

Package ‘krippendorffsalpha’

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Type Package

Title Measuring Agreement Using Krippendorff's Alpha Coefficient

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Author John Hughes

Maintainer John Hughes <drjphughesjr@gmail.com>

Suggests parallel, pbapply, testthat (>= 3.0.0)

Description Provides tools for applying Krippendorff's Alpha methodology <DOI:10.1080/19312450709336664>. The framework supports common and user-defined distance functions, and can accommodate any number of units, any number of coders, and missingness. Bootstrap inference is permitted, and the computation can be done in parallel.

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cartilage	<i>Data from an MRI study of hip cartilage in femoroacetabular impingement.</i>
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Description

This data frame has exactly two columns. The first column contains raw T2* values, the second column contrast-enhanced T2* values.

Usage

```
data(cartilage)
```

Format

A data frame having 323 rows and two columns

References

Nissi, M. J., Mortazavi, S., Hughes, J., Morgan, P., and Ellermann, J. (2015). T2* relaxation time of acetabular and femoral cartilage with and without intra-articular Gd-DTPA2 in patients with femoroacetabular impingement. *American Journal of Roentgenology*, **204**(6), W695.

confint.krippendorffsalph	<i>Compute a confidence interval for Krippendorff's Alpha.</i>
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Description

Compute a confidence interval for Krippendorff's Alpha.

Usage

```
## S3 method for class 'krippendorffsalph'
confint(object, parm = "alpha", level = 0.95, ...)
```

Arguments

object	an object of class "krippendorffsalph", the result of a call to krippendorffs.alpha .
parm	always ignored since there is only one parameter.
level	the desired confidence level for the interval. The default is 0.95.
...	additional arguments. These are passed to quantile .

Details

This function computes a bootstrap confidence interval for alpha, assuming that `krippendorffs.alpha` was called with `confint = TRUE`.

Value

A vector with entries giving lower and upper confidence limits. These will be labelled as $(1 - \text{level}) / 2$ and $1 - (1 - \text{level}) / 2$.

References

Nissi, M. J., Mortazavi, S., Hughes, J., Morgan, P., and Ellermann, J. (2015). T2* relaxation time of acetabular and femoral cartilage with and without intra-articular Gd-DTPA2 in patients with femoroacetabular impingement. *American Journal of Roentgenology*, **204**(6), W695.

See Also

[krippendorffs.alpha](#)

Examples

```
# Fit a subset of the cartilage data. Compute bootstrap confidence intervals
# using a bootstrap sample size of 1,000.

data(cartilage)
cartilage = as.matrix(cartilage[1:100, ])
fit.cart = krippendorffs.alpha(cartilage, level = "interval", confint = TRUE,
                              control = list(bootit = 1000, parallel = FALSE))
fit.cart$alpha.hat
confint(fit.cart, level = 0.99)
```

influence.krippendorffsalpha

Compute DFBETAs for units and/or coders.

Description

Compute DFBETAs for units and/or coders.

Usage

```
## S3 method for class 'krippendorffsalpha'
influence(model, units, coders, ...)
```

Arguments

model	a fitted model object, the result of a call to <code>krippendorffs.alpha</code> .
units	a vector of integers. A DFBETA will be computed for each of the corresponding units.
coders	a vector of integers. A DFBETA will be computed for each of the corresponding coders.
...	additional arguments. These are ignored.

Details

This function computes DFBETAs for one or more units and/or one or more coders.

Value

A list comprising at most two elements.

<code>dfbeta.units</code>	a vector containing DFBETAs for the units specified via argument <code>units</code> .
<code>dfbeta.coders</code>	a vector containing DFBETAs for the coders specified via argument <code>coders</code> .

References

Young, D. S. (2017). *Handbook of Regression Methods*. CRC Press.

Krippendorff, K. (2013). Computing Krippendorff's alpha-reliability. Technical report, University of Pennsylvania.

Examples

```
# The following data were presented in Krippendorff (2013).

nominal = matrix(c(1,2,3,3,2,1,4,1,2,NA,NA,NA,
                  1,2,3,3,2,2,4,1,2,5,NA,3,
                  NA,3,3,3,2,3,4,2,2,5,1,NA,
                  1,2,3,3,2,4,4,1,2,5,1,NA), 12, 4)
fit.nom = krippendorffs.alpha(nominal, level = "nominal", confint = FALSE)
summary(fit.nom)
(inf = influence(fit.nom, units = c(6, 11), coders = c(2, 3)))
```

interval.dist	<i>Compute the squared difference between two scores.</i>
---------------	---

Description

Compute the squared difference between two scores.

