

Package ‘vcmeta’

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Type Package

Title Varying Coefficient Meta-Analysis

Version 1.0.0

Description Implements functions for varying coefficient meta-analysis methods.

These methods do not assume effect size homogeneity. Subgroup effect size comparisons, general linear effect size contrasts, and linear models of effect sizes based on varying coefficient methods can be used to describe effect size heterogeneity. Varying coefficient meta-analysis methods do not require the unrealistic assumptions of the traditional fixed-effect and random-effects meta-analysis methods.

For details, see:

Bonett (2008) <[doi:10.1037/a0012868](https://doi.org/10.1037/a0012868)>,

Bonett (2009) <[doi:10.1037/a0016619](https://doi.org/10.1037/a0016619)>,

Bonett (2010) <[doi:10.1037/a0020142](https://doi.org/10.1037/a0020142)>,

Bonett & Price (2014) <[doi:10.1111/bmsp.12024](https://doi.org/10.1111/bmsp.12024)>,

Bonett & Price (2015) <[doi:10.1111/bmsp.12024](https://doi.org/10.1111/bmsp.12024)>,

Bonett (2020) <[doi:10.1111/bmsp.12189](https://doi.org/10.1111/bmsp.12189)>,

Bonett (2021) <[doi:10.1177/1094428120911088](https://doi.org/10.1177/1094428120911088)>.

URL <https://github.com/dgbonett/vcmeta>

BugReports <https://github.com/dgbonett/vcmeta/issues>

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`ci.fisher`*Fisher confidence interval for any type of correlation.*

Description

This function computes a confidence interval for any type correlation using an estimated correlation and its standard error. This function should be used with the `meta.ave.gen` function when the effect size is a correlation. Use the estimated average correlation and its standard error from `meta.ave.gen` (when the effect size is a correlation) in the `ci.fisher` function to obtain a more accurate confidence interval for the population average correlation.

Usage

```
ci.fisher(alpha, cor, se)
```

Arguments

<code>alpha</code>	alpha value for 1-alpha confidence
<code>cor</code>	estimate of correlation
<code>se</code>	standard error of estimated correlation

Value

A 2-element vector with lower and upper bounds of the confidence interval

Examples

```
ci.fisher(0.05, 0.50, .10)

# Should return:
# [1] 0.2802723 0.6699402
```

`cor.from.t`*Compute Pearson correlation between paired measurements from t statistic*

Description

This function computes the Pearson correlation between paired measurements using a reported paired-samples t statistic and other sample information. This correlation estimate is needed in several functions that analyze mean differences and standardized mean differences in paired-samples studies.

Usage

```
cor.from.t(m1, m2, sd1, sd2, t, n)
```

Arguments

m1	sample mean for measurement 1
m2	sample mean for measurement 2
sd1	sample standard deviation for measurement 1
sd2	sample standard deviation for measurement 2
t	value for paired-samples t-test
n	sample size

Value

Returns the sample Pearson correlation between the two paired measurements

Examples

```
cor.from.t(9.4, 9.8, 1.26, 1.40, 2.27, 30)

# Should return:
# [1] 0.7415209
```

meta.ave.agree	<i>Confidence interval for an average G-index agreement coefficient</i>
----------------	---

Description

Computes the estimate, standard error, and confidence interval for an G-index of agreement from two or more studies. This function assumes that two raters each provide a dichotomous rating to a sample of objects. The G-index of agreement is usually preferred to Cohen's kappa.

Usage

```
meta.ave.agree(alpha, f11, f12, f21, f22, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
f11	vector of frequencies in cell 1,1
f12	vector of frequencies in cell 1,2
f21	vector of frequencies in cell 2,1
f22	vector of frequencies in cell 2,2
bystudy	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If `bystudy` is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

Examples

```
f11 <- c(17, 28, 19)
f12 <- c(43, 56, 49)
f21 <- c(3, 5, 5)
f22 <- c(37, 54, 39)
meta.ave.agree(.05, f11, f12, f21, f22, bystudy = TRUE)

# Should return:
#           Estimate      SE      LL      UL
# Average 0.08657934 0.05312584 -0.01754538 0.1907041
# Study 1 0.07692308 0.09776752 -0.11469775 0.2685439
# Study 2 0.14285714 0.08163265 -0.01713992 0.3028542
# Study 3 0.03448276 0.09279245 -0.14738711 0.2163526
```

meta.ave.cor

Confidence interval for an average Pearson or partial correlation

Description

Computes the estimate, standard error, and confidence interval for an average Pearson or partial correlation. The sample correlations must be all Pearson correlations or all partial correlations. Use the `meta.ave.gen` function to meta-analysis any combination of Pearson, partial, or Spearman correlations.

Usage

```
meta.ave.cor(alpha, n, cor, q, bystudy = TRUE)
```

Arguments

<code>alpha</code>	alpha level for 1-alpha confidence
<code>n</code>	vector of sample sizes
<code>cor</code>	vector of sample correlations
<code>q</code>	number of control variables
<code>bystudy</code>	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If `bystudy` is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2008). "Meta-analytic interval estimation for bivariate correlations." *Psychological Methods*, **13**(3), 173–181. ISSN 1939-1463, doi: [10.1037/a0012868](https://doi.org/10.1037/a0012868), <https://doi.org/getdoi.cfm?doi=10.1037/a0012868>.

Examples

```
n <- c(55, 190, 65, 35)
cor <- c(.40, .65, .60, .45)
meta.ave.cor(.05, n, cor, 0, bystudy = TRUE)

# Should return:
#      Estimate      SE      LL      UL
# Average  0.525 0.05113361 0.4176678 0.6178816
# Study 1  0.400 0.11430952 0.1506943 0.6014699
# Study 2  0.650 0.04200694 0.5594086 0.7252465
# Study 3  0.600 0.08000000 0.4171458 0.7361686
# Study 4  0.450 0.13677012 0.1373507 0.6811071
```

meta.ave.cronbach *Confidence interval for an average Cronbach alpha reliability*

Description

Computes the estimate, standard error, and confidence interval for an average Cronbach reliability coefficient from two or more studies.

Usage

```
meta.ave.cronbach(alpha, n, rel, q, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
n	vector of sample sizes
rel	vector of sample reliabilities
q	number of measurements (e.g., items) used to compute each reliability
bystudy	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If `bystudy` is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2010). "Varying coefficient meta-analytic methods for alpha reliability." *Psychological Methods*, **15**(4), 368–385. ISSN 1939-1463, doi: [10.1037/a0020142](https://doi.org/10.1037/a0020142), [https://doi.org/getdoi.cfm?doi=10.1037/a0020142](https://doi.org/10.1037/a0020142).

