

Package: mvalpha (via r-universe)

June 18, 2026

Type Package

Title Krippendorff's Alpha for Multi-Valued Data

Version 0.6.0

Description Calculate Krippendorff's alpha for multi-valued data using the methods introduced by Krippendorff and Craggs (2016) [doi:10.1080/19312458.2016.1228863](https://doi.org/10.1080/19312458.2016.1228863). Nominal, ordinal, interval, and ratio data types are supported, with option to create bootstrapped estimates of alpha.

Roxygen list(markdown = TRUE)

License AGPL (>= 3)

Encoding UTF-8

URL <https://github.com/therealcfdrake/mvalpha>

BugReports <https://github.com/therealcfdrake/mvalpha/issues>

Depends R (>= 4.2.0)

RoxygenNote 7.3.3

LazyData true

Imports stats, utils, rlang, Rdpack, Rcpp, arrangements, stringr

Suggests testthat (>= 3.0.0)

RdMacros Rdpack

LinkingTo Rcpp

Config/testthat/edition 3

SystemRequirements OpenMP

Config/pak/sysreqs libgmp3-dev libicu-dev

Repository <https://eddelbuettel.r-universe.dev>

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RemoteUrl <https://github.com/eddelbuettel/mvalpha>

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|-----------|---------------------------|
| ex_table3 | <i>Published Examples</i> |
|-----------|---------------------------|

Description

These data represent examples found in the original paper describing the calculation of multi-valued Krippendorff’s alpha by Krippendorff and Craggs (2016).

Usage

ex_table3
 ex_table8a
 ex_table8b
 ex_table8c
 ex_table9a
 ex_table9b
 ex_table9c

Format

Each is a multi-valued nominal set with observers as columns and units as rows.

Source

[doi:10.1080/19312458.2016.1228863](https://doi.org/10.1080/19312458.2016.1228863)

References

Krippendorff K, Craggs R (2016). “The Reliability of Multi-Valued Coding of Data.” *Communication Methods and Measures*, **10**(4), 181–198. [doi:10.1080/19312458.2016.1228863](https://doi.org/10.1080/19312458.2016.1228863).

generate_mv_data *Generate Multi-Valued Data Sets*

Description

Generate Multi-Valued Data Sets

Usage

```
generate_mv_data(  
  type = "nominal",  
  n_units = 10,  
  n_observers = 3,  
  n_labels = 5,  
  tpr = 0.8,  
  fpr = 0.01,  
  card_pmf = c(0.3, 0.4, 0.2, 0.1),  
  p_missing = 0  
)
```

Arguments

| | |
|-------------|---|
| type | Data type. One of "nominal", "ordinal", "interval", or "ratio". |
| n_units | Number of units (rows) in data. |
| n_observers | Number of observers (cols) in data. |
| n_labels | Number of possible labels which could be applied to the data. |
| tpr | True Positive Rate. To generate the data, first a latent set of labels. tpr describes the probability that a latent label will be identified by each observer. |
| fpr | False Positive Rate. To generate the data, first a latent set of labels. fpr describes the probability that each observer includes each label not in the latent set. |
| card_pmf | Probability mass function describing the cardinality of observed label sets. Length should be less than or equal to n_labels. |
| p_missing | Proportion of observations that are randomly missing, indicated by NA. These are distinct from observations which are made, but the observer applied 0 labels, indicated by NULL. |

Value

A matrix with n_units rows and n_observers columns with list elements.

Examples

```
generate_mv_data()
```

mvalpha

Estimate Multi-Valued Krippendorff's Alpha

Description

mvalpha() calculates Krippendorff's alpha statistic when multi-valued observers are allowed to apply multiple values to an observation.