Usage

```
interval.dist(x, y)
```

Arguments

x	a score.
y	a score.

Details

This function computes the squared difference between two scores. This may be an appropriate distance function for the interval level of measurement. NA's are handled gracefully.

Value

$(x - y)^2$, or 0 if x or y is NA.

See Also

[nominal.dist](#), [ratio.dist](#)

krippendorffs.alpha *Apply Krippendorff's Alpha.*

Description

Apply Krippendorff's Alpha.

Usage

```
krippendorffs.alpha(
  data,
  level = c("interval", "nominal", "ordinal", "ratio"),
  confint = TRUE,
  verbose = FALSE,
  control = list()
)
```

Arguments

data	a matrix of scores. Each row corresponds to a unit, each column a coder.
level	the level of measurement, one of "nominal", "ordinal", "interval", or "ratio"; or a user-defined distance function.
confint	logical; if TRUE, a bootstrap sample is produced.
verbose	logical; if TRUE, various messages are printed to the console. Note that if confint = TRUE a progress bar (pblapply) is displayed (if possible) during the bootstrap computation.
control	a list of control parameters. bootit the size of the bootstrap sample. This applies when confint = TRUE. Defaults to 1,000.

`nodes` the desired number of nodes in the cluster.
`parallel` logical; if TRUE (the default), bootstrapping is done in parallel.
`type` one of the supported cluster types for `makeCluster`. Defaults to "SOCK".

Details

This is the package's flagship function. It applies the Krippendorff's Alpha methodology for nominal, ordinal, interval, or ratio levels of measurement, and, if desired, produces confidence intervals. Parallel computing is supported, when applicable.

If the level of measurement is nominal, the discrete metric (`nominal.dist`) is employed by default. If the level of measurement is interval or ordinal, the squared-difference distance function (`interval.dist`) is employed by default. (For the ordinal level of measurement, using the squared-difference distance function may be inappropriate, in which case the user should supply his/her own distance function.) If the level of measurement is ratio, a ratio distance function (`ratio.dist`) is applied. Alternatively, the user may supply his/her own distance function. Said function must handle NA's gracefully; see the above mentioned built-in distance functions for examples.

If argument `confint` is set to TRUE, bootstrapping is carried out. This is done by resampling, with replacement, the rows of data and then computing the alpha statistic for the resulting matrix. The elements of argument `control` are used to control the bootstrap computation.

Value

Function `krippendorffs.alpha` returns an object of class "krippendorffsalph", which is a list comprising the following elements.

<code>boot.sample</code>	when applicable, the bootstrap sample.
<code>call</code>	the matched call.
<code>coders</code>	the number of coders.
<code>alpha.hat</code>	the estimate of alpha.
<code>confint</code>	the value of argument <code>confint</code> .
<code>control</code>	the list of control parameters.
<code>data</code>	the matrix of scores.
<code>D.e</code>	the estimate of total variation.
<code>D.o</code>	the estimate of within-unit variation.
<code>level</code>	the level of measurement.
<code>units</code>	the number of units.
<code>verbose</code>	the value of argument <code>verbose</code> .

References

Krippendorff, K. (2013). Computing Krippendorff's alpha-reliability. Technical report, University of Pennsylvania.

Examples

```
# The following data were presented in Krippendorff (2013).

nominal = matrix(c(1,2,3,3,2,1,4,1,2,NA,NA,NA,
                  1,2,3,3,2,2,4,1,2,5,NA,3,
                  NA,3,3,3,2,3,4,2,2,5,1,NA,
                  1,2,3,3,2,4,4,1,2,5,1,NA), 12, 4)

nominal
fit.nom = krippendorffs.alpha(nominal, level = "nominal", confint = TRUE, verbose = TRUE,
                              control = list(bootit = 100, parallel = FALSE))

summary(fit.nom)
confint(fit.nom, level = 0.99)
```

nominal.dist	<i>Apply the discrete metric to two scores.</i>
--------------	---

Description

Apply the discrete metric to two scores.

Usage

```
nominal.dist(x, y)
```

Arguments

x	a score.
y	a score.

Details

This function applies the discrete metric to two scores. This may be an appropriate distance function for the nominal level of measurement. NA's are handled gracefully.

Value

0 if x is equal to y or if either is NA, 1 otherwise.

See Also

[interval.dist](#), [ratio.dist](#)

```
plot.krippendorffsalpha
```

Plot the results of a Krippendorff's Alpha analysis.

Description

Plot the results of a Krippendorff's Alpha analysis.

