods" (5th edition)

- feedback.freq (used in example 12.1)
- PPI.person (used in example 12.1)
- infection.site (used in examples 12.2, 12.3)
- district (used in examples 12.2, 12.3)
- drugYZ (used in example 12.4)
- side.effect (used in example 12.4)
- drugAB (used in example 12.5)
- side.effect.level (used in example 12.5)
- time.to.failure (used in example 12.6)
- cause (used in example 12.6)
- dose (used in examples 12.7, 12.8)
- dose.side.effect (used in example 12.7, 12.8)
- platelet.count (used in examples 12.9)
- spleen.size (used in example 12.9)
- last.digits (used in example 12.10)
- accidents (used in example 12.11)
- accidents.reduced (used in example 12.11)
- side.effect.new (used in example 12.12)
- side.effect.old (used in example 12.12)
- bronchitis (used in exercise 12.1)
- otitis.media (used in exercise 12.1)
- welsh.language (used in exercise 12.2)
- opportunities (used in exercise 12.2)
- diagnosis (used in exercise 12.3)
- position.played (used in exercise 12.3)
- PPI.person.2 (used in exercise 12.4)
- feedback.satisfaction (used in exercise 12.4)
- win.opinion (used in exercise 12.5)
- supporter (used in exercise 12.5)
- diabetes.status (used in exercise 12.6)
- ethnic.group (used in exercise 12.6)

- horse.wins (used in exercise 12.7)
- F1.wins (used in exercise 12.8)
- strokes (used in exercise 12.9)
- recurrent.visits (used in exercise 12.10)
- holes (used in exercise 12.11)
- first.response (used in exercise 12.12)
- second.response (used in exercise 12.12)

Usage

ch12

Format

ch12:

A list with 38 data vectors

Source

"Applied Nonparametric Statistical Methods" (5th edition)

ch13

Data used in Chapter 13

Description

Data used in Chapter 13 of "Applied Nonparametric Statistical Methods" (5th edition)

- physical.activity (used in examples 13.1, 13.2, exercise 13.2)
- tv.viewing (used in examples 13.1, 13.2, exercise 13.2)
- gender (used in examples 13.1, 13.2, exercise 13.2)
- machine (used in example 13.3)
- output.status (used in example 13.3)
- material.source (used in example 13.3)
- drug (used in example 13.4, section 13.2.5)
- side.effects (used in example 13.4, section 13.2.5)
- age.group (used in example 13.4, section 13.2.5)
- dose (used in examples 13.7, 13.8)
- dose.side.effect (used in examples 13.7, 13.8)
- alcohol (used in example 13.9)
- malformation (used in example 13.9)
- frequency (used in example 13.10)

- person (used in example 13.10)
- medicine (used in exercise 13.7, section 13.3.1)
- response (used in exercise 13.7, section 13.3.1)
- location (used in exercise 13.7, section 13.3.1)
- chemo.drug (used in example 13.12)
- chemo.side.effect (used in example 13.12)
- group (used in section 13.4)
- promoted (used in section 13.4)
- company (used in section 13.4)
- breakfast.eaten (used in exercise 13.3)
- VEL (used in exercise 13.3)
- boys.girls (used in exercise 13.3)
- cholesterol (used in exercise 13.4)
- SBP (used in exercise 13.4)
- schooling (used in exercise 13.5)
- abortion.attitude (used in exercise 13.5)
- PPI.ages (used in exercise 13.9)
- PPI.people (used in exercise 13.9)
- laid.off (used in exercises 13.10, 13.11)
- employee.ages (used in exercise 13.10)
- employee.ages.2 (used in exercise 13.11)

Usage

`ch13`

Format

`ch13:`

A list with 35 data vectors

Source

"Applied Nonparametric Statistical Methods" (5th edition)

ch14*Data used in Chapter 14*

Description

Data used in Chapter 14 of "Applied Nonparametric Statistical Methods" (5th edition)

- example14.2 (used in examples 14.2, 14.5)
- X14.4 (used in exercise 14.4)
- Y14.4 (used in exercise 14.4)

Usage

ch14

Format

ch14:

A list with 3 data vectors

Source

"Applied Nonparametric Statistical Methods" (5th edition)

ch15*Data used in Chapter 15*

Description

Data used in Chapter 15 of "Applied Nonparametric Statistical Methods" (5th edition)

- diet (used in section 15.3.5)
- BMI (used in section 15.3.1)
- wgt.VLCD (used in section 15.3.2)
- wgt.norm (used in section 15.3.2)
- opdiff (used in section 15.3.5)
- optime.VLCD (used in sections 15.3.3, 15.3.6)
- optime.norm (used in sections 15.3.3, 15.3.6)
- los.VLCD (used in section 15.3.6)
- los.norm (used in section 15.3.6)
- optime (used in section 15.3.4)
- los (used in section 15.3.4)

Usage

ch15

Format

ch15:

A list with 11 data vectors

Source

"Applied Nonparametric Statistical Methods" (5th edition)

ch3

*Data used in Chapter 3***Description**

Data used in Chapter 3 of "Applied Nonparametric Statistical Methods" (5th edition)

- sampleI (used in examples 3.1, 3.2, 3.3, exercise 3.17)
- sampleII (used in examples 3.1, 3.2, 3.3, exercise 3.17)
- heartrates1 (used in examples 3.4, 3.11)
- heartrates2 (used in examples 3.5, 3.6, 3.7)
- withties (used in example 3.8)
- tiedifrounded1 (used in example 3.8)
- tiedifrounded2 (used in example 3.8)
- ages (used in example 3.8, exercise 3.9)
- sampleA (used in example 3.12)
- sampleB (used in examples 3.12, 3.13)
- sampleA2 (used in example 3.12)
- sampleA3 (used in example 3.12)
- heartrates2a (used in example 3.14)
- heartrates2b (used in example 3.14)
- sampleIa (used in exercise 3.1)
- parkingtime (used in exercise 3.3)
- Svals (used in exercise 3.4)
- children (used in exercise 3.6)
- fishlengths (used in exercises 3.7, 3.11)
- sleeptime (used in exercise 3.10)
- weightloss (used in exercise 3.12)
- plants (used in exercise 3.13)
- birthprops (used in exercise 3.14)
- assembly (used in exercise 3.15)
- weightchange (used in exercise 3.16)

Usage

ch3

Format

ch3:

A list with 25 data vectors

Source

"Applied Nonparametric Statistical Methods" (5th edition)

ch4

*Data used in Chapter 4***Description**

Data used in Chapter 4 of "Applied Nonparametric Statistical Methods" (5th edition)

- breaks (used in example 4.2)
- ages (used in example 4.4)
- precipitation (used in example 4.13)
- tosses1 (used in example 4.14)
- tosses2 (used in example 4.14)
- tosses3 (used in example 4.14)
- births (used in example 4.15)
- times.as.degrees (used in example 4.16)
- dates.as.degrees (used in example 4.17)
- waiting.time (used in exercise 4.2)
- visiting.supporters (used in exercise 4.3)
- days.waiting (used in exercise 4.8)
- rainfall.by.latitude (used in exercise 4.9)
- points (used in exercise 4.10)
- rainfall.DRC (used in exercise 4.11)
- piped.water.DRC (used in exercise 4.12)
- accident.bearings (used in exercise 4.13)
- board.angles (used in exercise 4.14)
- arrow.angles (used in exercise 4.15)
- football.results (used in exercise 4.17)

Usage

ch4

Format

ch4:

A list with 20 data vectors

Source

"Applied Nonparametric Statistical Methods" (5th edition)

ch5*Data used in Chapter 5*

Description

Data used in Chapter 5 of "Applied Nonparametric Statistical Methods" (5th edition)

- LVF (used in example 5.1, exercise 6.2)
- RVF (used in example 5.1, exercise 6.2)
- arithmetic (used in example 5.2)
- bp (used in example 5.3)
- bp.incorrect (used in example 5.3)
- yr0910 (used in example 5.10)
- yr1314 (used in example 5.10)
- bp.diff (used in exercise 5.1)
- LabI (used in exercise 5.2)
- LabII (used in exercise 5.2)
- parent (used in exercise 5.4)
- online (used in exercise 5.5)
- lectures (used in exercise 5.5)
- additiveA (used in exercise 5.9)
- additiveB (used in exercise 5.9)
- round2 (used in exercise 5.10)
- round3 (used in exercise 5.10)
- pollA (used in exercise 5.11)
- pollB (used in exercise 5.11)
- kHz0.125 (used in exercise 5.12)
- kHz0.25 (used in exercise 5.12)

Usage

ch5

Format

ch5:

A list with 21 data vectors

Source

"Applied Nonparametric Statistical Methods" (5th edition)

ch6

*Data used in Chapter 6***Description**

Data used in Chapter 6 of "Applied Nonparametric Statistical Methods" (5th edition)

- groupA (used in examples 6.1, 6.2, 6.3, 6.10, 6.17)
- groupB (used in examples 6.1, 6.2, 6.3, 6.10, 6.17)
- groupA.sch2 (used in example 6.4)
- groupB.sch2 (used in example 6.4)
- groupA.sch2.grp (used in example 6.5)
- groupB.sch2.grp (used in example 6.5)
- males (used in examples 6.7, 6.8)
- females (used in examples 6.7, 6.8)
- sampleI (used in example 6.9)
- sampleII (used in example 6.9)
- typeA (used in examples 6.11, 6.12, 6.13, exercises 6.11, 6.12)
- typeB (used in examples 6.11, 6.12, 6.13, exercises 6.11, 6.12)
- groupI (used in example 6.14)
- groupII (used in example 6.14)
- groupI.trimmed (used in example 6.14)
- groupI.amended (used in example 6.14)
- salivaF (used in examples 6.15, 6.16)
- salivaM (used in examples 6.15, 6.16)
- sex (used in example 6.18)
- temp.H (used in exercise 6.1)
- temp.L (used in exercise 6.1)