Examples

```
n <- c(583, 470, 546, 680)
rel <- c(.91, .89, .90, .89)
meta.ave.cronbach(.05, n, rel, 10, bystudy = TRUE)

# Should return:
#           Estimate      SE      LL      UL
# Average  0.8975 0.003256081 0.8911102 0.9038592
# Study 1  0.9100 0.005566064 0.8985763 0.9204108
# Study 2  0.8900 0.007579900 0.8743616 0.9041013
# Study 3  0.9000 0.006391375 0.8868623 0.9119356
# Study 4  0.8900 0.006297549 0.8771189 0.9018203
```

meta.ave.gen

Confidence interval for an average of any parameter

Description

Computes the estimate, standard error, and confidence interval for an average of any type of parameter.

Usage

```
meta.ave.gen(alpha, est, se, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
est	vector of parameter estimates
se	vector of standard errors
bystudy	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If bystudy is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

Examples

```
est <- c(.022, .751, .421, .287, .052, .146, .562, .904)
se <- c(.124, .464, .102, .592, .864, .241, .252, .318)
meta.ave.gen(.05, est, se, bystudy = TRUE)

# Should return:
#      Estimate      SE      LL      UL
# Average 0.393125 0.1561622 0.08705266 0.6991973
# Study 1 0.022000 0.1240000 -0.22103553 0.2650355
# Study 2 0.751000 0.4640000 -0.15842329 1.6604233
# Study 3 0.421000 0.1020000 0.22108367 0.6209163
# Study 4 0.287000 0.5920000 -0.87329868 1.4472987
# Study 5 0.052000 0.8640000 -1.64140888 1.7454089
# Study 6 0.146000 0.2410000 -0.32635132 0.6183513
# Study 7 0.562000 0.2520000 0.06808908 1.0559109
# Study 8 0.904000 0.3180000 0.28073145 1.5272685
```

meta.ave.gen.cc	<i>Confidence interval for an average effect size using a constant coefficient model</i>
-----------------	--

Description

Computes the estimate, standard error, and confidence interval for a weighted average effect using the constant coefficient (fixed-effect) meta-analysis model. The weighted average estimate will be biased regardless of number of studies or sample size per study and the actual confidence interval coverage probability can be much smaller than the specified confidence level when the true effect sizes are not identical across studies.

Usage

```
meta.ave.gen.cc(alpha, est, se, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
est	vector of parameter estimates
se	vector of standard errors
bystudy	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If `bystudy` is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

- Hedges LV, Olkin I (1985). *Statistical methods for meta-analysis*. Academic Press, New York. ISBN 01-233-63802.
- Borenstein M, Hedges LV, Higgins JP, Rothstein HR (2009). *Introduction to meta-analysis*. Wiley, New York.

Examples

```
est <- c(.022, .751, .421, .287, .052, .146, .562, .904)
se <- c(.124, .464, .102, .592, .864, .241, .252, .318)
meta.ave.gen.cc(.05, est, se, bystudy = TRUE)
```

```
# Should return:
```

```
#           Estimate      SE      LL      UL
# Average   0.3127916 0.06854394 0.17844794 0.4471352
# Study 1   0.0220000 0.12400000 -0.22103553 0.2650355
# Study 2   0.7510000 0.46400000 -0.15842329 1.6604233
# Study 3   0.4210000 0.10200000 0.22108367 0.6209163
# Study 4   0.2870000 0.59200000 -0.87329868 1.4472987
# Study 5   0.0520000 0.86400000 -1.64140888 1.7454089
# Study 6   0.1460000 0.24100000 -0.32635132 0.6183513
# Study 7   0.5620000 0.25200000 0.06808908 1.0559109
# Study 8   0.9040000 0.31800000 0.28073145 1.5272685
```

```
meta.ave.gen.rc
```

Confidence interval for an average effect size using a random coefficient model

Description

Computes the estimate, standard error, and confidence interval for a a weighted average effect using the random coefficient (random-effects) meta-analysis model. An estimate of effect-size heterogeneity (tau-squared) is also computed. The random coefficient model assumes that the studies in the meta-analysis are a random sample from some definable superpopulation of studies. This assumption is difficult to justify. The weighted average estimate will be biased regardless of number of studies or sample size per study and the actual confidence interval coverage probability can much smaller than the specified confidence level if the effect sizes are correlated with the weights. This method also assume that the true effects sizes in the superpopulation of studies have a normal distribution. A large number of studies, each with a large sample size, is required to assess the superpopulation normality assumption. The traditional confidence interval for the population tau-squared is hypersensitive to very minor and difficult to detect violations of the superpopulation normality assumption.

Usage

```
meta.ave.gen.rc(alpha, est, se, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
est	vector of parameter estimates
se	vector of standard errors
bystudy	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If `bystudy` is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

- Hedges LV, Olkin I (1985). *Statistical methods for meta-analysis*. Academic Press, New York. ISBN 01-233-63802.
- Borenstein M, Hedges LV, Higgins JP, Rothstein HR (2009). *Introduction to meta-analysis*. Wiley, New York.

Examples

```
est <- c(.022, .751, .421, .287, .052, .146, .562, .904)
se <- c(.124, .464, .102, .592, .864, .241, .252, .318)
meta.ave.gen.rc(.05, est, se, bystudy = TRUE)
```

```
# Should return:
#           Estimate      SE      LL      UL
# Tau-squared 0.03772628 0.0518109 0.00000000 0.1392738
# Average     0.35394806 0.1155239 0.12752528 0.5803708
# Study 1     0.02200000 0.1240000 -0.22103553 0.2650355
# Study 2     0.75100000 0.4640000 -0.15842329 1.6604233
# Study 3     0.42100000 0.1020000 0.22108367 0.6209163
# Study 4     0.28700000 0.5920000 -0.87329868 1.4472987
# Study 5     0.05200000 0.8640000 -1.64140888 1.7454089
# Study 6     0.14600000 0.2410000 -0.32635132 0.6183513
# Study 7     0.56200000 0.2520000 0.06808908 1.0559109
# Study 8     0.90400000 0.3180000 0.28073145 1.5272685
```

meta.ave.mean.ps

Confidence interval for an average mean difference from paired-samples studies

Description

Computes the estimate, standard error, and confidence interval for an average mean difference from two or more paired-samples studies. A Satterthwaite adjustment to the degrees of freedom is used to improve the accuracy of the confidence interval for the average effect size. Equal variances within or across studies is not assumed.

Usage

```
meta.ave.mean.ps(alpha, m1, m2, sd1, sd2, cor, n, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for measurement 1
m2	vector of sample means for measurement 2
sd1	vector of sample SDs for measurement 1
sd2	vector of sample SDs for measurement 2
cor	vector of sample correlations for paired measurements
n	vector of sample sizes
bystudy	logical to also return each study estimate (TRUE) or not

Value

A matrix. First row is the overall average estimate. If bystudy is TRUE also returns 1 row per study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- df - degrees of freedom

References

Bonett DG (2009). "Meta-analytic interval estimation for standardized and unstandardized mean differences." *Psychological Methods*, **14**(3), 225–238. ISSN 1939-1463, doi: [10.1037/a0016619](https://doi.org/10.1037/a0016619), <https://doi.org/10.1037/a0016619>.