Usage

```
mvalpha(data, type = "nominal", verbose = TRUE, n_boot = NULL, n_threads = 1)
```

Arguments

| | |
|-----------|--|
| data | a data frame containing a list column for each observer. Each row represents an observation unit, and each cell contains a vector of 0 to w unique values, where w is the number of unique values found in the data set. NA values are used to represent missing observations and NULL values represent the empty set, {}, of responses. |
| type | a string describing the data type of the label set. This can be "nominal", "ordinal", "interval", or "ratio" and is used to select the appropriate distance metric. |
| verbose | a logical value which toggles whether status updates are printed to the console while alpha is being calculated. |
| n_boot | an integer representing the number of bootstrap estimates to calculate for mvDo. The default, NULL, will not generate additional estimates. |
| n_threads | an integer describing the number of cores to allocate to parallelization. |

Value

An object of class mvalpha

References

Krippendorff K, Craggs R (2016). "The Reliability of Multi-Valued Coding of Data." *Communication Methods and Measures*, **10**(4), 181–198. doi:10.1080/19312458.2016.1228863.

Examples

```
library(mvalpha)

### replicate example from Table 3 in Krippendorff and Craggs (2016) with bootstrapped estimates

# View data
ex_table3

# # Estimate alpha
# x <- mvalpha(ex_table3, verbose = TRUE, n_boot = 500)
```

```
#  
# # View result  
# x  
#  
# # View the unique values observed in the data  
# x$values  
#  
# # View the unique labels used to code the data  
# x$labels  
#  
# # Histogram of bootstrapped estimates  
# hist(x$bootstrap_mvalpha)
```

new_mvalpha

Create new mvalpha class object

Description

Wrapper for creating mvalpha class object.

Usage

```
new_mvalpha(  
  mvalpha,  
  type,  
  mvDo,  
  mvDe,  
  bootstrap_mvalpha,  
  unique_cardinalities,  
  units,  
  observers,  
  labels,  
  values,  
  values_by_unit,  
  dist_CK,  
  p_CK,  
  data  
)
```

Arguments

| | |
|---------|---|
| mvalpha | Multi-valued alpha estimate |
| type | a string describing the data type of the label set. This can be "nominal", "ordinal", "interval", or "ratio" and is used to select the appropriate distance metric. |
| mvDo | Observed disagreement |
| mvDe | Expected disagreement |

| | |
|-----------------------------------|--|
| <code>bootstrap_mvalpha</code> | Bootstrap estimates of <code>mvalpha</code> |
| <code>unique_cardinalities</code> | Numeric vector of the unique cardinalities observed in the data |
| <code>units</code> | Names of units |
| <code>observers</code> | Names of observers |
| <code>labels</code> | Unique labels used in data |
| <code>values</code> | Unique values used in data |
| <code>values_by_unit</code> | Table of values by unit |
| <code>dist_CK</code> | Distance matrix for label sets C and K |
| <code>p_CK</code> | Probability matrix for label sets C and K |
| <code>data</code> | a data frame containing a list column for each observer. Each row represents an observation unit, and each cell contains a vector of 0 to <code>w</code> unique values, where <code>w</code> is the number of unique values found in the data set. NA values are used to represent missing observations and NULL values represent the empty set, {}, of responses. |

Value

an `mvalpha` object

| | |
|----------------------------|-----------------------------------|
| <code>print.mvalpha</code> | <i>Print mvalpha class object</i> |
|----------------------------|-----------------------------------|

Description

Print generic

Usage

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

Arguments

| | |
|------------------|-----------------------------|
| <code>x</code> | <code>mvalpha</code> object |
| <code>...</code> | additional parameters |

Value

invisibly returns the alpha estimate of an `mvalpha` object

Description

Find the intersection and set difference(s) of two sets all at once and more efficiently than calling `base::intersect()` and `base::setdiff()` separately. Based on this [stackoverflow answer](https://stackoverflow.com/a/72631719) <https://stackoverflow.com/a/72631719>

Usage

```
set_ops(A, B, type)
```

Arguments

| | |
|------|---|
| A, B | sets (vectors) of elements |
| type | a string describing the data type of the label set. This can be "nominal", "ordinal", "interval", or "ratio" and is used to select the appropriate distance metric. |

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