

# Package ‘jgsbook’

June 20, 2024

**Type** Package

**Title** Package of the German Book ``Statistik mit R und RStudio" by  
Joerg grosse Schlarmann

**Description** All datasets and functions used in the german book ``Statistik mit R und RStudio" by grosse Schlarmann (2010-2024) <<https://www.produnis.de/R/>>.

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|            |                              |
|------------|------------------------------|
| compare.lm | <i>Compare Linear Models</i> |
|------------|------------------------------|

---

## Description

This function fits and compares several models (linear, quadratic, cubic, exponential, logarithmic, sigmoidal, power, logistic) to a given set of dependent and independent variables. It returns either a summary of the models with their R-squared values or predicted values based on the models.

## Usage

```
compare.lm(dep, ind, predict = FALSE, steps = 0.01)
```

## Arguments

|         |                                                                                                                 |
|---------|-----------------------------------------------------------------------------------------------------------------|
| dep     | A numeric vector representing the dependent variable.                                                           |
| ind     | A numeric vector representing the independent variable.                                                         |
| predict | Logical. If TRUE, the function returns predicted values for each model. Defaults to FALSE.                      |
| steps   | Numeric. The step size for generating x-values for predictions. Only used if predict is TRUE. Defaults to 0.01. |

## Value

A data frame. If predict is FALSE, returns a data frame with the R-squared values for each model. If predict is TRUE, returns a data frame with the original data and predicted values for each model.

**Examples**

```
x <- c(6, 9, 12, 14, 30, 35, 40, 47, 51, 55, 60)
y <- c(14, 28, 50, 70, 89, 94, 90, 75, 59, 44, 27)
compare.lm(y, x)
compare.lm(y, x, predict=TRUE)
```

---

epa

*Datatable of the epa Example*

---

**Description**

Datatable of the epa Example

**Usage**

```
data(epa)
```

**Format**

A data frame with 620 observations in 6 variables

**Details**

Variables in the dataset:

- sex. a factor with levels m w d, giving the proband's sex
- age. a numeric vector
- cms. a numeric vector
- risk. a dichotome vector, 0 = not at risk, 1 = at risk
- expert. a dichotome vector of expert's decision, 0 = not at risk, 1 = at risk
- decu. a dichotome vector, 0 = no decubitus, 1 = decubitus

**Source**

<https://www.produnis.de/R/>

---

Faktorenbogen

*Datatable of the Faktorenbogen Example for factor analysis*

---

**Description**

Datatable of the Faktorenbogen Example for factor analysis

**Usage**

```
data(Faktorenbogen)
```

**Format**

A data frame with 150 observations in 14 variables

**Details**

Variables in the dataset:

- gender. a factor with levels female male other, giving the proband's gender
- age. a numeric vector of proband's age in years
- A. Item A of the questionnaire, numeric
- B. Item B of the questionnaire, numeric
- C. Item C of the questionnaire, numeric
- D. Item D of the questionnaire, numeric
- E. Item E of the questionnaire, numeric
- F. Item F of the questionnaire, numeric
- G. Item G of the questionnaire, numeric
- H. Item H of the questionnaire, numeric
- I. Item I of the questionnaire, numeric
- J. Item J of the questionnaire, numeric
- K. Item K of the questionnaire, numeric
- L. Item L of the questionnaire, numeric

**Source**

<https://www.produnis.de/R/>

---

|           |                                 |
|-----------|---------------------------------|
| freqTable | <i>create a frequency table</i> |
|-----------|---------------------------------|

---

**Description**

returns a frequency table with absolute and relative frequencies and cumulated frequencies

**Usage**

```
freqTable(werte)
```

**Arguments**

werte            factor with observed data

**Value**

dataframe table

**Examples**

```
x <- ceiling(stats::rnorm(20))
freqTable(x)
```

---

|              |                                          |
|--------------|------------------------------------------|
| kenngroessen | <i>create a tibble with kenngroessen</i> |
|--------------|------------------------------------------|

---

**Description**

returns a tibble with all kenngroessen

**Usage**

```
kenngroessen(werte)
```

**Arguments**

werte            numeric vector

**Value**

tibble with all kenngroessen

**Examples**

```
x <- ceiling(stats::rnorm(20))
kenngroessen(x)
```

---

|              |                                                              |
|--------------|--------------------------------------------------------------|
| KIbinomial_a | <i>compute confidence intervall for binomial proportions</i> |
|--------------|--------------------------------------------------------------|

---

**Description**

returns borders and length of confidence intervall for binomial proportions

**Usage**

```
KIbinomial_a(p, n, alpha)
```

**Arguments**

|       |                        |
|-------|------------------------|
| p     | proportion obeserved   |
| n     | number of observations |
| alpha | error niveau           |