Examples

```
m1 <- c(53, 60, 53, 57)
m2 <- c(55, 62, 58, 61)
sd1 <- c(4.1, 4.2, 4.5, 4.0)
sd2 <- c(4.2, 4.7, 4.9, 4.8)
cor <- c(.7, .7, .8, .85)
n <- c(30, 50, 30, 70)
meta.ave.mean.ps(.05, m1, m2, sd1, sd2, cor, n, bystudy = TRUE)

# Should return:
#      Estimate      SE      LL      UL      df
# Average  -3.25 0.2471557 -3.739691 -2.7603091 112.347
# Study 1   -2.00 0.5871400 -3.200836 -0.7991639  29.000
# Study 2   -2.00 0.4918130 -2.988335 -1.0116648  49.000
# Study 3   -5.00 0.5471136 -6.118973 -3.8810270  29.000
```

```
# Study 4   -4.00  0.3023716  -4.603215  -3.3967852  69.000
```

meta.ave.mean2	<i>Confidence interval for an average mean difference from 2-group studies</i>
----------------	--

Description

Computes the estimate, standard error, and confidence interval for an average mean difference from two or more 2-group studies. A Satterthwaite adjustment to the degrees of freedom is used to improve the accuracy of the confidence intervals. Equal variances within or across studies is not assumed.

Usage

```
meta.ave.mean2(alpha, m1, m2, sd1, sd2, n1, n2, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for group 1
m2	vector of sample means for group 2
sd1	vector of sample SDs for group 1
sd2	vector of sample SDs for group 2
n1	vector of group 1 sample size
n2	vector of group 2 sample sizes
bystudy	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If bystudy is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- df - degrees of freedom

References

Bonett DG (2009). "Meta-analytic interval estimation for standardized and unstandardized mean differences." *Psychological Methods*, **14**(3), 225–238. ISSN 1939-1463, doi: [10.1037/a0016619](https://doi.org/10.1037/a0016619), <https://doi.apa.org/getdoi.cfm?doi=10.1037/a0016619>.

Examples

```

m1 <- c(7.4, 6.9)
m2 <- c(6.3, 5.7)
sd1 <- c(1.72, 1.53)
sd2 <- c(2.35, 2.04)
n1 <- c(40, 60)
n2 <- c(40, 60)
meta.ave.mean2(.05, m1, m2, sd1, sd2, n1, n2, bystudy = TRUE)

# Should return:
#           Estimate      SE      LL      UL      df
# Average    1.15 0.2830183 0.5904369 1.709563 139.41053
# Study 1    1.10 0.4604590 0.1819748 2.018025  71.46729
# Study 2    1.20 0.3292036 0.5475574 1.852443 109.42136

```

meta.ave.meanratio.ps *Confidence interval for an average mean ratio from paired-samples studies*

Description

Computes the estimate, standard error, and confidence interval for an geometric average mean ratio from two or more paired-samples studies. A Satterthwaite adjustment to the degrees of freedom is used to improve the accuracy of the confidence interval for the average effect size. Equal variances within or across studies is not assumed.

Usage

```
meta.ave.meanratio.ps(alpha, m1, m2, sd1, sd2, cor, n, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for measurement 1
m2	vector of sample means for measurement 2
sd1	vector of sample SDs for measurement 1
sd2	vector of sample SDs for measurement 2
cor	vector of sample correlations for paired measurements
n	vector of sample sizes
bystudy	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If `bystudy` is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- exp(Estimate) - the exponentiated estimate (the geometric mean)
- exp(LL) - lower limit of the exponentiated confidence interval
- exp(UL) - upper limit of the exponentiated confidence interval
- df - degrees of freedom

Examples

```
m1 <- c(53, 60, 53, 57)
m2 <- c(55, 62, 58, 61)
sd1 <- c(4.1, 4.2, 4.5, 4.0)
sd2 <- c(4.2, 4.7, 4.9, 4.8)
cor <- c(.7, .7, .8, .85)
n <- c(30, 50, 30, 70)
meta.ave.meanratio.ps(.05, m1, m2, sd1, sd2, cor, n, bystudy = TRUE)

# Should return:
#           Estimate          SE          LL          UL
# Average -0.05695120 0.004350863 -0.06558008 -0.04832231
# Study 1 -0.03704127 0.010871086 -0.05927514 -0.01480740
# Study 2 -0.03278982 0.008021952 -0.04891054 -0.01666911
# Study 3 -0.09015110 0.009779919 -0.11015328 -0.07014892
# Study 4 -0.06782260 0.004970015 -0.07773750 -0.05790769
#           exp(Estimate) exp(LL) exp(UL)      df
# Average    0.9446402 0.9365240 0.9528266 103.0256
# Study 1    0.9636364 0.9424474 0.9853017  29.0000
# Study 2    0.9677419 0.9522663 0.9834691  49.0000
# Study 3    0.9137931 0.8956968 0.9322550  29.0000
# Study 4    0.9344262 0.9252073 0.9437371  69.0000
```

meta.ave.meanratio2 *Confidence interval for an average mean ratio from 2-group studies*

Description

Computes the estimate, standard error, and confidence interval for an geometric average mean ratio from two or more 2-group studies. A Satterthwaite adjustment to the degrees of freedom is used to improve the accuracy of the confidence intervals. Equal variances within or across studies is not assumed.