- DMF.M (used in exercise 6.3)
- DMFF (used in exercise 6.3)
- weight.diabetic (used in exercise 6.4)
- weight.normal (used in exercise 6.4)
- cooling.time.standard (used in exercise 6.5)
- cooling.time.cheap (used in exercise 6.5)
- wait.1979 (used in exercise 6.6)
- wait.1983 (used in exercise 6.6)
- activity.boys (used in exercise 6.7)
- activity.girls (used in exercise 6.7)
- time.withoutLD (used in exercises 6.13, 6.14)
- time.withLD (used in exercises 6.13, 6.14)
- doseI (used in exercise 6.15)
- doseII (used in exercise 6.15)
- doseI.2 (used in exercise 6.15)
- travel (used in exercise 6.16)
- politics (used in exercise 6.16)
- twins (used in exercise 6.17)

Usage

ch6

Format

ch6:

A list with 39 data vectors

Source

"Applied Nonparametric Statistical Methods" (5th edition)

Description

Data used in Chapter 7 of "Applied Nonparametric Statistical Methods" (5th edition)

- affordability (used in example 7.1, exercise 7.16)
- regions (used in example 7.1, exercise 7.16)
- age (used in example 7.2)
- positions (used in example 7.2)
- dementia.age (used in examples 7.3, 7.9)
- features (used in examples 7.3, 7.9)
- time (used in examples 7.4, 7.5)
- surgeon (used in examples 7.4, 7.5)
- pulse (used in example 7.6)
- student (used in example 7.6)
- time.period (used in example 7.6)
- nodes (used in example 7.7)
- treatment (used in example 7.7)
- block (used in example 7.7)
- outcome (used in example 7.8)
- member (used in example 7.8)
- climb (used in example 7.8)
- procedure.time (used in example 7.10)
- team.member (used in example 7.10)
- sentences (used in exercise 7.2)
- author (used in exercise 7.2)
- head.width (used in exercise 7.4)
- species (used in exercise 7.4)
- braking.distance (used in exercise 7.5)
- speed (used in exercise 7.5)
- platelet.count (used in exercise 7.6)
- spleen.size (used in exercise 7.6)
- liver.weight (used in exercise 7.7)
- dose (used in exercise 7.7)
- house (used in exercise 7.7)
- mark (used in exercise 7.8)

- scheme (used in exercise 7.8)
- candidate (used in exercise 7.8)
- prem.contractions (used in exercise 7.9)
- drug (used in exercise 7.9)
- patient (used in exercise 7.9)
- births (used in exercise 7.11)
- week (used in exercise 7.11)
- weekday (used in exercise 7.11)
- names.recalled (used in exercise 7.12)
- group (used in exercise 7.12)
- medical.student (used in exercise 7.12)
- soc.media.use (used in exercise 7.14)
- participant (used in exercise 7.14)
- day (used in exercise 7.14)
- braking.distance.2 (used in exercise 7.15)
- initial.speed (used in exercise 7.15)

Usage

ch7

Format

ch7:

A list with 47 data vectors

Source

"Applied Nonparametric Statistical Methods" (5th edition)

ch8

Data used in Chapter 8

Description

Data used in Chapter 8 of "Applied Nonparametric Statistical Methods" (5th edition)

- plant.weight (used in example 8.2)
- growth.hormone (used in examples 8.6, 8.7)
- undersoil.heating (used in examples 8.6, 8.7)
- plant.weight.2 (used in example 8.6)
- plant.weight.3 (used in examples 8.4, 8.5)

- plant.weight.4 (used in example 8.7)
- sequence (used in example 8.9)
- periodI (used in example 8.9)
- periodII (used in example 8.9)
- sentences (used in example 8.10)
- authors (used in example 8.10)
- prey.preference (used in example 8.11)
- prey (used in example 8.11)
- larva (used in example 8.11)
- game.time (used in exercise 8.3)
- experience (used in exercise 8.3)
- game (used in exercise 8.3)
- periodI.mistakes.AB (used in exercise 8.6)
- periodII.mistakes.AB (used in exercise 8.6)
- periodI.mistakes.BA (used in exercise 8.6)
- periodII.mistakes.BA (used in exercise 8.6)
- periodI.time.AB (used in exercise 8.7)
- periodII.time.AB (used in exercise 8.7)
- periodI.time.BA (used in exercise 8.7)
- periodII.time.BA (used in exercise 8.7)
- seizure.score (used in exercises 8.8, 8.9)
- hospital (used in exercises 8.8, 8.9)
- silver.content (used in exercise 8.10)
- dynasty (used in exercise 8.10)

Usage

ch8

Format

ch8:

A list with 29 data vectors

Source

"Applied Nonparametric Statistical Methods" (5th edition)

ch9

Data used in Chapter 9

Description

Data used in Chapter 9 of "Applied Nonparametric Statistical Methods" (5th edition)

- symp.survtime (used in examples 9.1, 9.3)
- symp.censor (used in examples 9.1, 9.3)
- asymp.survtime (used in examples 9.1, 9.3)
- asymp.censor (used in examples 9.1, 9.3)
- sampleI.survtime (used in following example 9.3, example 9.4)
- sampleI.censor (used in example 9.4)
- sampleII.survtime (used in example 9.4)
- sampleII.survtime.2 (used in following example 9.3)
- sampleII.censor (used in example 9.4)
- samplesAB.survtime (used in example 9.6)
- samplesAB.censor (used in example 9.6)
- samplesAB (used in example 9.6)
- samplesXYZ.survtime (used in example 9.7)
- samplesXYZ.censor (used in example 9.7)
- samplesXYZ (used in example 9.7)
- boys.toothtime (used in exercise 9.2)
- girls.toothtime (used in exercise 9.2)
- regimeA.survtime (used in exercises 9.5, 9.6)
- regimeA.censor (used in exercises 9.5, 9.6)
- regimeB.survtime (used in exercises 9.5, 9.6)
- regimeB.censor (used in exercises 9.5, 9.6)
- bulbA (used in exercise 9.8)
- bulbB (used in exercise 9.8)

Usage

ch9

Format

ch9:

A list with 23 data vectors

Source

"Applied Nonparametric Statistical Methods" (5th edition)

<code>chisqtest.ANSM</code>	<i>Perform Chi-squared test</i>
-----------------------------	---------------------------------

Description

`chisqtest.ANSM()` is a wrapper for `chisq.test()` from the `stats` package - performs the Chi-squared test and is used in chapters 12 and 13 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
chisqtest.ANSM(
  x,
  y = NULL,
  p = NULL,
  cont.corr = TRUE,
  max.exact.cases = 10,
  nsims.mc = 1e+05,
  seed = NULL,
  do.exact = TRUE,
  do.asymp = FALSE,
  do.mc = FALSE
)
```

Arguments

<code>x</code>	Factor of same length as <code>y</code> , or table
<code>y</code>	Factor of same length as <code>x</code> (or <code>NULL</code> if <code>x</code> is table) (defaults to <code>NULL</code>)
<code>p</code>	Vector of probabilities (expressed as numbers between 0 and 1 and summing to 1) of same length as <code>x</code> or <code>NULL</code> (defaults to <code>NULL</code>)
<code>cont.corr</code>	Boolean indicating whether or not to use continuity correction (defaults to <code>TRUE</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to 10)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 100000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to <code>NULL</code>)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to <code>TRUE</code>)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to <code>FALSE</code>)
<code>do.mc</code>	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to <code>FALSE</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 12.1 from "Applied Nonparametric Statistical Methods" (5th edition)
chisqtest.ANSM(ch12$feedback.freq, ch12$PPI.person, do.exact = FALSE, do.asymp = TRUE)

# Exercise 13.7 from "Applied Nonparametric Statistical Methods" (5th edition)
chisqtest.ANSM(ch13$medicine[ch13$location == "Rural"],
               ch13$response[ch13$location == "Rural"], seed = 1)
```

cochran.q

Perform Cochran Q test

Description

`cochran.q()` performs the Cochran Q test and is used in chapter 7 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
cochran.q(
  y,
  groups,
  blocks,
  max.exact.perms = 1e+05,
  nsims.mc = 1e+05,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE
)
```

Arguments

<code>y</code>	Binary vector of same length as <code>groups</code> , <code>blocks</code>
<code>groups</code>	Factor of same length as <code>y</code> , <code>blocks</code> with levels such that <code>length(y) == nlevels(groups)</code> * <code>nlevels(blocks)</code>
<code>blocks</code>	Factor of same length as <code>y</code> , <code>groups</code> with levels such that <code>length(y) == nlevels(groups)</code> * <code>nlevels(blocks)</code>
<code>max.exact.perms</code>	Maximum number of permutations allowed for exact calculations (defaults to 100000)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 100000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to NULL)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 7.8 from "Applied Nonparametric Statistical Methods" (5th edition)
cochran.q(ch7$outcome, ch7$climb, ch7$member, do.exact = FALSE, do.asymp = TRUE)

# Exercise 7.14 from "Applied Nonparametric Statistical Methods" (5th edition)
cochran.q(ch7$soc.media.use, ch7$participant, ch7$day, do.exact = FALSE, do.asymp = TRUE)
```

