

Package ‘Keng’

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Title Knock Errors Off Nice Guesses

Version 2024.11.25

Description Miscellaneous functions and data used in Qingyao's psychological research and teaching. Keng currently has a built-in dataset `depress`, and could (1) scale a vector, (2) test the significance and compute the cut-off values of Pearson's r without raw data, (3) compare `lm()`'s fitted outputs using R-squared, `f_squared`, post-hoc power, and PRE (Proportional Reduction in Error, also called partial R-squared or partial Eta-squared). (4) Calculate PRE from partial correlation, Cohen's f , or `f_squared`. (5) Compute the post-hoc power for one or a set of predictors in regression analysis without raw data, (6) Plan the sample size for one or a set of predictors in regression analysis.

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Encoding UTF-8

RoxygenNote 7.3.2

Imports stats

Suggests knitr, rmarkdown, car, effectsize, testthat (>= 3.0.0)

Config/testthat/edition 3

URL <https://github.com/qyaozh/Keng>

BugReports <https://github.com/qyaozh/Keng/issues>

Depends R (>= 2.10)

LazyData true

VignetteBuilder knitr

NeedsCompilation no

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Contents

calc_PRE	2
compare_lm	3
cut_r	4
depress	5
power_lm	6
Scale	7
test_r	8

Index	10
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calc_PRE	<i>Calculate PRE from Cohen's f, f_squared, or partial correlation</i>
----------	--

Description

Calculate PRE from Cohen's f, f_squared, or partial correlation

Usage

```
calc_PRE(f = NULL, f_squared = NULL, r_p = NULL)
```

Arguments

f	Cohen's f. Cohen (1988) suggested ≥ 0.1 , ≥ 0.25 , and ≥ 0.40 as cut-off values of f for small, medium, and large effect sizes, respectively.
f_squared	Cohen's f_squared. Cohen (1988) suggested ≥ 0.02 , ≥ 0.15 , and ≥ 0.35 as cut-off values of f for small, medium, and large effect sizes, respectively.
r_p	Partial correlation.

Value

A list including PRE, r_p (partial correlation), Cohen's f_squared, and f.

References

Cohen, J. (1988). *Statistical power analysis for the behavioral sciences* (2nd ed.). Routledge.

Examples

```
calc_PRE(f = 0.1)
calc_PRE(f_squared = 0.02)
calc_PRE(r_p = 0.2)
```

compare_lm	<i>Compare lm()'s fitted outputs using PRE and R-squared.</i>
------------	---

Description

Compare lm()'s fitted outputs using PRE and R-squared.

Usage

```
compare_lm(
  fitC = NULL,
  fitA = NULL,
  n = NULL,
  PC = NULL,
  PA = NULL,
  SSEC = NULL,
  SSEA = NULL
)
```

Arguments

fitC	The result of lm() of the Compact model (model C).
fitA	The result of lm() of the Augmented model (model A).
n	Sample size of the model C or model A. model C and model A must use the same sample, and hence have the same sample size.
PC	The number of parameters in model C.
PA	The number of parameters in model A. PA must be larger than PC.
SSEC	The Sum of Squared Errors (SSE) of model C.
SSEA	The Sum of Squared Errors of model A.

Details

compare_lm() compare model A with model C using *PRE* (Proportional Reduction in Error) , *R-squared*, *f_squared*, and post-hoc power. *PRE* is partial R-squared (called partial Eta-squared in Anova). There are two ways of using compare_lm(). The first is giving compare_lm() fitC and fitA. The second is giving n, PC, PA, SSEC, and SSEA. The first way is more convenient, and it minimizes precision loss by omitting copying-and-pasting. Note that the *F*-tests for *PRE* and that for R-squared change are equivalent. Please refer to Judd et al. (2017) for more details about *PRE*, and refer to Aberson (2019) for more details about *f_squared* and post-hoc power.

Value

A matrix with 11 rows and 4 columns. The first column reports information for baseline model (intercept-only model), the second for model C, the third for model A, and the fourth for the change (model A vs. model C). *SSE* (Sum of Squared Errors) and *df* of *SSE* for baseline model, model C,

model A, and change (model A vs. model C) are reported in row 1 and row 2. The information in the fourth column are all for the change; put differently, These results could quantify the effect of one or a set of new parameters model A has but model C doesn't. If fitC and fitA are not inferior to the intercept-only model, *R-squared*, *Adjusted R-squared*, *PRE*, *PRE_adjusted*, and *f_squared* for the full model (compared with the baseline model) are reported for model C and model A. If model C or model A has at least one predictor, *F* -test with *p*, and post-hoc power would be computed for the corresponding full model.