Usage

```
## S3 method for class 'krippendorffsalpha'
plot(
  x,
  y = NULL,
  level = 0.95,
  type = 7,
  density = TRUE,
  lty.density = 1,
  lty.estimate = 1,
  lty.interval = 2,
  col.density = "black",
  col.estimate = "orange",
  col.interval = "blue",
  lwd.density = 3,
  lwd.estimate = 3,
  lwd.interval = 3,
  ...
)
```

Arguments

x	an object of class "krippendorffsalpha", the result of a call to <code>krippendorffs.alpha</code> .
y	always ignored.
level	the desired confidence level for the interval. The default is 0.95.
type	the method used to compute sample quantiles. This argument is passed to <code>quantile</code> . The default is 7.
density	logical; if TRUE, a kernel density estimate is plotted.
lty.density	the line type for the kernel density estimate. The default is 1.
lty.estimate	the line type for the estimate of alpha. The default is 1.
lty.interval	the line type for the confidence limits. The default is 2.
col.density	the color for the kernel density estimate. The default is black.
col.estimate	the color for the estimate of alpha. The default is orange.
col.interval	the color for the confidence limits. The default is blue.
lwd.density	the line width for the kernel density estimate. The default is 3.

lwd.estimate the line width for the estimate of alpha. The default is 3.
 lwd.interval the line width for the confidence limits. The default is 3.
 ... additional arguments. These are passed to [hist](#).

Details

This function plots the results of a Krippendorff's Alpha analysis, assuming that [krippendorffs.alpha](#) was called with `confint = TRUE`.

References

Krippendorff, K. (2013). Computing Krippendorff's alpha-reliability. Technical report, University of Pennsylvania.

Examples

```
# The following data were presented in Krippendorff (2013).

nominal = matrix(c(1,2,3,3,2,1,4,1,2,NA,NA,NA,
                  1,2,3,3,2,2,4,1,2,5,NA,3,
                  NA,3,3,3,2,3,4,2,2,5,1,NA,
                  1,2,3,3,2,4,4,1,2,5,1,NA), 12, 4)
fit.nom = krippendorffs.alpha(nominal, level = "nominal", confint = TRUE, verbose = TRUE,
                              control = list(bootit = 1000, parallel = FALSE))
dev.new()
plot(fit.nom, main = "Results for Nominal Data", xlab = "Bootstrap Estimates", density = FALSE)
```

<code>ratio.dist</code>	<i>Apply a ratio distance function to two scores.</i>
-------------------------	---

Description

Apply a ratio distance function to two scores.

Usage

```
ratio.dist(x, y)
```

Arguments

x a score.
 y a score.

Details

This function applies a ratio distance function to two scores. This may be an appropriate distance function for the ratio level of measurement. NA's are handled gracefully.

Value

$(x - y)^2 / (x + y)^2$, or 0 if x or y is NA.

See Also

[interval.dist](#), [nominal.dist](#)

summary.krippendorffsalpha

Print a summary of a Krippendorff's Alpha fit.

Description

Print a summary of a Krippendorff's Alpha fit.

Usage

```
## S3 method for class 'krippendorffsalpha'  
summary(object, conf.level = 0.95, digits = 4, ...)
```

Arguments

object	an object of class "krippendorffsalpha", the result of a call to krippendorffs.alpha .
conf.level	the confidence level for the confidence intervals. The default is 0.95.
digits	the number of significant digits to display. The default is 4.
...	additional arguments. These are passed to quantile .

Details

This function prints a summary of the fit. First the values of the control parameters (defaults and/or values supplied in the call) are printed. Then a table of estimates is shown. If applicable, the table includes confidence intervals.

References

Nissi, M. J., Mortazavi, S., Hughes, J., Morgan, P., and Ellermann, J. (2015). T2* relaxation time of acetabular and femoral cartilage with and without intra-articular Gd-DTPA2 in patients with femoroacetabular impingement. *American Journal of Roentgenology*, **204**(6), W695.

See Also

[krippendorffs.alpha](#)

Examples

```
# Fit a subset of the cartilage data. Compute bootstrap confidence intervals
# using a bootstrap sample size of 1,000. Display a summary of the results,
# including a 99% confidence interval. Also plot the results.

data(cartilage)
cartilage = as.matrix(cartilage[1:100, ])
fit.cart = krippendorffs.alpha(cartilage, level = "interval", confint = TRUE,
                              control = list(bootit = 1000, parallel = FALSE))
summary(fit.cart, conf.level = 0.99)
dev.new()
plot(fit.cart, xlim = c(0.7, 0.9), xlab = "Bootstrap Estimates",
     main = "Results for Cartilage Data")
```

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