cohen.kappa

*Calculate Cohen's kappa***Description**

cohen.kappa() calculates Cohen's kappa and is used in chapter 10 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
cohen.kappa(
  y1,
  y2,
  blocks = NULL,
  alternative = c("two.sided", "less", "greater"),
  CI.width = 0.95,
  max.exact.cases = 10,
  nsims.mc = 1e+05,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE,
  do.CI = FALSE,
  do.mc = FALSE
)
```

Arguments

y1	Factor of same length as y2, blocks and same levels as y2 and (if blocks not NULL) with 2 levels
y2	Factor of same length as y1, blocks and same levels as y1 and (if blocks not NULL) with 2 levels
blocks	Factor of same length as y1, y2 or NULL (defaults to NULL)
alternative	Type of alternative hypothesis (defaults to two.sided)
CI.width	Confidence interval width (defaults to 0.95)

max.exact.cases	Maximum number of cases allowed for exact calculations (defaults to 10)
nsims.mc	Number of Monte Carlo simulations to be performed (defaults to 100000)
seed	Random number seed to be used for Monte Carlo simulations (defaults to NULL)
do.asymp	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
do.exact	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)
do.CI	Boolean indicating whether or not to perform confidence interval calculations (defaults to FALSE)
do.mc	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to FALSE)

Value

An ANSMstat object with the results from applying the function

Examples

```
# Example 10.11 from "Applied Nonparametric Statistical Methods" (5th edition)
cohen.kappa(ch10$dentistA, ch10$dentistB, do.asymp = TRUE, do.exact = FALSE,
            alternative = "greater")

# Example 10.12 from "Applied Nonparametric Statistical Methods" (5th edition)
cohen.kappa(ch10$questionnaire, ch10$demonstration, ch10$items)
```

conover

Perform Conover test using standard or squared ranks

Description

conover() performs the Conover test using standard or squared ranks and is used in chapters 6 and 7 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
conover(
  x,
  y,
  H0 = NULL,
  alternative = c("two.sided", "less", "greater"),
  abs.ranks = FALSE,
  max.exact.perms = 5e+06,
  nsims.mc = 10000,
  seed = NULL,
```

```

do.asymp = FALSE,
do.exact = TRUE,
do.mc = FALSE
)

```

Arguments

x	Numeric vector of same length as y
y	Factor of same length as x
H0	Null hypothesis value (defaults to NULL)
alternative	Type of alternative hypothesis (defaults to two.sided)
abs.ranks	Boolean indicating whether absolute ranks to be used instead of squared ranks (defaults to FALSE)
max.exact.perms	Maximum number of permutations allowed for exact calculations (defaults to 5000000)
nsims.mc	Number of Monte Carlo simulations to be performed (defaults to 10000)
seed	Random number seed to be used for Monte Carlo simulations (defaults to NULL)
do.asymp	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
do.exact	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)
do.mc	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to FALSE)

Value

An ANSMtest object with the results from applying the function

Examples

```

# Example 6.13 from "Applied Nonparametric Statistical Methods" (5th edition)
conover(ch6$typeA, ch6$typeB, do.exact = FALSE, do.asymp = TRUE)

# Exercise 7.15 from "Applied Nonparametric Statistical Methods" (5th edition)
conover(ch7$braking.distance.2, ch7$initial.speed, do.exact = FALSE, do.asymp = TRUE)

```

control.median *Perform Control median test*

Description

`control.median()` performs the Control median test and is used in chapters 6 and 9 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
control.median(
  x,
  y,
  H0 = NULL,
  alternative = c("two.sided", "less", "greater"),
  CI.width = 0.95,
  max.exact.cases = 1000,
  nsims.mc = 10000,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE,
  do.CI = TRUE
)
```

Arguments

<code>x</code>	Numeric vector
<code>y</code>	Numeric vector
<code>H0</code>	Null hypothesis value (defaults to <code>NULL</code>)
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>CI.width</code>	Confidence interval width (defaults to <code>0.95</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to <code>1000</code>)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to <code>10000</code>)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to <code>NULL</code>)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to <code>FALSE</code>)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to <code>TRUE</code>)
<code>do.CI</code>	Boolean indicating whether or not to perform confidence interval calculations (defaults to <code>TRUE</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 6.9 from "Applied Nonparametric Statistical Methods" (5th edition)
control.median(ch6$sampleI, ch6$sampleII, alternative = "greater")

# Exercise 9.8 from "Applied Nonparametric Statistical Methods" (5th edition)
control.median(ch9$bulbA, ch9$bulbB, alternative = "greater", nsims = 1000)
```

cox.stuart

Perform Cox-Stuart test

Description

`cox.stuart()` performs the Cox-Stuart test and is used in chapters 4 and 10 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
cox.stuart(
  x,
  alternative = c("two.sided", "less", "greater"),
  cont.corr = TRUE,
  max.exact.cases = 1e+07,
  do.asymp = FALSE,
  do.exact = TRUE
)
```

Arguments

<code>x</code>	Numeric vector
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>cont.corr</code>	Boolean indicating whether or not to use continuity correction (defaults to <code>TRUE</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to <code>10000000</code>)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to <code>FALSE</code>)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to <code>TRUE</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 4.13 from "Applied Nonparametric Statistical Methods" (5th edition)
cox.stuart(ch4$precipitation)

# Exercise 10.5 from "Applied Nonparametric Statistical Methods" (5th edition)
cox.stuart(app1$McDelta[order(ch10$death.year)], alternative = "less")
```

cramer.von.mises *Perform Cramer-von Mises test*

Description

cramer.von.mises() performs the Cramer-von Mises test and is used in chapter 6 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
cramer.von.mises(x, y, alternative = c("two.sided", "less", "greater"))
```

Arguments

x	Numeric vector
y	Numeric vector
alternative	Type of alternative hypothesis (defaults to two.sided)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 6.16 from "Applied Nonparametric Statistical Methods" (5th edition)
cramer.von.mises(ch6$salivaF, ch6$salivaM)
cramer.von.mises(ch6$salivaF, ch6$salivaM, alternative = "greater")
```

<code>fishertest.ANSM</code>	<i>Perform Fisher exact test</i>
------------------------------	----------------------------------

Description

`fishertest.ANSM()` is a wrapper for `fisher.test()` from the `stats` package - performs the Fisher exact test and is used in chapters 6, 12 and 13 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
fishertest.ANSM(
  x,
  y,
  H0 = NULL,
  alternative = c("two.sided", "less", "greater"),
  max.exact.cases = 10000,
  do.exact = TRUE
)
```

Arguments

<code>x</code>	Numeric vector or factor
<code>y</code>	Numeric vector or factor
<code>H0</code>	Null hypothesis value (defaults to <code>NULL</code>)
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to 10000)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to <code>TRUE</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 6.7 from "Applied Nonparametric Statistical Methods" (5th edition)
fishertest.ANSM(ch6$males, ch6$females)

# Exercise 13.10 from "Applied Nonparametric Statistical Methods" (5th edition)
fishertest.ANSM(ch13$laid.off, ch13$employee.ages)
```

<code>friedman</code>	<i>Perform Friedman test</i>
-----------------------	------------------------------

Description

`friedman()` performs the Friedman test and is used in chapter 7 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
 friedman(
  y,
  groups,
  blocks,
  use.Iman.Davenport = FALSE,
  max.exact.perms = 1e+05,
  nsims.mc = 1e+05,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE
)
```

Arguments

<code>y</code>	Numeric vector of same length as groups, blocks
<code>groups</code>	Factor of same length as <code>y</code> , blocks with levels such that <code>length(y) == nlevels(groups)</code> * <code>nlevels(blocks)</code>
<code>blocks</code>	Factor of same length as <code>y</code> , groups with levels such that <code>length(y) == nlevels(groups)</code> * <code>nlevels(blocks)</code>
<code>use.Iman.Davenport</code>	Boolean indicating whether or not to use Iman and Davenport approximation (defaults to FALSE)
<code>max.exact.perms</code>	Maximum number of permutations allowed for exact calculations (defaults to 100000)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 100000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to NULL)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 7.6 from "Applied Nonparametric Statistical Methods" (5th edition)
friedman(ch7$pulse, ch7$time.period, ch7$student, do.exact = FALSE, do.asymp = TRUE)

# Exercise 7.12 from "Applied Nonparametric Statistical Methods" (5th edition)
friedman(ch7$names.recalled, ch7$group, ch7$medical.student, use.Iman.Davenport = TRUE,
do.exact = FALSE, do.asymp = TRUE)
```

friedman.lsd

Perform Least Significant Differences test after the Friedman test

Description

`friedman.lsd()` performs the Least Significant Differences test after the Friedman test and is used in chapter 8 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
 friedman.lsd(y, groups, blocks, ids)
```

Arguments

<code>y</code>	Numeric vector of same length as <code>groups</code> , <code>blocks</code>
<code>groups</code>	Factor of same length as <code>y</code> , <code>blocks</code> with levels such that <code>length(y) == nlevels(groups)</code> * <code>nlevels(blocks)</code>
<code>blocks</code>	Factor of same length as <code>y</code> , <code>groups</code> with levels such that <code>length(y) == nlevels(groups)</code> * <code>nlevels(blocks)</code>
<code>ids</code>	Vector of length 2 with elements both levels of <code>groups</code>

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 8.11 from "Applied Nonparametric Statistical Methods" (5th edition)
friedman.lsd(ch8$prey.preference, ch8$prey, ch8$larva, c("Cyclops", "Anopheles"))

# from "Applied Nonparametric Statistical Methods" (5th edition)
```