References

- Aberson, C. L. (2019). *Applied power analysis for the behavioral sciences*. Routledge.
- Judd, C. M., McClelland, G. H., & Ryan, C. S. (2017). *Data analysis: A model Comparison approach to regression, ANOVA, and beyond*. Routledge.

Examples

```
x1 <- rnorm(193)
x2 <- rnorm(193)
y <- 0.3 + 0.2*x1 + 0.1*x2 + rnorm(193)
dat <- data.frame(y, x1, x2)
# Fix intercept to constant 1 using I().
fit1 <- lm(I(y - 1) ~ 0, dat)
# Free intercept.
fit2 <- lm(y ~ 1, dat)
compare_lm(fit1, fit2)
# One predictor.
fit3 <- lm(y ~ x1, dat)
compare_lm(fit2, fit3)
# Fix intercept to 0.3 using offset().
intercept <- rep(0.3, 193)
fit4 <- lm(y ~ 0 + x1 + offset(intercept), dat)
compare_lm(fit4, fit3)
# Two predictors.
fit5 <- lm(y ~ x1 + x2, dat)
compare_lm(fit2, fit5)
compare_lm(fit3, fit5)
# Fix slope of x2 to 0.05 using offset().
fit6 <- lm(y ~ x1 + offset(0.05*x2), dat)
compare_lm(fit6, fit5)
```

cut_r

Cut-off values of r given the sample size n.

Description

Cut-off values of r given the sample size n.

Usage

```
cut_r(n)
```

Arguments

n Sample size of the r .

Details

Given n and p , t and then r could be determined. The formula used could be found in `test_r()`'s documentation.

Value

A data.frame including the cut-off values of r at the significance levels of $p = 0.1, 0.05, 0.01, 0.001$. r with the absolute value larger than the cut-off value is significant at the corresponding significance level.

Examples

```
cut_r(193)
```

depress	<i>Depression and Coping</i>
---------	------------------------------

Description

A subset of data from a research about depression and coping.

Usage

```
depress
```

Format

depress:

A data frame with 94 rows and 237 columns:

id Participant id

class Class

grade Grade

elite Elite classes

intervene 0 = Control group, 1 = Intervention group

gender 0 = girl, 1 = boy

age Age in year

cope1i1p Cope scale, Time1, Item1, Problem-focused coping, 1 = very seldom, 5 = very often

cope1i3a Cope scale, Time1, Item3, Avoidance coping

cope1i5e cope scale, Time1, Item5, Emotion-focused coping

cope2i1p Cope scale, Time2, Item1, Problem-focused coping

depr1i1 Depression scale, Time1, Item1, 1 = very seldom, 5 = always

ecr1avo ECR-RS scale, Item1, attachment avoidance, 1 = very disagree, 7 = very agree

ecr2anx ECR-RS scale, Item2, attachment anxiety

dm1 Depression, Mean, Time1

pm1 Problem-focused coping, Mean, Time1

em1 Emotion-focused coping, Mean, Time1

am1 Avoidance coping, Mean, Time1

avo Attachment avoidance, Mean

anx Attachment anxiety, Mean

Source

Keng package.

power_lm	<i>Compute the post-hoc power and/or plan the sample size for one or a set of predictors in linear regression</i>
----------	---

Description

Compute the post-hoc power and/or plan the sample size for one or a set of predictors in linear regression

Usage

```
power_lm(PRE = 0.02, PC = 0, PA = 1, power = 0.8, sig.level = 0.05, n = NULL)
```

Arguments

PRE	Proportional Reduction in Error. <i>PRE</i> = The square of partial correlation. Cohen (1988) suggested ≥ 0.02 , ≥ 0.13 , and ≥ 0.26 as cut-off values of PRE for small, medium, and large effect sizes, respectively.
PC	Number of parameters of model C (compact model) without focal predictors of interest.
PA	Number of parameters of model A (augmented model) with focal predictors of interest.
power	Expected statistical power for effects of focal predictors.
sig.level	Expected significance level for effects of focal predictors.
n	The current sample size. If <i>n</i> is given, the post-hoc power would be computed.