Usage

```
meta.ave.meanratio2(alpha, m1, m2, sd1, sd2, n1, n2, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for group 1
m2	vector of sample means for group 2
sd1	vector of sample SDs for group 1
sd2	vector of sample SDs for group 2
n1	vector of group 1 sample size
n2	vector of group 2 sample sizes
bystudy	logical to also return each study estimate (TRUE) or not

Value

A matrix. First row is the overall average estimate. If bystudy is TRUE also returns 1 row per study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- exp(Estimate) - the exponentiated estimate (the geometric mean)
- exp(LL) - lower limit of the exponentiated confidence interval
- exp(UL) - upper limit of the exponentiated confidence interval
- df - degrees of freedom

References

Bonett DG, Price RM (2020). "Confidence intervals for ratios of means and medians." *Journal of Educational and Behavioral Statistics*, **45**(6), 750–770. ISSN 1076-9986, doi: [10.3102/1076998620934125](https://doi.org/10.3102/1076998620934125), <https://journals.sagepub.com/doi/10.3102/1076998620934125>.

Examples

```
m1 <- c(7.4, 6.9)
m2 <- c(6.3, 5.7)
sd1 <- c(1.7, 1.5)
sd2 <- c(2.3, 2.0)
n1 <- c(40, 20)
n2 <- c(40, 20)
meta.ave.meanratio2(.05, m1, m2, sd1, sd2, n1, n2, bystudy = TRUE)

# Should return:
#      Estimate      SE      LL      UL exp(Estimate)
```

```
# Average 0.1759928 0.05738065 0.061437186 0.2905484 1.192429
# Study 1 0.1609304 0.06820167 0.024749712 0.2971110 1.174603
# Study 2 0.1910552 0.09229675 0.002986265 0.3791242 1.210526
#      exp(LL)  exp(UL)      df
# Average 1.063364 1.337161 66.26499
# Study 1 1.025059 1.345965 65.69929
# Study 2 1.002991 1.461004 31.71341
```

meta.ave.odds

Confidence interval for average odds ratio from 2-group studies

Description

Computes the estimate, standard error, and confidence interval for a geometric mean odds ratio from two or more studies.

Usage

```
meta.ave.odds(alpha, f1, f2, n1, n2, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
f1	vector of group 1 event counts
f2	vector of group 2 event counts
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
bystudy	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If bystudy is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- exp(Estimate) - the exponentiated estimate
- exp(LL) - lower limit of the exponentiated confidence interval
- exp(UL) - upper limit of the exponentiated confidence interval

References

Bonett DG, Price RM (2015). “Varying coefficient meta-analysis methods for odds ratios and risk ratios.” *Psychological Methods*, **20**(3), 394–406. ISSN 1939-1463, doi: [10.1037/met0000032](https://doi.org/10.1037/met0000032), <https://doi.org/10.1037/met0000032>.

Examples

```
n1 <- c(204, 201, 932, 130, 77)
n2 <- c(106, 103, 415, 132, 83)
f1 <- c(24, 40, 93, 14, 5)
f2 <- c(12, 9, 28, 3, 1)
meta.ave.odds(.05, f1, f2, n1, n2, bystudy = TRUE)

# Should return:
#           Estimate      SE      LL      UL
# Average 0.86211102 0.2512852 0.36960107 1.3546210
# Study 1 0.02581353 0.3700520 -0.69947512 0.7511022
# Study 2 0.91410487 0.3830515 0.16333766 1.6648721
# Study 3 0.41496672 0.2226089 -0.02133877 0.8512722
# Study 4 1.52717529 0.6090858 0.33338907 2.7209615
# Study 5 1.42849472 0.9350931 -0.40425414 3.2612436
#           exp(Estimate) exp(LL) exp(UL)
# Average      2.368155 1.4471572 3.875292
# Study 1      1.026150 0.4968460 2.119335
# Study 2      2.494541 1.1774342 5.284997
# Study 3      1.514320 0.9788873 2.342625
# Study 4      4.605150 1.3956902 15.194925
# Study 5      4.172414 0.6674745 26.081952
```

meta.ave.path	<i>Confidence interval for an average slope coefficient in a general linear model or a path model.</i>
---------------	--

Description

Computes the estimate, standard error, and confidence interval for an average slope coefficient from a general linear model (ANOVA, ANCOVA, multiple regression) or a path model.

Usage

```
meta.ave.path(alpha, n, slope, se, s, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
n	vector of sample sizes

slope	vector of slope estimates
se	vector of slope standard errors
s	number of predictors of the response variable
bystudy	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If `bystudy` is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

Examples

```
n <- c(75, 85, 250, 160)
slope <- c(1.57, 1.38, 1.08, 1.25)
se <- c(.658, .724, .307, .493)
meta.ave.path(.05, n, slope, se, 2, bystudy = TRUE)

# Should return:
#      Estimate      SE      LL      UL      df
# Average    1.32 0.2844334 0.75994528 1.880055 263.1837
# Study 1    1.57 0.6580000 0.25830097 2.881699  72.0000
# Study 2    1.38 0.7240000 -0.06026664 2.820267  82.0000
# Study 3    1.08 0.3070000 0.47532827 1.684672 247.0000
# Study 4    1.25 0.4930000 0.27623174 2.223768 157.0000
```

meta.ave.pbcor

Confidence interval for an average point-biserial correlation

Description

Computes the estimate, standard error, and confidence interval for an average point-biserial correlation from two or more studies. Two types of point-biserial correlations can be meta-analyzed. One type uses an unweighted variance and is appropriate in 2-group experimental designs. The other type uses a weighted variance and is appropriate in 2-group nonexperimental designs with simple random sampling within each group. This function requires all point-biserial correlations to be of the same type. Use the `meta.ave.gen` function to meta-analyze any combination of biserial correlation types.

Usage

```
meta.ave.pbcor(alpha, m1, m2, sd1, sd2, n1, n2, type, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for group 1
m2	vector of sample means for group 2
sd1	vector of sample SDs for group 1
sd2	vector of sample SDs for group 2
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
type	<ul style="list-style-type: none"> • set to 1 for weighted variance • set to 2 for unweighted variance
bystudy	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If bystudy is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2020). "Point-biserial correlation: Interval estimation, hypothesis testing, meta-analysis, and sample size determination." *British Journal of Mathematical and Statistical Psychology*, **73**(S1), 113–144. ISSN 0007-1102, doi: [10.1111/bmsp.12189](https://doi.org/10.1111/bmsp.12189), <https://onlinelibrary.wiley.com/doi/10.1111/bmsp.12189>.