<code>gehan.wilcoxon</code>	<i>Perform Gehan-Wilcoxon test</i>
-----------------------------	------------------------------------

Description

`gehan.wilcoxon()` performs the Gehan-Wilcoxon test and is used in chapter 9 of "Applied Non-parametric Statistical Methods" (5th edition)

Usage

```
gehan.wilcoxon(
  x,
  y,
  x.c,
  y.c,
  alternative = c("two.sided", "less", "greater"),
  max.exact.perms = 1e+05,
  nsims.mc = 1e+05,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE
)
```

Arguments

<code>x</code>	Numeric vector of same length as <code>y</code> , <code>x.c</code> , <code>y.c</code>
<code>y</code>	Numeric vector of same length as <code>x</code> , <code>x.c</code> , <code>y.c</code>
<code>x.c</code>	Binary vector of same length as <code>x</code> , <code>y</code> , <code>x.c</code>
<code>y.c</code>	Binary vector of same length as <code>x</code> , <code>y</code> , <code>y.c</code>
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>max.exact.perms</code>	Maximum number of permutations allowed for exact calculations (defaults to 100000)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 100000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to NULL)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 9.1 from "Applied Nonparametric Statistical Methods" (5th edition)
gehan.wilcoxon(ch9$symp.survtime, ch9$asymp.survtime,
  ch9$symp.censor, ch9$asymp.censor, alternative = "less",
  do.exact = FALSE, do.asymp = TRUE)

# Exercise 9.5 from "Applied Nonparametric Statistical Methods" (5th edition)
gehan.wilcoxon(ch9$regimeA.survtime, ch9$regimeB.survtime,
  ch9$regimeA.censor, ch9$regimeB.censor, do.exact = FALSE, do.asymp = TRUE)
```

`hettmansperger.elmore` *Perform Hettmansperger and Elmore interaction test*

Description

`hettmansperger.elmore()` performs the Hettmansperger and Elmore interaction test and is used in chapter 8 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
hettmansperger.elmore(
  y,
  factor.a,
  factor.b,
  nsims.mc = 1000,
  seed = NULL,
  do.asymp = TRUE,
  do.mc = FALSE,
  median.polish = FALSE
)
```

Arguments

<code>y</code>	Numeric vector of same length as <code>factor.a</code> , <code>factor.b</code>
<code>factor.a</code>	Factor of same length as <code>y</code> , <code>factor.b</code>
<code>factor.b</code>	Factor of same length as <code>y</code> , <code>factor.a</code>
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 1000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to NULL)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to TRUE)
<code>do.mc</code>	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to FALSE)
<code>median.polish</code>	Boolean indicating whether or not to use median polish (defaults to FALSE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 8.6 from "Applied Nonparametric Statistical Methods" (5th edition)
hettmansperger.elmore(ch8$plant.weight.2, ch8$growth.hormone, ch8$undersoil.heating)

# Exercise 8.3 from "Applied Nonparametric Statistical Methods" (5th edition)
hettmansperger.elmore(ch8$game.time, ch8$experience, ch8$game)
```

hodges.ajne

Perform Hodges-Ajne test

Description

`hodges.ajne()` performs the Hodges-Ajne test and is used in chapter 4 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
hodges.ajne(x, alternative = c("two.sided"), minx = 0, maxx = 360)
```

Arguments

<code>x</code>	Numeric vector
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>c("two.sided")</code>)
<code>minx</code>	Minimum value for <code>x</code> (defaults to <code>0</code>)
<code>maxx</code>	Maximum value for <code>x</code> (defaults to <code>360</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 4.16 from "Applied Nonparametric Statistical Methods" (5th edition)
hodges.ajne(ch4$times.as.degrees)

# Exercise 4.14 from "Applied Nonparametric Statistical Methods" (5th edition)
hodges.ajne(ch4$board.angles)
```

`jonckheere.terpstra` *Perform Jonckheere-Terpstra test*

Description

`jonckheere.terpstra()` performs the Jonckheere-Terpstra test and is used in chapters 7, 8 and 12 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
jonckheere.terpstra(
  x,
  g,
  alternative = c("less", "greater"),
  max.exact.cases = 15,
  nsims.mc = 10000,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE,
  do.mc = FALSE,
  do.asymp.ties.adjust = TRUE
)
```

Arguments

<code>x</code>	Numeric vector or factor of same length as <code>g</code>
<code>g</code>	Factor of same length as <code>x</code>
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>c("less", "greater")</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to 15)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 10000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to <code>NULL</code>)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to <code>FALSE</code>)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to <code>TRUE</code>)
<code>do.mc</code>	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to <code>FALSE</code>)
<code>do.asymp.ties.adjust</code>	Boolean indicating whether or not to use adjustment for ties in data (defaults to <code>TRUE</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 7.3 from "Applied Nonparametric Statistical Methods" (5th edition)
jonckheere.terpstra(ch7$dementia.age, ch7$features, alternative = "greater",
do.exact = FALSE, do.asymp = TRUE, do.asymp.ties.adjust = FALSE)

# Exercise 12.6 from "Applied Nonparametric Statistical Methods" (5th edition)
jonckheere.terpstra(ch12$ethnic.group, ch12$diabetes.status, do.exact = FALSE, do.asymp = TRUE)
```

kendall.concordance *Calculate Kendall's concordance*

Description

`kendall.concordance()` calculates Kendall's concordance and is used in chapter 10 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
kendall.concordance(
  y,
  groups,
  blocks,
  max.exact.perms = 1e+05,
  nsims.mc = 1e+05,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE
)
```

Arguments

<code>y</code>	Numeric vector of same length as groups, blocks
<code>groups</code>	Factor of same length as <code>y</code> , blocks with levels such that <code>length(y) == nlevels(groups)</code> * <code>nlevels(blocks)</code>
<code>blocks</code>	Factor of same length as <code>y</code> , groups with levels such that <code>length(y) == nlevels(groups)</code> * <code>nlevels(blocks)</code>
<code>max.exact.perms</code>	Maximum number of permutations allowed for exact calculations (defaults to 100000)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 10000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to <code>NULL</code>)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)

Value

An ANSMstat object with the results from applying the function

Examples

```
# Exercise 10.11 from "Applied Nonparametric Statistical Methods" (5th edition)
kendall.concordance(ch10$marks, ch10$script, do.exact = FALSE, do.asymp = TRUE)
kendall.concordance(ch10$marks, ch10$examiner, ch10$script, do.exact = FALSE, do.asymp = TRUE)
```

kendall.tau

*Perform Kendall's tau***Description**

`kendall.tau()` performs the Kendall's tau and is used in chapter 10 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
kendall.tau(
  x,
  y,
  alternative = c("two.sided", "less", "greater"),
  max.exact.cases = 10,
  nsims.mc = 1e+05,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE,
  do.mc = FALSE
)
```

Arguments

<code>x</code>	Numeric vector of same length as <code>y</code>
<code>y</code>	Numeric vector of same length as <code>x</code>
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to 10)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 100000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to <code>NULL</code>)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to <code>FALSE</code>)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to <code>TRUE</code>)
<code>do.mc</code>	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to <code>FALSE</code>)

Value

An ANSMstat object with the results from applying the function

Examples

```
# Example 10.8 from "Applied Nonparametric Statistical Methods" (5th edition)
kendall.tau(ch10$death.year, app1$McDelta, alternative = "greater",
do.asymp = TRUE, do.exact = FALSE)

# Example 10.9 from "Applied Nonparametric Statistical Methods" (5th edition)
kendall.tau(ch10$Canadian, ch10$Australian)
```

kruskal.wallis	<i>Perform Kruskal-Wallis test</i>
----------------	------------------------------------

Description

`kruskal.wallis()` performs the Kruskal-Wallis test and is used in chapters 7 and 12 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
kruskal.wallis(
  x,
  g,
  max.exact.cases = 15,
  nsims.mc = 10000,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE,
  do.mc = FALSE
)
```

Arguments

<code>x</code>	Numeric vector or factor of same length as <code>g</code>
<code>g</code>	Factor of same length as <code>x</code>
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to 15)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 10000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to NULL)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)
<code>do.mc</code>	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to FALSE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 7.1 from "Applied Nonparametric Statistical Methods" (5th edition)
kruskal.wallis(ch7$affordability, ch7$regions, do.exact = FALSE, do.asymp = TRUE)

# Exercise 7.16 from "Applied Nonparametric Statistical Methods" (5th edition)
kruskal.wallis(ch7$affordability, ch7$regions)
```

kruskal.wallis.lsd

Perform Least Significant Differences test after the Kruskal-Wallis test

Description

kruskal.wallis.lsd() performs the Least Significant Differences test after the Kruskal-Wallis test and is used in chapter 8 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
kruskal.wallis.lsd(x, g, ids)
```

Arguments

x	Numeric vector of same length as g
g	Factor of same length as x
ids	Vector of length 2 with elements both levels of g