**Value**

confidence intervall

**Examples**

```
KIbinomial_a(0.35, 150, 0.05)
```

---

|              |                                                                            |
|--------------|----------------------------------------------------------------------------|
| KIbinomial_u | <i>compute confidence intervall for difference of binomial proportions</i> |
|--------------|----------------------------------------------------------------------------|

---

**Description**

returns borders and length of confidence intervall for difference of binomial proportions

**Usage**

```
KIbinomial_u(p1, n1, p2, n2, alpha)
```

**Arguments**

|       |                                   |
|-------|-----------------------------------|
| p1    | proportion obeserved in group 1   |
| n1    | number of observations in group 1 |
| p2    | proportion obeserved in group 2   |
| n2    | number of observations in group 2 |
| alpha | error niveau                      |

**Value**

confidence intervall

**Examples**

```
KIbinomial_u(0.25, 100, 0.4, 150, 0.05)
```

---

|            |                                                                         |
|------------|-------------------------------------------------------------------------|
| KInormal_a | <i>compute confidence intervall for mean of normal distributed data</i> |
|------------|-------------------------------------------------------------------------|

---

**Description**

returns borders and length of confidence intervall for mean of normal distributed data

**Usage**

```
KInormal_a(xquer, s, n, alpha)
```

**Arguments**

|       |                                     |
|-------|-------------------------------------|
| xquer | mean of obeserved data              |
| s     | standard deviation of observed data |
| n     | number of observations              |
| alpha | error niveau                        |

**Value**

confidence intervall

**Examples**

```
KInormal_a(400, 20, 100, 0.05)
```

---

|            |                                                                         |
|------------|-------------------------------------------------------------------------|
| KInormal_u | <i>compute confidence intervall for mean of normal distributed data</i> |
|------------|-------------------------------------------------------------------------|

---

**Description**

returns a data.frame with borders and length of confidence intervall for mean of normal distributed data

**Usage**

```
KInormal_u(x1, s1, n1, x2, s2, n2, alpha)
```

**Arguments**

|       |                                                |
|-------|------------------------------------------------|
| x1    | mean of obeserved data in group 1              |
| s1    | standard deviation of observed data in group 1 |
| n1    | number of observations in group 1              |
| x2    | mean of obeserved data in group 2              |
| s2    | standard deviation of observed data in group 2 |
| n2    | number of observations in group 2              |
| alpha | error niveau                                   |

**Value**

data.frame of confidence intervall

**Examples**

```
KInormal_u(2.22, 0.255, 13, 2.7, 0.306, 10 , 0.05)
```

---

|             |                                                                                                                                                                    |
|-------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| lon.lat.osm | <i>get longitude and altitude from an address using OpenStreetMap's API at <a href="http://nominatim.openstreetmap.org">http://nominatim.openstreetmap.org</a></i> |
|-------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------|

---

**Description**

get longitude and altitude from an address using OpenStreetMap's API at <http://nominatim.openstreetmap.org>

**Usage**

```
lon.lat.osm(address = NULL)
```



**Arguments**

address            a character of an address

**Value**

a data.frame containig "address", "lon", "lat"

**Examples**

```
lon.lat.osm("Eiffeltower")
```

---

MarioANOVA

*Datatable of the SuperMario Example for Friedman-ANOVA*

---

**Description**

Datatable of the SuperMario Example for Friedman-ANOVA

**Usage**

```
data(MarioANOVA)
```

**Format**

A data frame with 47 observations in 8 variables

**Details**

Variables in the dataset:

- Name. The characters' name
- Alter. The characters' age in years
- Kingdom. The characters' home
- Geschlecht. The characters' gender (männlich = male, weiblich = female)
- BadGuy. Whether the character is a bad guy, logical
- t1. Measure at time 1
- t2. Measure at time 2
- t3. Measure at time 3

**Source**

<https://www.produnis.de/R/>

---

Messwiederholung

*Datatable of the Messwiederholung Example for ANOVA*

---

**Description**

Datatable of the Messwiederholung Example for ANOVA

**Usage**

```
data(Messwiederholung)
```

**Format**

A data frame with 200 observations in 4 variables

**Details**

Variables in the dataset:

- Name. The first name of the probands.
- t1. Measure at time 1
- t2. Measure at time 2
- t3. Measure at time 3

**Source**

<https://www.produnis.de/R/>

---

mma

*Dataset of a work sampling study*

---

**Description**

Dataset of a work sampling study

**Usage**

```
data(mma)
```

**Format**

A data frame with 9768 observations in 6 variables.