Examples

```
m1 <- c(21.9, 23.1, 19.8)
m2 <- c(16.1, 17.4, 15.0)
sd1 <- c(3.82, 3.95, 3.67)
sd2 <- c(3.21, 3.30, 3.02)
n1 <- c(40, 30, 24)
n2 <- c(40, 28, 25)
meta.ave.pbcor(.05, m1, m2, sd1, sd2, n1, n2, 2, bystudy = TRUE)

# Should return:
#      Estimate      SE      LL      UL
# Average 0.6159094 0.04363432 0.5230976 0.6942842
# Study 1 0.6349786 0.06316796 0.4842098 0.7370220
```

```
# Study 2 0.6160553 0.07776700 0.4255342 0.7380898
# Study 3 0.5966942 0.08424778 0.3903883 0.7283966
```

 meta.ave.plot

Forest plot for average effect sizes

Description

Generates a forest plot to visualize effect sizes estimates and overall averages from the meta.ave functions in vcmeta. If the column exp(Estimate) is present, this function plots the exponentiated effect size and CI found in columns exp(Estimate), exp(LL), and exp(UL). Otherwise, this function plots the effect size and CI found in the columns Estimate, LL, and UL.

Usage

```
meta.ave.plot(
  result,
  reference_line = NULL,
  diamond_height = 0.2,
  ggtheme = ggplot2::theme_classic()
)
```

Arguments

result	• a result matrix from any of the replicate functions in vcmeta
reference_line	Optional x-value for a reference line. Only applies if focus is 'Difference' or 'Both'. Defaults to NULL, in which case a reference line is not drawn.
diamond_height	• Optional height of the diamond representing average effect size. Only applies if focus is 'Average' or 'Both'. Defaults to 0.2
ggtheme	• optional ggplot2 theme object; defaults to theme_classic()

Value

Returns a ggplot object. If stored, can be further customized via the ggplot API

Examples

```
# Plot results from meta.ave.mean2
m1 <- c(7.4, 6.9)
m2 <- c(6.3, 5.7)
sd1 <- c(1.72, 1.53)
sd2 <- c(2.35, 2.04)
n1 <- c(40, 60)
n2 <- c(40, 60)
result <- meta.ave.mean2(.05, m1, m2, sd1, sd2, n1, n2, bystudy = TRUE)
meta.ave.plot(result, reference_line = 0)
```

```

# Plot results from meta.ave.meanratio2
# Note that this plots the exponentiated effect size and CI
m1 <- c(53, 60, 53, 57)
m2 <- c(55, 62, 58, 61)
sd1 <- c(4.1, 4.2, 4.5, 4.0)
sd2 <- c(4.2, 4.7, 4.9, 4.8)
cor <- c(.7, .7, .8, .85)
n <- c(30, 50, 30, 70)
result <- meta.ave.meanratio.ps(.05, m1, m2, sd1, sd2, cor, n, bystudy = TRUE)
myplot <- meta.ave.plot(result, reference_line = 1)
myplot

# Change x-scale to log2
library(ggplot2)
myplot <- myplot + scale_x_continuous(
  trans = 'log2',
  limits = c(0.75, 1.25),
  name = "Estimated Ratio of Means, Log2 Scale"
)
myplot

```

meta.ave.prop.ps	<i>Confidence interval for an average proportion difference in paired-samples studies</i>
------------------	---

Description

Computes the estimate, standard error, and confidence interval for an average proportion difference from two or more studies.

Usage

```
meta.ave.prop.ps(alpha, f11, f12, f21, f22, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
f11	vector of frequencies in cell 1,1
f12	vector of frequencies in cell 1,2
f21	vector of frequencies in cell 2,1
f22	vector of frequencies in cell 2,2
bystudy	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If `bystudy` is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG, Price RM (2014). "Meta-analysis methods for risk differences." *British Journal of Mathematical and Statistical Psychology*, **67**(3), 371–387. ISSN 00071102, doi: [10.1111/bmsp.12024](https://onlinelibrary.wiley.com/doi/10.1111/bmsp.12024), <https://onlinelibrary.wiley.com/doi/10.1111/bmsp.12024>.

Examples

```
f11 <- c(17, 28, 19)
f12 <- c(43, 56, 49)
f21 <- c(3, 5, 5)
f22 <- c(37, 54, 39)
meta.ave.prop.ps(.05, f11, f12, f21, f22, bystudy = TRUE)

# Should return:
#      Estimate      SE      LL      UL
# Average 0.3809573 0.03000016 0.3221581 0.4397565
# Study 1 0.3921569 0.05573055 0.2829270 0.5013867
# Study 2 0.3517241 0.04629537 0.2609869 0.4424614
# Study 3 0.3859649 0.05479300 0.2785726 0.4933572
```

meta.ave.prop2	<i>Confidence interval for an average proportion difference in 2-group studies</i>
----------------	--

Description

Computes the estimate, standard error, and confidence interval for an average proportion difference from two or more studies.

Usage

```
meta.ave.prop2(alpha, f1, f2, n1, n2, bystudy = TRUE)
```


Arguments

alpha	alpha level for 1-alpha confidence
f1	vector of group 1 event counts
f2	vector of group 2 event counts
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
bystudy	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If `bystudy` is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG, Price RM (2014). "Meta-analysis methods for risk differences." *British Journal of Mathematical and Statistical Psychology*, **67**(3), 371–387. ISSN 00071102, doi: [10.1111/bmsp.12024](https://onlinelibrary.wiley.com/doi/10.1111/bmsp.12024), <https://onlinelibrary.wiley.com/doi/10.1111/bmsp.12024>.

Examples

```
n1 <- c(204, 201, 932, 130, 77)
n2 <- c(106, 103, 415, 132, 83)
f1 <- c(24, 40, 93, 14, 5)
f2 <- c(12, 9, 28, 3, 1)
meta.ave.prop2(.05, f1, f2, n1, n2, bystudy = TRUE)

# Should return:
#           Estimate      SE      LL      UL
# Average 0.0567907589 0.01441216 2.854345e-02 0.08503807
# Study 1 0.0009888529 0.03870413 -7.486985e-02 0.07684756
# Study 2 0.1067323481 0.04018243 2.797623e-02 0.18548847
# Study 3 0.0310980338 0.01587717 -2.064379e-05 0.06221671
# Study 4 0.0837856174 0.03129171 2.245499e-02 0.14511624
# Study 5 0.0524199553 0.03403926 -1.429577e-02 0.11913568
```

meta.ave.propratio2 *Confidence interval for an average proportion ratio from 2-group studies*

Description

Computes the estimate, standard error, and confidence interval for a geometric mean proportion ratio from two or more studies.

Usage

```
meta.ave.propratio2(alpha, f1, f2, n1, n2, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
f1	vector of group 1 event counts
f2	vector of group 2 event counts
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
bystudy	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If bystudy is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- exp(Estimate) - the exponentiated estimate
- exp(LL) - lower limit of the exponentiated confidence interval
- exp(UL) - upper limit of the exponentiated confidence interval

References

Price RM, Bonett DG (2008). "Confidence intervals for a ratio of two independent binomial proportions." *Statistics in Medicine*, **27**(26), 5497–5508. ISSN 02776715, doi: [10.1002/sim.3376](https://doi.org/10.1002/sim.3376), <https://onlinelibrary.wiley.com/doi/10.1002/sim.3376>.