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 8.10 from "Applied Nonparametric Statistical Methods" (5th edition)
kruskal.wallis.lsd(ch8$sentences, ch8$authors, c("Vulliamy", "Queen"))

# Exercise 8.8 from "Applied Nonparametric Statistical Methods" (5th edition)
kruskal.wallis.lsd(ch8$seizure.score, ch8$hospital, c("HospitalA", "HospitalC"))
```

`kruskal.wallis.vdW` *Perform Kruskal-Wallis test with van der Waerden scores*

Description

`kruskal.wallis.vdW()` performs the Kruskal-Wallis test with van der Waerden scores and is used in chapter 7 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
kruskal.wallis.vdW(
  x,
  g,
  max.exact.cases = 15,
  nsims.mc = 10000,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE
)
```

Arguments

<code>x</code>	Numeric vector of same length as <code>g</code>
<code>g</code>	Factor of same length as <code>x</code>
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to 15)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 10000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to <code>NULL</code>)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to <code>FALSE</code>)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to <code>TRUE</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 7.2 from "Applied Nonparametric Statistical Methods" (5th edition)
kruskal.wallis.vdW(ch7$age, ch7$positions)
kruskal.wallis.vdW(ch7$age, ch7$positions, do.exact = FALSE, do.asymp = TRUE)
```

kstest.ANSMPerform Smirnov test and Kolgomorov test

Description

`kstest.ANSM()` is a wrapper for `ks.test()` from the `stats` package - performs the Smirnov test and Kolgomorov test and is used in chapters 4, 6 and 9 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
kstest.ANSM(
  x,
  y,
  ...,
  alternative = c("two.sided", "less", "greater"),
  max.exact.cases = 1000,
  do.asymp = FALSE,
  do.exact = TRUE
)
```

Arguments

<code>x</code>	Numeric vector
<code>y</code>	Numeric vector or a character string naming a cumulative distribution function or an actual cumulative distribution function
<code>...</code>	For the default method of <code>ks.test</code> , parameters of the distribution specified (as a character string) by <code>y</code> . Otherwise, further arguments to be passed to or from methods
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to 1000)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Exercise 4.3 from "Applied Nonparametric Statistical Methods" (5th edition)
kstest.ANSM(ch4$visiting.supporters, "pexp", rate = 2600)

# Exercise 9.2 from "Applied Nonparametric Statistical Methods" (5th edition)
kstest.ANSM(ch9$boys.toothtime, ch9$girls.toothtime)
```

lik.ratio

Perform Likelihood ratio test

Description

`lik.ratio()` performs the Likelihood ratio test and is used in chapters 12 and 13 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
lik.ratio(
  x,
  y,
  max.exact.cases = 10,
  nsims.mc = 1e+05,
  seed = NULL,
  do.exact = TRUE,
  do.asymp = FALSE,
  do.mc = FALSE
)
```

Arguments

<code>x</code>	Factor of same length as <code>y</code>
<code>y</code>	Factor of same length as <code>x</code>
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to 10)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 100000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to <code>NULL</code>)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to <code>TRUE</code>)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to <code>FALSE</code>)
<code>do.mc</code>	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to <code>FALSE</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 12.2 from "Applied Nonparametric Statistical Methods" (5th edition)
lik.ratio(ch12$infection.site, ch12$district, do.exact = FALSE, do.asymp = TRUE)

# Example 13.12 from "Applied Nonparametric Statistical Methods" (5th edition)
chemo.side.effect.3 <- ch13$chemo.side.effect
levels(chemo.side.effect.3) <- list("Side-effect" = c("Hair loss",
  "Visual impairment", "Hair loss & Visual impairment"), "None" = "None")
lik.ratio(ch13$chemo.drug, chemo.side.effect.3, seed = 1)
```

lilliefors*Performs Lilliefors test of Normality***Description**

`lilliefors()` performs Lilliefors test of Normality and is used in chapters 4, 5 and 6 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
lilliefors(x, alternative = c("two.sided"), nsims.mc = 10000, seed = NULL)
```

Arguments

<code>x</code>	Numeric vector
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>c("two.sided")</code>)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 10000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to <code>NULL</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 4.4 from "Applied Nonparametric Statistical Methods" (5th edition)
lilliefors(ch4$ages, seed = 1)

# Exercise 6.15 from "Applied Nonparametric Statistical Methods" (5th edition)
lilliefors(ch6$doseI.2, seed = 1, nsims = 1000)
```

linear.by.linear *Perform Linear by linear association test*

Description

`linear.by.linear()` performs the Linear by linear association test and is used in chapter 13 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
linear.by.linear(
  x,
  y,
  u = NULL,
  v = NULL,
  nsims.mc = 1e+05,
  seed = NULL,
  do.asymp = FALSE,
  do.mc = TRUE
)
```

Arguments

<code>x</code>	Factor of same length as <code>y</code>
<code>y</code>	Factor of same length as <code>x</code>
<code>u</code>	Numeric vector of length equal to number of levels of <code>x</code> or <code>NULL</code> (defaults to <code>NULL</code>)
<code>v</code>	Numeric vector of length equal to number of levels of <code>y</code> or <code>NULL</code> (defaults to <code>NULL</code>)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to <code>100000</code>)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to <code>NULL</code>)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to <code>FALSE</code>)
<code>do.mc</code>	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to <code>TRUE</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 13.8 from "Applied Nonparametric Statistical Methods" (5th edition)
linear.by.linear(ch13$dose, ch13$dose.side.effect, do.mc = FALSE, do.asymp = TRUE)
```

```
# Exercise 13.4 from "Applied Nonparametric Statistical Methods" (5th edition)
linear.by.linear(ch13$SBP, ch13$cholesterol, seed = 1)
```

logoddsratio.2x2 *Perform Log odds ratio test*

Description

`logoddsratio.2x2()` performs the Log odds ratio test and is used in chapter 13 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
logoddsratio.2x2(
  x,
  y,
  max.exact.cases = 10,
  nsims.mc = 1e+05,
  seed = NULL,
  do.exact = TRUE,
  do.asymp = FALSE,
  do.mc = FALSE
)
```

Arguments

<code>x</code>	Binary factor of same length as <code>y</code>
<code>y</code>	Binary factor of same length as <code>x</code>
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to 10)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 100000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to <code>NULL</code>)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to <code>TRUE</code>)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to <code>FALSE</code>)
<code>do.mc</code>	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to <code>FALSE</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Exercise 13.2 from "Applied Nonparametric Statistical Methods" (5th edition)
#logoddsratio.2x2(ch13$physical.activity[ch13$gender == "Boy"],
# ch13$tv.viewing[ch13$gender == "Boy"], do.exact = FALSE, do.asymp = TRUE)
#logoddsratio.2x2(ch13$physical.activity[ch13$gender == "Girl"],
# ch13$tv.viewing[ch13$gender == "Girl"], do.exact = FALSE, do.asymp = TRUE)
```

logrank

Perform logrank test

Description

`logrank()` performs the logrank test and is used in chapter 9 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
logrank(
  x,
  censored,
  groups,
  score.censored = TRUE,
  max.exact.perms = 1e+05,
  nsims.mc = 10000,
  seed = NULL
)
```

Arguments

<code>x</code>	Numeric vector of same length as <code>censored</code> , <code>groups</code>
<code>censored</code>	Binary vector of same length as <code>x</code> , <code>groups</code>
<code>groups</code>	Factor of same length as <code>x</code> , <code>censored</code>
<code>score.censored</code>	Boolean indicating whether or not to score censored values (defaults to TRUE)
<code>max.exact.perms</code>	Maximum number of permutations allowed for exact calculations (defaults to 100000)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 10000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to NULL)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 9.6 from "Applied Nonparametric Statistical Methods" (5th edition)
logrank(ch9$samplesAB.survtime, ch9$samplesAB.censor, ch9$samplesAB, score.censored = FALSE)

# Exercise 9.7 from "Applied Nonparametric Statistical Methods" (5th edition)
logrank(ch9$samplesXYZ.survtime, ch9$samplesXYZ.censor, ch9$samplesXYZ)
```

mantel.haenszel *Perform Mantel-Haenszel test*

Description

`mantel.haenszel()` performs the Mantel-Haenszel test and is used in chapter 13 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
mantel.haenszel(x, y, z, do.asymp = TRUE)
```

Arguments

x	Binary factor of same length as y, z
y	Binary factor of same length as x, z
z	Factor of same length as x, y
do.asymp	Boolean indicating whether or not to perform asymptotic calculations (defaults to TRUE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 13.4 from "Applied Nonparametric Statistical Methods" (5th edition)
mantel.haenszel(ch13$drug, ch13$side.effects, ch13$age.group)

# from "Applied Nonparametric Statistical Methods" (5th edition)
```

med.test*Perform Median test*

Description

`med.test()` performs the Median test and is used in chapters 6 and 7 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
med.test(
  x,
  y,
  H0 = NULL,
  alternative = c("two.sided", "less", "greater"),
  CI.width = 0.95,
  max.exact.cases = 1000,
  do.asymp = FALSE,
  do.exact = TRUE,
  do.CI = TRUE
)
```