**Details**

Variables in the dataset:

- day. a vector, giving the number of the observation day
- time. a factor giving the time of observation
- ward. a factor giving the ward under observation
- qual. a factor giving the qualification of the nurse
- category. a factor of qualification categories
- action. a factor giving the observed action

**Source**

<https://www.produnis.de/R/>

---

Nachtwachen

*Dataset of the German Nachtwachen study*

---

**Description**

Dataset of the German Nachtwachen study

**Usage**

`data(Nachtwachen)`

**Format**

A data frame with 276 observations in 37 variables.

**Source**

<https://www.produnis.de/R/>

nw

*Dataset of the German Nachtwachen study with labelled variables*

---

**Description**

Dataset of the German Nachtwachen study, labelled version

**Usage**

```
data(nw)
```

**Format**

A data frame with 276 observations in 37 variables.

**Source**

<https://www.produnis.de/R/>

---

OrdinalSample

*Datatable of an Ordinal Sample*

---

**Description**

Datatable of an Ordinal Sample

**Usage**

```
data(OrdinalSample)
```

**Format**

A data frame with 415 observations in 4 variables.

**Details**

Variables in the dataset:

- *Konflikt*. a numeric vector giving the potential of conflicts.
- *Zufriedenh*. a numeric vector giving the satisfaction of workers
- *Geschlecht*. a factor of proband's sex, 1 = male, 2=female
- *Stimmung*. an ordinal factor of proband's mood

**Source**

<https://www.produnis.de/R/>

---

pairwise.chisq.test     *Pairwise Chi-Square Tests*

---

### Description

This function performs pairwise Chi-Square tests for two factors.

### Usage

```
pairwise.chisq.test(A, B, p.adjust.method = "bonferroni")
```

### Arguments

A                     A factor with two or more levels. The first variable.  
B                     A factor with two or more levels. The second variable.  
p.adjust.method     A string specifying the method for adjusting p-values. Default is "bonferroni".

### Details

This function creates all possible pairs of levels of factor B and performs a Chi-Square test for each pair of B on variable A. The p-values are adjusted according to the specified method. #' This function is created for educational purposes only. For exact p-values, consider using `reporttools::pairwise.fisher.test()`.

### Value

A data frame with the results of the pairwise Chi-Square tests. Includes the groups, Chi-Square statistic, degrees of freedom, p-values, adjusted p-values, and significance stars.

### Examples

```
set.seed(123)
A <- factor(sample(c("Male", "Female"), 100, replace = TRUE))
B <- factor(sample(c("Location1", "Location2", "Location3"), 100, replace = TRUE))
pairwise.chisq.test(A, B, "holm")
```

---

pf8                     *Dataset of the PF8 example.*

---

### Description

This is the dataset of the PF8 example.

### Usage

```
data(pf8)
```

**Format**

A data frame with 731 observations in 16 variables.

**Source**

<https://www.produnis.de/R/>

---

Pflegeberufe

*Matrix of Pflegeberufe by Isfort et al. 2018*

---

**Description**

Matrix of Pflegeberufe by Isfort et al. 2018

**Usage**

```
data(Pflegeberufe)
```

**Format**

A matrix with 9 cols (years) and 5 rows (nursing profession).

**Author(s)**

Isfort et al. 2018 (Pflegethermometer)

**Source**

<https://www.produnis.de/R/>

---

sens.spec

*compute sensitivity and specificity*

---

**Description**

returns sensitivity specificity, negativ-predictive-value, positiv-predictive-value

**Usage**

```
sens.spec(rp, rn, fp, fn)
```

**Arguments**

|    |                                           |
|----|-------------------------------------------|
| rp | number of true-positive (richtig-positiv) |
| rn | number of true-negative (richtig-negativ) |
| fp | number of false-positive (falsch-positiv) |
| fn | number of false-negative (falsch-negativ) |

**Value**

a data.frame with sens, spec, ppw, npw

**Examples**

```
sens.spec(40, 17, 85, 4)
```

---

|        |                                                                                |
|--------|--------------------------------------------------------------------------------|
| ztrans | <i>z-Transformation by given numbers, with <math>z = (x - \mu) / sd</math></i> |
|--------|--------------------------------------------------------------------------------|

---

**Description**

z-Transformation by given numbers, with  $z = (x - \mu) / sd$

**Usage**

```
ztrans(x, mu = 0, sd = 1)
```

**Arguments**

|    |                              |
|----|------------------------------|
| x  | a value to transform         |
| mu | the given mu                 |
| sd | the given standard deviation |

**Value**

the z-transformed value

**Examples**

```
ztrans(120, mu=118, sd=20)
```

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