Examples

```

n1 <- c(204, 201, 932, 130, 77)
n2 <- c(106, 103, 415, 132, 83)
f1 <- c(24, 40, 93, 14, 5)
f2 <- c(12, 9, 28, 3, 1)
meta.ave.propratio2(.05, f1, f2, n1, n2, bystudy = TRUE)

# Should return:
#           Estimate      SE      LL      UL
# Average 0.84705608 0.2528742 0.35143178 1.3426804
# Study 1 0.03604257 0.3297404 -0.61023681 0.6823220
# Study 2 0.81008932 0.3442007 0.13546839 1.4847103
# Study 3 0.38746839 0.2065227 -0.01730864 0.7922454
# Study 4 1.49316811 0.6023296 0.31262374 2.6737125
# Study 5 1.50851199 0.9828420 -0.41782290 3.4348469
#   exp(Estimate)  exp(LL)  exp(UL)
# Average 2.332769 1.4211008 3.829294
# Study 1 1.036700 0.5432222 1.978466
# Study 2 2.248109 1.1450730 4.413686
# Study 3 1.473246 0.9828403 2.208350
# Study 4 4.451175 1.3670071 14.493677
# Study 5 4.520000 0.6584788 31.026662

```

meta.ave.semipart	<i>Confidence interval for an average semipartial correlation</i>
-------------------	---

Description

Computes the estimate, standard error, and confidence interval for an average semipartial correlation from two or more studies.

Usage

```
meta.ave.semipart(alpha, n, cor, r2, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
n	vector of sample sizes
cor	vector of sample semipartial correlations
r2	vector of squared multiple correlations for full model
bystudy	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If `bystudy` is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

Examples

```
n <- c(128, 97, 210, 217)
cor <- c(.35, .41, .44, .39)
r2 <- c(.29, .33, .36, .39)
meta.ave.semipart(.05, n, cor, r2, bystudy = TRUE)

# Should return:
#      Estimate      SE      LL      UL
# Average  0.3975 0.03221240 0.3325507 0.4586965
# Study 1  0.3500 0.07175200 0.2023485 0.4820930
# Study 2  0.4100 0.07886080 0.2447442 0.5521076
# Study 3  0.4400 0.05146694 0.3338366 0.5351410
# Study 4  0.3900 0.05085271 0.2860431 0.4848830
```

meta.ave.slope

Confidence interval for an average slope coefficient

Description

Computes the estimate, standard error, and confidence interval for an average slope coefficient in a simple linear regression model. A Satterthwaite adjustment to the degrees of freedom is used to improve the accuracy of the confidence interval.

Usage

```
meta.ave.slope(alpha, n, cor, sdy, sdx, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
n	vector of sample sizes
cor	vector of sample correlations
sdy	vector of SDs of y
sdx	vector of SDs of x
bystudy	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If `bystudy` is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- df - degrees of freedom

Examples

```
n <- c(45, 85, 50, 60)
cor <- c(.24, .35, .16, .20)
sdy <- c(12.2, 14.1, 11.7, 15.9)
sdX <- c(1.34, 1.87, 2.02, 2.37)
meta.ave.slope(.05, n, cor, sdy, sdX, bystudy = TRUE)

# Should return:
#      Estimate      SE      LL      UL      df
# Average 1.7731542 0.4755417 0.8335021 2.712806 149.4777
# Study 1 2.1850746 1.3084468 -0.4536599 4.823809 43.0000
# Study 2 2.6390374 0.7262491 1.1945573 4.083518 83.0000
# Study 3 0.9267327 0.8146126 -0.7111558 2.564621 48.0000
# Study 4 1.3417722 0.8456799 -0.3510401 3.034584 58.0000
```

meta.ave.spear

Confidence interval for an average Spearman correlation

Description

Computes the estimate, standard error, and confidence interval for an average Spearman correlation. The Spearman correlation is preferred to the Pearson correlation if the relation between the two quantitative variables is monotonic rather than linear or if the bivariate normality assumption is not plausible.

Usage

```
meta.ave.spear(alpha, n, cor, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
n	vector of sample sizes
cor	vector of sample Spearman correlations
bystudy	logical to also return each study estimate (TRUE) or not

Value

Returns a matrix. The first row is the average estimate across all studies. If `bystudy` is true, there is 1 additional row for each study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2008). "Meta-analytic interval estimation for bivariate correlations." *Psychological Methods*, **13**(3), 173–181. ISSN 1939-1463, doi: [10.1037/a0012868](https://doi.org/10.1037/a0012868), <https://doi.org/getdoi.cfm?doi=10.1037/a0012868>.

Examples

```
n <- c(150, 200, 300, 200, 350)
cor <- c(.14, .29, .16, .21, .23)
meta.ave.spear(.05, n, cor, bystudy = TRUE)

# Should return:
#           Estimate      SE      LL      UL
# Average  0.206 0.02944265 0.14763960 0.2629309
# Study 1  0.140 0.08031750 -0.02151639 0.2943944
# Study 2  0.290 0.06492643 0.15476515 0.4145671
# Study 3  0.160 0.05635101 0.04689807 0.2690514
# Study 4  0.210 0.06776195 0.07187439 0.3402225
# Study 5  0.230 0.05069710 0.12690280 0.3281809
```

meta.ave.stdmean.ps *Confidence interval for an average standardized mean difference from paired-samples studies*

Description

Computes the estimate, standard error, and confidence interval for an average standardized mean difference from two or more paired-samples studies. Unweighted variances and single group variance are options for the standardizer. Equal variances within or across studies is not assumed.

Usage

```
meta.ave.stdmean.ps(alpha, m1, m2, sd1, sd2, cor, n, stdzr, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for measurement 1
m2	vector of sample means for measurement 2
sd1	vector of sample SDs for measurement 1
sd2	vector of sample SDs for measurement 2
cor	vector of sample correlations for paired measurements
n	vector of sample sizes
stdzr	set to 0 for square root average variance standardizer set to 1 for measurement 1 SD standardizer set to 2 for measurement 2 SD standardizer
bystudy	logical to also return each study estimate (TRUE) or not

Value

A matrix. First row is the overall average estimate. If bystudy is TRUE also returns 1 row per study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2009). "Meta-analytic interval estimation for standardized and unstandardized mean differences." *Psychological Methods*, **14**(3), 225–238. ISSN 1939-1463, doi: [10.1037/a0016619](https://doi.org/10.1037/a0016619), <https://doi.apa.org/getdoi.cfm?doi=10.1037/a0016619>.