Arguments

<code>x</code>	Numeric vector of same length as <code>y</code>
<code>y</code>	Numeric vector, or factor of same length as <code>x</code>
<code>H0</code>	Null hypothesis value (defaults to <code>NULL</code>)
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>CI.width</code>	Confidence interval width (defaults to <code>0.95</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to <code>1000</code>)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to <code>FALSE</code>)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to <code>TRUE</code>)
<code>do.CI</code>	Boolean indicating whether or not to perform confidence interval calculations (defaults to <code>TRUE</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 6.7 from "Applied Nonparametric Statistical Methods" (5th edition)
med.test(ch6$males, ch6$females)

# Example 7.5 from "Applied Nonparametric Statistical Methods" (5th edition)
med.test(ch7$time, ch7$surgeon, do.exact = FALSE, do.asymp = TRUE)
```

mood	<i>Perform Mood test</i>
------	--------------------------

Description

`mood()` performs the Mood test and is used in chapter 6 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
mood(
  x,
  y,
  H0 = NULL,
  alternative = c("two.sided", "less", "greater"),
  max.exact.cases = 25,
  do.asymp = FALSE,
  do.exact = TRUE
)
```

Arguments

<code>x</code>	Numeric vector
<code>y</code>	Numeric vector
<code>H0</code>	Null hypothesis value (defaults to <code>NULL</code>)
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to 25)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to <code>FALSE</code>)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to <code>TRUE</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 6.12 from "Applied Nonparametric Statistical Methods" (5th edition)
mood(ch6$typeA, ch6$typeB)
mood(ch6$typeA, ch6$typeB, do.exact = FALSE, do.asymp = TRUE)
```

`moses.extreme.reactions`

Perform Moses test for extreme reactions

Description

`moses.extreme.reactions()` performs the Moses test for extreme reactions and is used in chapter 6 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
moses.extreme.reactions(
  x,
  y,
  H0 = NULL,
  max.exact.cases = 1000,
  do.exact = TRUE
)
```

Arguments

<code>x</code>	Numeric vector
<code>y</code>	Numeric vector
<code>H0</code>	Null hypothesis value (defaults to <code>NULL</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to 1000)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to <code>TRUE</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 6.14 from "Applied Nonparametric Statistical Methods" (5th edition)
moses.extreme.reactions(ch6$groupI.amended, ch6$groupII)
moses.extreme.reactions(ch6$groupI.amended, ch6$groupII)
```

noether*Calculate Noether approximation*

Description

`noether()` calculates the Noether approximation and is used in chapter 5 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
noether(p1, alpha = 0.05, power = 0.9)
```

Arguments

p1	Probability (expressed as a number between 0 and 1)
alpha	Level of significance (expressed as number between 0 and 1) (defaults to 0.05)
power	Power (expressed as number between 0 and 1) (defaults to 0.9)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Exercise 5.8 from "Applied Nonparametric Statistical Methods" (5th edition)
noether(p1 = 0.7534, alpha = 0.05, power = 0.9)

# Exercise 5.16 from "Applied Nonparametric Statistical Methods" (5th edition)
noether(p1 = 0.8, alpha = 0.025, power = 0.9)
```

normal.scores.test*Perform Normal Scores test*

Description

`normal.scores.test()` performs the Normal Scores test and is used in chapters 6 and 8 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
normal.scores.test(
  x,
  y,
  H0 = NULL,
  alternative = c("two.sided", "less", "greater"),
  max.exact.cases = 25,
  do.asymp = FALSE,
  do.exact = TRUE
)
```

Arguments

x	Numeric vector
y	Numeric vector
H0	Null hypothesis value (defaults to NULL)
alternative	Type of alternative hypothesis (defaults to two.sided)
max.exact.cases	Maximum number of cases allowed for exact calculations (defaults to 25)
do.asymp	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
do.exact	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 5.8 from "Applied Nonparametric Statistical Methods" (5th edition)
normal.scores.test(ch6$groupA, ch6$groupB, do.exact = FALSE, do.asymp = TRUE)

# Exercise 6.15 from "Applied Nonparametric Statistical Methods" (5th edition)
normal.scores.test(ch6$doseI, ch6$doseII)
```

oddsratio.2x2diff *Perform test for difference in odds ratios*

Description

oddsratio.2x2diff() performs the test for difference in odds ratios and is used in chapter 13 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
oddsratio.2x2diff(
  x,
  y,
  z,
  alternative = c("two.sided", "less", "greater"),
  CI.width = 0.95,
  max.exact.perms = 1e+06,
  nsims.mc = 1e+05,
  seed = NULL,
  do.exact = TRUE,
  do.asymp = FALSE,
  do.mc = FALSE,
  do.CI = TRUE
)
```

Arguments

x	Binary factor of same length as y, z
y	Binary factor of same length as x, z
z	Binary factor of same length as x, y
alternative	Type of alternative hypothesis (defaults to two.sided)
CI.width	Confidence interval width (defaults to 0.95)
max.exact.perms	Maximum number of permutations allowed for exact calculations (defaults to 1000000)
nsims.mc	Number of Monte Carlo simulations to be performed (defaults to 100000)
seed	Random number seed to be used for Monte Carlo simulations (defaults to NULL)
do.exact	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)
do.asymp	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
do.mc	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to FALSE)
do.CI	Boolean indicating whether or not to perform confidence interval calculations (defaults to TRUE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 13.2 from "Applied Nonparametric Statistical Methods" (5th edition)
oddsratio.2x2diff(ch13$physical.activity, ch13$tv.viewing, ch13$gender,
  do.exact = FALSE, do.asymp = TRUE)
```

```
oddsratio.2x2diff(ch13$physical.activity, ch13$tv.viewing, ch13$gender,
  do.exact = FALSE, do.mc = TRUE, seed = 1, nsims = 10000)
```

pearson

*Calculate Pearson correlation***Description**

`pearson()` calculates the Pearson correlation and is used in chapters 10 and 11 of "Applied Non-parametric Statistical Methods" (5th edition)

Usage

```
pearson(
  x,
  y,
  alternative = c("two.sided", "less", "greater"),
  max.exact.cases = 10,
  nsims.mc = 1e+05,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE,
  do.mc = FALSE
)
```

Arguments

<code>x</code>	Numeric vector of same length as <code>y</code>
<code>y</code>	Numeric vector of same length as <code>x</code>
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to 10)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 100000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to <code>NULL</code>)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)
<code>do.mc</code>	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to FALSE)

Value

An ANSMstat object with the results from applying the function

Examples

```
# Section 10.1.2 from "Applied Nonparametric Statistical Methods" (5th edition)
pearson(ch10$q1, ch10$q2, alternative = "greater", do.asymp = TRUE, do.exact = FALSE)

# Example 11.2 from "Applied Nonparametric Statistical Methods" (5th edition)
pearson(ch11$parentlimit, ch11$reportedtime - 1 * ch11$parentlimit, alternative = "two.sided")
```

pearson.beta

Calculate Pearson beta

Description

`pearson.beta()` calculates the Pearson beta and is used in chapter 11 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
pearson.beta(
  y,
  x,
  H0 = NULL,
  alternative = c("two.sided", "less", "greater"),
  CI.width = 0.95,
  max.exact.cases = 10,
  nsims.mc = 1e+05,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE,
  do.CI = FALSE,
  do.mc = FALSE
)
```

Arguments

<code>y</code>	Numeric vector of same length as <code>x</code>
<code>x</code>	Numeric vector of same length as <code>y</code>
<code>H0</code>	Null hypothesis value (defaults to <code>NULL</code>)
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>CI.width</code>	Confidence interval width (defaults to <code>0.95</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to <code>10</code>)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to <code>100000</code>)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to <code>NULL</code>)

do.asymp	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
do.exact	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)
do.CI	Boolean indicating whether or not to perform confidence interval calculations (defaults to FALSE)
do.mc	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to FALSE)

Value

An ANSMstat object with the results from applying the function

Examples

```
# Example 11.2 from "Applied Nonparametric Statistical Methods" (5th edition)
pearson.beta(ch11$reportedtime, ch11$parentlimit, H0 = 1)
pearson.beta(ch11$reportedtime[1:6], ch11$parentlimit[1:6], H0 = 1)
```

peto.wilcoxon	<i>Perform Peto-Wilcoxon test</i>
----------------------	-----------------------------------

Description

`peto.wilcoxon()` performs the Peto-Wilcoxon test and is used in chapter 9 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
peto.wilcoxon(
  x,
  y,
  x.c,
  y.c,
  alternative = c("two.sided", "less", "greater"),
  max.exact.perms = 1e+05,
  nsims.mc = 10000,
  seed = NULL
)
```

Arguments

x	Numeric vector of same length as y, x.c, y.c
y	Numeric vector of same length as x, x.c, y.c
x.c	Binary vector of same length as x, y, x.c

```

y.c           Binary vector of same length as x, y, y.c
alternative   Type of alternative hypothesis (defaults to two.sided)
max.exact.perms
               Maximum number of permutations allowed for exact calculations (defaults to
               100000)
nsims.mc      Number of Monte Carlo simulations to be performed (defaults to 10000)
seed          Random number seed to be used for Monte Carlo simulations (defaults to NULL)

```

Value

An ANSMtest object with the results from applying the function

Examples

```

# Example 9.4 from "Applied Nonparametric Statistical Methods" (5th edition)
peto.wilcoxon(ch9$sampleI.survtime, ch9$sampleII.survtime,
               ch9$sampleI.censor, ch9$sampleII.censor, alternative = "less")

```

pitman

Perform Pitman test

Description

pitman() performs the Pitman test and is used in chapter 3 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```

pitman(
  x,
  H0 = NULL,
  alternative = c("two.sided", "less", "greater"),
  CI.width = 0.95,
  max.exact.cases = 1000,
  nsims.mc = 10000,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE,
  do.CI = TRUE
)