Value

A list with 4 items: (1) `post`, the post-hoc F-test, `lambda` (non-centrality parameter), and power for sample size n ; (2) `minimum`, the minimum sample size required for focal predictors to reach the expected statistical power and significance level; (3) `prior`, a `data.frame` including `n_i`, `PC`, `PA`, `df_A_i`, `F_i`, `p_i`, `lambda_i`, `power_i`. `_i` indicates these statistics are the intermediate iterative results. Each row of `prior` presents results for one possible sample size n_i . Given `n_i`, `df_A_i`, `F_i`, `p_i`, `lambda_i` and `power_i` would be computed accordingly. (4) A plot of power against sample size n . The cut-off value of n for expected statistical power and expected significance level `sig.level` is annotated on the plot.

References

Cohen, J. (1988). *Statistical power analysis for the behavioral sciences* (2nd ed.). Routledge.

Examples

```
power_lm()
```

Scale

Scale a vector

Description

Scale a vector

Usage

```
Scale(x, m = NULL, sd = NULL, oadvances = NULL)
```

Arguments

<code>x</code>	The original vector.
<code>m</code>	The expected Mean of the scaled vector.
<code>sd</code>	The expected Standard Deviation (unit) of the scaled vector.
<code>oadvances</code>	The distance the Origin of <code>x</code> advances by.

Details

To scale `x`, its origin, or unit (`sd`), or both, could be changed.

If `m = 0` or `NULL`, and `sd = NULL`, `x` would be mean-centered.

If `m` is a non-zero number, and `sd = NULL`, the mean of `x` would be transformed to `m`.

If `m = 0` or `NULL`, and `sd = 1`, `x` would be standardized to be its z-score with $m = 0$ and $m = 1$.

The standardized score is not necessarily the z-score. If neither `m` nor `sd` is `NULL`, `x` would be standardized to be a vector whose mean and standard deviation would be `m` and `sd`, respectively.

To standardize x , the mean and standard deviation of x are needed and computed, for which the missing values of x are removed if any.

If `oadvances` is not `NULL`, the origin of x will advance with the standard deviation being unchanged. In this case, `Scale()` could be used to pick points in simple slope analysis for moderation models. Note that when `oadvances` is not `NULL`, `m` and `sd` must be `NULL`.

Value

The scaled vector.

Examples

```
(x <- rnorm(10, 5, 2))
# Mean-center x.
Scale(x)
# Transform the mean of x to 3.
Scale(x, m = 3)
# Transform x to its z-score.
Scale(x, sd = 1)
# Standardize x with m = 100 and sd = 15.
Scale(x, m = 100, sd = 15)
# The origin of x advances by 3.
Scale(x, oadvances = 3)
```

test_r

Test r using the t-test and Fisher's z given r and n.

Description

Test r using the t -test and Fisher's z given r and n .

Usage

```
test_r(r, n)
```

Arguments

<code>r</code>	Pearson's correlation.
<code>n</code>	Sample size of r .

Details

To test the significance of the r using one-sample t -test, the SE of the r is determined by the following formula: $SE = \sqrt{(1 - r^2)/(n - 2)}$. Another way is transforming r to Fisher's z using the following formula: $fz = \text{atanh}(r)$ with the SE of fz being $\sqrt{(n - 3)}$. Note that Fisher's z is commonly used to compare two Pearson's correlations from independent samples. Fisher's transformation is presented here only for satisfying the curiosity of users interested in the difference of t -test and Fisher's transformation.

Value

A list including r , t -test of r (SE_r , t , p_r), 95% CI of r based on t -test ($LLCI_r_t$, $ULCI_r_t$), fz (Fisher's z) of r , z -test of Fisher's z (SE_{fz} , z , p_{fz}), and 95% CI of r derived from fz . Note that the returned CI of r may be out of r 's valid range $[-1, 1]$. This "error" is deliberately left to users, who should correct the CI manually when reporting.

Examples

```
test_r(0.2, 193)

# compare the p-values of t-test and Fisher's transformation
for (i in seq(30, 200, 10)) {
  cat(c(
    "n =", i, ", ",
    format(
      abs(test_r(0.2, i)[[1]][4] - test_r(0.2, i)[[2]][4]),
      nsmall = 12, scientific = FALSE),
    fill = TRUE)
  )
}
```

Index

* **datasets**
 depress, 5

calc_PRE, 2
compare_lm, 3
cut_r, 4

depress, 5

power_lm, 6

Scale, 7

test_r, 8