Examples

```
m1 <- c(23.9, 24.1)
m2 <- c(25.1, 26.9)
sd1 <- c(1.76, 1.58)
sd2 <- c(2.01, 1.76)
cor <- c(.78, .84)
n <- c(25, 30)
meta.ave.stdmean.ps(.05, m1, m2, sd1, sd2, cor, n, 1, bystudy = TRUE)

# Should return:
#           Estimate      SE      LL      UL
# Average -1.1931045 0.1568034 -1.500433 -0.8857755
# Study 1 -0.6818182 0.1773785 -1.029474 -0.3341628
# Study 2 -1.7721519 0.2586234 -2.279044 -1.2652594
```

meta.ave.stdmean2 *Confidence interval for an average standardized mean difference from 2-group studies*

Description

Computes the estimate, standard error, and confidence interval for an average standardized mean difference from two or more 2-group studies. Unweighted variances, weighted variances, and single group variance are options for the standardizer. Equal variances within or across studies is not assumed.

Usage

```
meta.ave.stdmean2(alpha, m1, m2, sd1, sd2, n1, n2, stdzr, bystudy = TRUE)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for group 1
m2	vector of sample means for group 2
sd1	vector of sample SDs for group 1
sd2	vector of sample SDs for group 2
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
stdzr	<ul style="list-style-type: none"> • set to 0 for square root average variance standardizer • set to 1 for group 1 SD standardizer • set to 2 for group 2 SD standardizer • set to 3 for square root weighted variance standardizer
bystudy	logical to also return each study estimate (TRUE) or not

Value

A matrix. First row is the overall average estimate. If bystudy is TRUE also returns 1 row per study. The matrix has the following columns:

- Estimate - the estimated effect size
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2009). "Meta-analytic interval estimation for standardized and unstandardized mean differences." *Psychological Methods*, **14**(3), 225–238. ISSN 1939-1463, doi: [10.1037/a0016619](https://doi.org/10.1037/a0016619), <https://doi.apa.org/getdoi.cfm?doi=10.1037/a0016619>.

Examples

```

m1 <- c(21.9, 23.1, 19.8)
m2 <- c(16.1, 17.4, 15.0)
sd1 <- c(3.82, 3.95, 3.67)
sd2 <- c(3.21, 3.30, 3.02)
n1 <- c(40, 30, 24)
n2 <- c(40, 28, 25)
meta.ave.stdmean2(.05, m1, m2, sd1, sd2, n1, n2, 0, bystudy = TRUE)

# Should return:
#      Estimate      SE      LL      UL
# Average 1.526146 0.1734341 1.1862217 1.866071
# Study 1 1.643894 0.2629049 1.1286100 2.159178
# Study 2 1.566132 0.3056278 0.9671126 2.165152
# Study 3 1.428252 0.3289179 0.7835848 2.072919

```

meta.lc.agree

Confidence interval for a linear contrast of G-index coefficients

Description

Computes the estimate, standard error, and confidence interval for a linear contrast of G-index of agreement coefficients from two or more studies. This function assumes that two raters each provide a dichotomous rating for a sample of objects.

Usage

```
meta.lc.agree(alpha, f11, f12, f21, f22, v)
```

Arguments

alpha	alpha level for 1-alpha confidence
f11	vector of frequencies in cell 1,1
f12	vector of frequencies in cell 1,2
f21	vector of frequencies in cell 2,1
f22	vector of frequencies in cell 2,2
v	vector of contrast coefficients

Value

Returns 1-row matrix with the estimate for the contrast. The matrix has the following columns:

- Estimate - estimate of linear contrast
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

Examples

```
f11 <- c(17, 28, 19)
f12 <- c(43, 56, 49)
f21 <- c(3, 5, 5)
f22 <- c(37, 54, 39)
v <- c(.5, .5, -1)
meta.lc.agree(.05, f11, f12, f21, f22, v)

# Should return:
#           Estimate      SE      LL      UL
# Contrast 0.07692783 0.1138407 -0.1461958 0.3000515
```

meta.lc.gen

Confidence interval for a linear contrast of effect sizes

Description

Computes the estimate, standard error, and confidence interval for a linear contrast of any type of effect size from two or more studies.

Usage

```
meta.lc.gen(alpha, est, se, v)
```

Arguments

alpha	alpha level for 1-alpha confidence
est	vector of sample estimates
se	vector of standard errors
v	vector of contrast coefficients

Value

Returns 1-row matrix with the estimate for the contrast. The matrix has the following columns:

- Estimate - estimate of linear contrast
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

Examples

```

est <- c(.55, .59, .44, .48, .26, .19)
se <- c(.054, .098, .029, .084, .104, .065)
v <- c(.5, .5, -.25, -.25, -.25, -.25)
meta.lc.gen(.05, est, se, v)

# Should return:
#           Estimate      SE      LL      UL
# Contrast  0.2275 0.06755461 0.0950954 0.3599046

```

meta.lc.mean.ps	<i>Confidence interval for a linear contrast of mean differences from paired-samples studies</i>
-----------------	--

Description

Computes the estimate, standard error, and confidence interval for a linear contrast of paired-samples mean differences from two or more studies. A Satterthwaite adjustment to the degrees of freedom is used to improve the accuracy of the confidence interval. Equal variances within or across studies is not assumed.

Usage

```
meta.lc.mean.ps(alpha, m1, m2, sd1, sd2, cor, n, v)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for group 1
m2	vector of sample means for group 2
sd1	vector of sample SDs for group 1
sd2	vector of sample SDs for group 2
cor	vector of sample correlations for paired measurements
n	vector of sample sizes
v	vector of contrast coefficients

Value

Returns 1-row matrix with the estimate for the contrast. The matrix has the following columns:

- Estimate - estimate of linear contrast
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- df - degrees of freedom

References

Bonett DG (2009). “Meta-analytic interval estimation for standardized and unstandardized mean differences.” *Psychological Methods*, **14**(3), 225–238. ISSN 1939-1463, doi: [10.1037/a0016619](https://doi.org/10.1037/a0016619), <https://doi.org/10.1037/a0016619>.