```

Arguments

<code>x</code>	Numeric vector
<code>H0</code>	Null hypothesis value (defaults to NULL)
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>CI.width</code>	Confidence interval width (defaults to 0.95)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to 1000)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 10000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to NULL)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)
<code>do.CI</code>	Boolean indicating whether or not to perform confidence interval calculations (defaults to TRUE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 3.11 from "Applied Nonparametric Statistical Methods" (5th edition)
pitman(ch3$heartrates1, 70, "greater", do.exact = FALSE, do.asymp = TRUE)

# Exercise 3.17 from "Applied Nonparametric Statistical Methods" (5th edition)
pitman(ch3$sampleII, 110, do.exact = FALSE, do.asymp = TRUE)
```

`print.ANSMstat` *Prints an ANSMstat object*

Description

`print.ANSMstat()` prints the output contained in an ANSMstat object

Usage

```
## S3 method for class 'ANSMstat'
print(x, ...)
```

Arguments

<code>x</code>	An ANSMstat object
<code>...</code>	Further arguments relevant to the default <code>print</code> function

Value

No return value, called to display results

print.ANSMtest	<i>Prints an ANSMtest object</i>
----------------	----------------------------------

Description

`print.ANSMtest()` prints the output contained in an ANSMtest object

Usage

```
## S3 method for class 'ANSMtest'  
print(x, ...)
```

Arguments

x	An ANSMtest object
...	Further arguments relevant to the default <code>print</code> function

Value

No return value, called to display results

rng.test	<i>Perform Range test</i>
----------	---------------------------

Description

`rng.test()` performs the Range test and is used in chapter 4 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
rng.test(x, alternative = c("two.sided"), minx = 0, maxx = 360)
```

Arguments

x	Numeric vector
alternative	Type of alternative hypothesis (defaults to <code>c("two.sided")</code>)
minx	Minimum value for x (defaults to 0)
maxx	Maximum value for x (defaults to 360)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 4.17 from "Applied Nonparametric Statistical Methods" (5th edition)
rng.test(ch4$dates.as.degrees)

# Exercise 4.13 from "Applied Nonparametric Statistical Methods" (5th edition)
rng.test(ch4$accident.bearings)
```

runs.2cat

*Perform Runs test for two categories***Description**

`runs.2cat()` performs the Runs test for two categories and is used in chapters 4, 5 and 6 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
runs.2cat(
  x,
  alternative = c("two.sided", "less", "greater"),
  cont.corr = TRUE,
  do.asymp = FALSE,
  do.exact = TRUE
)
```

Arguments

<code>x</code>	Vector with two unique values
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>cont.corr</code>	Boolean indicating whether or not to use continuity correction (defaults to <code>TRUE</code>)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to <code>FALSE</code>)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to <code>TRUE</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 4.14 from "Applied Nonparametric Statistical Methods" (5th edition)
runs.2cat(ch4$tosses1, do.exact = FALSE, do.asymp = TRUE)

# Exercise 6.17 from "Applied Nonparametric Statistical Methods" (5th edition)
runs.2cat(ch6$twins, alternative = "greater")
```

runs.ncat

Perform Runs test for three or more categories

Description

`runs.ncat()` performs the Runs test for three or more categories and is used in chapters 4 and 7 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
runs.ncat(
  x,
  alternative = c("two.sided", "less", "greater"),
  cont.corr = TRUE,
  nsims.mc = 1e+05,
  seed = NULL,
  do.asymp = TRUE,
  do.mc = FALSE
)
```

Arguments

<code>x</code>	Vector or factor
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>cont.corr</code>	Boolean indicating whether or not to use continuity correction (defaults to TRUE)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 100000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to NULL)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to TRUE)
<code>do.mc</code>	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to FALSE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 4.15 from "Applied Nonparametric Statistical Methods" (5th edition)
runs.ncat(ch4$births, alternative = "less")

# Exercise 7.16 from "Applied Nonparametric Statistical Methods" (5th edition)
runs.ncat(ch7$regions[order(ch7$affordability)], alternative = "less")
```

sgn.test

Perform Sign test

Description

`sgn.test()` performs the Sign test and is used in chapters 3, 4, 5 and 6 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
sgn.test(
  x,
  H0 = NULL,
  alternative = c("two.sided", "less", "greater"),
  cont.corr = TRUE,
  CI.width = 0.95,
  max.exact.cases = 1e+06,
  do.asymp = FALSE,
  do.exact = TRUE,
  do.CI = TRUE
)
```

Arguments

<code>x</code>	Numeric vector, or binary factor and <code>H0</code> is <code>NULL</code>
<code>H0</code>	Null hypothesis value (defaults to <code>NULL</code>)
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>cont.corr</code>	Boolean indicating whether or not to use continuity correction (defaults to <code>TRUE</code>)
<code>CI.width</code>	Confidence interval width (defaults to <code>0.95</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to <code>1000000</code>)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to <code>FALSE</code>)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to <code>TRUE</code>)
<code>do.CI</code>	Boolean indicating whether or not to perform confidence interval calculations (defaults to <code>TRUE</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 3.1 from "Applied Nonparametric Statistical Methods" (5th edition)
sgn.test(ch3$sampleI, 110)

# Exercise 6.2 from "Applied Nonparametric Statistical Methods" (5th edition)
sgn.test(ch5$LVF - ch5$RVF, 0)
```

shapirotest.ANSM *Perform Shapiro-Wilk test of Normality*

Description

shapirotest.ANSM() is a wrapper for shapiro.test() from the stats package - performs the Shapiro-Wilk test of Normality and is used in chapters 4 and 5 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
shapirotest.ANSM(x, alternative = c("two.sided"))
```

Arguments

x	Numeric vector
alternative	Type of alternative hypothesis (defaults to c("two.sided"))

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 4.4 from "Applied Nonparametric Statistical Methods" (5th edition)
shapirotest.ANSM(ch4$ages)

# Example 5.3 from "Applied Nonparametric Statistical Methods" (5th edition)
shapirotest.ANSM(ch5$bp.incorrect)
```

siegel.tukey	<i>Perform Siegel-Tukey test</i>
--------------	----------------------------------

Description

`siegel.tukey()` performs the Siegel-Tukey test using mean or median shift and is used in chapter 6 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
siegel.tukey(
  x,
  y,
  H0 = NULL,
  alternative = c("two.sided", "less", "greater"),
  mean.shift = FALSE,
  cont.corr = TRUE,
  max.exact.cases = 1000,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE
)
```

Arguments

<code>x</code>	Numeric vector
<code>y</code>	Numeric vector
<code>H0</code>	Null hypothesis value (defaults to <code>NULL</code>)
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>mean.shift</code>	Boolean indicating whether mean shift to be used instead of median shift (defaults to <code>FALSE</code>)
<code>cont.corr</code>	Boolean indicating whether or not to use continuity correction (defaults to <code>TRUE</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to 1000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to <code>NULL</code>)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to <code>FALSE</code>)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to <code>TRUE</code>)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Exercise 6.11 from "Applied Nonparametric Statistical Methods" (5th edition)
siegel.tukey(ch6$typeA, ch6$typeB, mean.shift = TRUE)

# Exercise 6.16 from "Applied Nonparametric Statistical Methods" (5th edition)
siegel.tukey(ch6$travel, ch6$politics)
```

spearman

Calculate Spearman correlation

Description

`spearman()` calculates the Spearman correlation and is used in chapter 10 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
spearman(
  x,
  y,
  alternative = c("two.sided", "less", "greater"),
  max.exact.cases = 10,
  nsims.mc = 1e+05,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE,
  do.mc = FALSE
)
```

Arguments

<code>x</code>	Numeric vector of same length as <code>y</code>
<code>y</code>	Numeric vector of same length as <code>x</code>
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to 10)
<code>nsims.mc</code>	Number of Monte Carlo simulations to be performed (defaults to 100000)
<code>seed</code>	Random number seed to be used for Monte Carlo simulations (defaults to <code>NULL</code>)
<code>do.asymp</code>	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
<code>do.exact</code>	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)
<code>do.mc</code>	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to FALSE)

Value

An ANSMstat object with the results from applying the function

Examples

```
# Example 10.2 from "Applied Nonparametric Statistical Methods" (5th edition)
spearman(ch10$q1, ch10$q2, alternative = "greater", do.asymp = TRUE, do.exact = FALSE)

# Exercise 10.1 from "Applied Nonparametric Statistical Methods" (5th edition)
spearman(ch10$ERA, ch10$ESMS, do.exact = FALSE)
```

spearman.beta

Calculate Spearman beta

Description

`spearman.beta()` calculates the Spearman beta and is used in chapter 11 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
spearman.beta(
  y,
  x,
  H0 = NULL,
  alternative = c("two.sided", "less", "greater"),
  CI.width = 0.95,
  max.exact.cases = 10,
  nsims.mc = 1e+05,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE,
  do.CI = FALSE,
  do.mc = FALSE
)
```

Arguments

<code>y</code>	Numeric vector of same length as <code>x</code>
<code>x</code>	Numeric vector of same length as <code>y</code>
<code>H0</code>	Null hypothesis value (defaults to <code>NULL</code>)
<code>alternative</code>	Type of alternative hypothesis (defaults to <code>two.sided</code>)
<code>CI.width</code>	Confidence interval width (defaults to <code>0.95</code>)
<code>max.exact.cases</code>	Maximum number of cases allowed for exact calculations (defaults to <code>10</code>)

nsims.mc	Number of Monte Carlo simulations to be performed (defaults to 100000)
seed	Random number seed to be used for Monte Carlo simulations (defaults to NULL)
do.asymp	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
do.exact	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)
do.CI	Boolean indicating whether or not to perform confidence interval calculations (defaults to FALSE)
do.mc	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to FALSE)

Value

An ANSMstat object with the results from applying the function

Examples

```
# Example 11.3 from "Applied Nonparametric Statistical Methods" (5th edition)
spearman.beta(ch11$reportedtime, ch11$parentlimit, H0 = 1)
spearman.beta(ch11$reportedtime, ch11$parentlimit, H0 = 1, do.CI = TRUE)
```

theil.kendall	<i>Calculate Theil-Kendall beta</i>
---------------	-------------------------------------

Description

theil.kendall() calculates the Theil-Kendall beta and is used in chapter 11 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
theil.kendall(
  y,
  x,
  H0 = NULL,
  do.abbreviated = FALSE,
  do.alpha = FALSE,
  alternative = c("two.sided", "less", "greater"),
  CI.width = 0.95,
  max.exact.cases = 10,
  nsims.mc = 1e+05,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE,
  do.CI = FALSE,
  do.mc = FALSE
)
```

Arguments

y	Numeric vector of same length as x
x	Numeric vector of same length as y
H0	Null hypothesis value (defaults to NULL)
do.abbreviated	Boolean indicating whether or not to use abbreviated Theil procedure (defaults to FALSE)
do.alpha	Boolean indicating whether or not to report estimate of alpha (defaults to FALSE)
alternative	Type of alternative hypothesis (defaults to two.sided)
CI.width	Confidence interval width (defaults to 0.95)
max.exact.cases	Maximum number of cases allowed for exact calculations (defaults to 10)
nsims.mc	Number of Monte Carlo simulations to be performed (defaults to 100000)
seed	Random number seed to be used for Monte Carlo simulations (defaults to NULL)
do.asymp	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
do.exact	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)
do.CI	Boolean indicating whether or not to perform confidence interval calculations (defaults to FALSE)
do.mc	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to FALSE)

Value

An ANSMstat object with the results from applying the function

Examples

```
# Example 11.6 from "Applied Nonparametric Statistical Methods" (5th edition)
theil.kendall(ch11$reportedtime, ch11$parentlimit, do.alpha = TRUE)

# Exercise 11.10 from "Applied Nonparametric Statistical Methods" (5th edition)
theil.kendall(ch11$N.Scotland, ch11$SW.England)
```

wilcoxon.mann.whitney *Perform Wilcoxon-Mann-Whitney test*

Description

wilcoxon.mann.whitney() performs the Wilcoxon-Mann-Whitney test and is used in chapters 6, 8, 9 and 12 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
wilcoxon.mann.whitney(
  x,
  y,
  H0 = NULL,
  alternative = c("two.sided", "less", "greater"),
  cont.corr = TRUE,
  CI.width = 0.95,
  max.exact.cases = 1000,
  nsims.mc = 1e+05,
  seed = NULL,
  do.asymp = FALSE,
  do.exact = TRUE,
  do.mc = FALSE,
  do.CI = TRUE
)
```

Arguments

x	Numeric vector, or factor with same levels as y
y	Numeric vector, or factor with same levels as x
H0	Null hypothesis value (defaults to NULL)
alternative	Type of alternative hypothesis (defaults to two.sided)
cont.corr	Boolean indicating whether or not to use continuity correction (defaults to TRUE)
CI.width	Confidence interval width (defaults to 0.95)
max.exact.cases	Maximum number of cases allowed for exact calculations (defaults to 1000)
nsims.mc	Number of Monte Carlo simulations to be performed (defaults to 100000)
seed	Random number seed to be used for Monte Carlo simulations (defaults to NULL)
do.asymp	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
do.exact	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)
do.mc	Boolean indicating whether or not to perform Monte Carlo calculations (defaults to FALSE)
do.CI	Boolean indicating whether or not to perform confidence interval calculations (defaults to TRUE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Examples 6.1 and 6.2 from "Applied Nonparametric Statistical Methods" (5th edition)
wilcoxon.mann.whitney(ch6$groupA, ch6$groupB)

# Exercise 12.4 from "Applied Nonparametric Statistical Methods" (5th edition)
wilcoxon.mann.whitney(ch12$feedback.satisfaction[ch12$PPI.person.2 == "Representative"],
                      ch12$feedback.satisfaction[ch12$PPI.person.2 == "Researcher"],
                      do.exact = FALSE, do.asymp = TRUE)
```

wilcoxon.signedrank *Perform Wilcoxon signed-rank test*

Description

wilcoxon.signedrank() performs the Wilcoxon signed-rank test and is used in chapters 3, 4 and 5 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
wilcoxon.signedrank(
  x,
  H0 = NULL,
  alternative = c("two.sided", "less", "greater"),
  cont.corr = TRUE,
  CI.width = 0.95,
  max.exact.cases = 1000,
  do.asymp = FALSE,
  do.exact = TRUE,
  do.CI = TRUE
)
```

Arguments

x	Numeric vector
H0	Null hypothesis value (defaults to NULL)
alternative	Type of alternative hypothesis (defaults to two.sided)
cont.corr	Boolean indicating whether or not to use continuity correction (defaults to TRUE)
CI.width	Confidence interval width (defaults to 0.95)
max.exact.cases	Maximum number of cases allowed for exact calculations (defaults to 1000)
do.asymp	Boolean indicating whether or not to perform asymptotic calculations (defaults to FALSE)
do.exact	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)
do.CI	Boolean indicating whether or not to perform confidence interval calculations (defaults to TRUE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Example 3.4 from "Applied Nonparametric Statistical Methods" (5th edition)
wilcoxon.signedrank(ch3$heartrates1, 70, "greater")

# Exercise 5.12 from "Applied Nonparametric Statistical Methods" (5th edition)
wilcoxon.signedrank(ch5$kHz0.125 - ch5$kHz0.25, 0)
```

zelen

Perform Zelen test

Description

zelen() performs the Zelen test and is used in chapter 13 of "Applied Nonparametric Statistical Methods" (5th edition)

Usage

```
zelen(x, y, z, max.exact.perms = 1e+06, do.exact = TRUE)
```

Arguments

x	Binary factor of same length as y, z
y	Binary factor of same length as x, z
z	Factor of same length as x, y
max.exact.perms	Maximum number of permutations allowed for exact calculations (defaults to 1000000)
do.exact	Boolean indicating whether or not to perform exact calculations (defaults to TRUE)

Value

An ANSMtest object with the results from applying the function

Examples

```
# Section 13.2.5 from "Applied Nonparametric Statistical Methods" (5th edition)
zelen(ch13$drug, ch13$side.effects, ch13$age.group)

# Example 13.3 from "Applied Nonparametric Statistical Methods" (5th edition)
zelen(ch13$machine, ch13$output.status, ch13$material.source)
```

Index

- * datasets
 - app1, 4
 - ch10, 9
 - ch11, 11
 - ch12, 12
 - ch13, 13
 - ch14, 15
 - ch15, 15
 - ch3, 16
 - ch4, 17
 - ch5, 18
 - ch6, 19
 - ch7, 21
 - ch8, 22
 - ch9, 24
- ansari.bradley, 3
- app1, 4
- binom, 5
- blomqvist, 6
- bowker, 7
- breslow.day, 8
- bs, 9
- ch10, 9
- ch11, 11
- ch12, 12
- ch13, 13
- ch14, 15
- ch15, 15
- ch3, 16
- ch4, 17
- ch5, 18
- ch6, 19
- ch7, 21
- ch8, 22
- ch9, 24
- chisqtest.ANSM, 25
- cochran.q, 26
- cohen.kappa, 27
- conover, 28
- control.median, 30
- cox.stuart, 31
- cramer.von.mises, 32
- fishertest.ANSM, 33
- friedman, 34
- friedman.lsd, 35
- gehan.wilcoxon, 36
- hettmansperger.elmore, 37
- hodges.ajne, 38
- jonckheere.terpstra, 39
- kendall.concordance, 40
- kendall.tau, 41
- kruskal.wallis, 42
- kruskal.wallis.lsd, 43
- kruskal.wallis.vdW, 44
- kstest.ANSM, 45
- lik.ratio, 46
- lilliefors, 47
- linear.by.linear, 48
- logoddsratio.2x2, 49
- logrank, 50
- mantel.haenszel, 51
- med.test, 52
- mood, 53
- moses.extreme.reactions, 54
- noether, 55
- normal.scores.test, 55
- oddsratio.2x2diff, 56
- pearson, 58

pearson.beta, 59
peto.wilcoxon, 60
pitman, 61
print.ANSMstat, 62
print.ANSMtest, 63

rng.test, 63
runs.2cat, 64
runs.ncat, 65

sgn.test, 66
shapirotest.ANSM, 67
siegel.tukey, 68
spearman, 69
spearman.beta, 70

theil.kendall, 71

wilcoxon.mann.whitney, 72
wilcoxon.signedrank, 74

zelen, 75