Examples

```
m1 <- c(53, 60, 53, 57)
m2 <- c(55, 62, 58, 61)
sd1 <- c(4.1, 4.2, 4.5, 4.0)
sd2 <- c(4.2, 4.7, 4.9, 4.8)
cor <- c(.7, .7, .8, .85)
n <- c(30, 50, 30, 70)
v <- c(.5, .5, -.5, -.5)
meta.lc.mean.ps(.05, m1, m2, sd1, sd2, cor, n, v)

# Should return:
#      Estimate      SE      LL      UL      df
# Contrast      2.5 0.4943114 1.520618 3.479382 112.347
```

meta.lc.mean1

Confidence interval a for a linear contrast of means

Description

Computes the estimate, standard error, and confidence interval for a linear contrast of means from two or more studies. This function will use either an unequal variance (recommended) or an equal variance method. A Satterthwaite adjustment to the degrees of is used with the unequal variance method.

Usage

```
meta.lc.mean1(alpha, m, sd, n, v, eqvar = FALSE)
```

Arguments

alpha	alpha level for 1-alpha confidence
m	vector of sample means
sd	vector of sample standard deviations
n	vector of sample sizes
v	vector of contrast coefficients
eqvar	<ul style="list-style-type: none"> • FALSE for unequal variance method • TRUE for equal variance method

Value

Returns 1-row matrix with the estimate for the contrast. The matrix has the following columns:

- Estimate - estimate of linear contrast
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- df - degrees of freedom

References

Snedecor GW, Cochran WG (1980). *Statistical methods*, 7th edition. ISU University Pres, Ames, Iowa.

Examples

```
m <- c(33.5, 37.9, 38.0, 44.1)
sd <- c(3.84, 3.84, 3.65, 4.98)
n <- c(10, 10, 10, 10)
v <- c(.5, .5, -.5, -.5)
meta.lc.mean1(.05, m, sd, n, v, eqvar = FALSE)

# Should return:
#      Estimate      SE      LL      UL      df
# Contrast    -5.35 1.300136 -7.993583 -2.706417 33.52169
```

meta.lc.mean2	<i>Confidence interval for a linear contrast of mean differences from 2-group studies</i>
---------------	---

Description

Computes the estimate, standard error, and confidence interval for a linear contrast of 2-group mean differences from two or more studies. A Satterthwaite adjustment to the degrees of freedom is used to improve the accuracy of the confidence interval. Equal variances within or across studies is not assumed.

Usage

```
meta.lc.mean2(alpha, m1, m2, sd1, sd2, n1, n2, v)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for group 1
m2	vector of sample means for group 2
sd1	vector of sample SDs for group 1
sd2	vector of sample SDs for group 2
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
v	vector of contrast coefficients

Value

Returns 1-row matrix with the estimate for the contrast. The matrix has the following columns:

- Estimate - estimate of the linear contrast
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- df - degrees of freedom

References

Bonett DG (2009). "Meta-analytic interval estimation for standardized and unstandardized mean differences." *Psychological Methods*, **14**(3), 225–238. ISSN 1939-1463, doi: [10.1037/a0016619](https://doi.org/10.1037/a0016619), <https://doi.org/10.1037/a0016619>.

Examples

```
m1 <- c(45.1, 39.2, 36.3, 34.5)
m2 <- c(30.0, 35.1, 35.3, 36.2)
sd1 <- c(10.7, 10.5, 9.4, 11.5)
sd2 <- c(12.3, 12.0, 10.4, 9.6)
n1 <- c(40, 20, 50, 25)
n2 <- c(40, 20, 48, 26)
v <- c(.5, .5, -.5, -.5)
meta.lc.mean2(.05, m1, m2, sd1, sd2, n1, n2, v)

# Should return:
#      Estimate      SE      LL      UL      df
# Contrast      9.95 2.837787 4.343938 15.55606 153.8362
```

meta.lc.meanratio.ps *Confidence interval for a log-linear contrast of mean ratios from paired-samples studies*

Description

Computes the estimate, standard error, and confidence interval for a log-linear contrast of paired-sample mean ratios from two or more studies. A Satterthwaite adjustment to the degrees of freedom is used to improve the accuracy of the confidence interval. Equal variances within or across studies is not assumed.

Usage

```
meta.lc.meanratio.ps(alpha, m1, m2, sd1, sd2, cor, n, v)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for group 1
m2	vector of sample means for group 2
sd1	vector of sample SDs for group 1
sd2	vector of sample SDs for group 2
cor	vector of sample correlations for paired measurements
n	vector of sample sizes
v	vector of contrast coefficients

Value

Returns 1-row matrix with the estimate for the contrast. The matrix has the following columns:

- Estimate - estimate of log-linear contrast
- SE - standard error of log-linear contrast
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- exp(Estimate) - exponentiated estimate of log-linear function
- exp(LL) - lower limit of the exponentiated confidence interval
- exp(UL) - upper limit of the exponentiated confidence interval
- df - degrees of freedom

References

Bonett DG, Price RM (2020). "Confidence intervals for ratios of means and medians." *Journal of Educational and Behavioral Statistics*, **45**(6), 750–770. ISSN 1076-9986, doi: [10.3102/1076998620934125](https://doi.org/10.3102/1076998620934125), <https://journals.sagepub.com/doi/10.3102/1076998620934125>.

Examples

```

m1 <- c(53, 60, 53, 57)
m2 <- c(55, 62, 58, 61)
sd1 <- c(4.1, 4.2, 4.5, 4.0)
sd2 <- c(4.2, 4.7, 4.9, 4.8)
cor <- c(.7, .7, .8, .85)
n <- c(30, 50, 30, 70)
v <- c(.5, .5, -.5, -.5)
meta.lc.meanratio.ps(.05, m1, m2, sd1, sd2, cor, n, v)

# Should return:
#           Estimate          SE          LL          UL exp(Estimate)
# Contrast 0.0440713 0.008701725 0.02681353 0.06132907    1.045057
#           exp(LL) exp(UL)          df
# Contrast 1.027176 1.063249 103.0256

```

meta.lc.meanratio2 *Confidence interval for a log-linear contrast of mean ratios from 2-group studies*

Description

Computes the estimate, standard error, and confidence interval for a log-linear contrast of 2-group mean ratio from two or more studies. A Satterthwaite adjustment to the degrees of freedom is used to improve the accuracy of the confidence interval. Equal variances within or across studies is not assumed.

Usage

```
meta.lc.meanratio2(alpha, m1, m2, sd1, sd2, n1, n2, v)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for group 1
m2	vector of sample means for group 2
sd1	vector of sample SDs for group 1
sd2	vector of sample SDs for group 2
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
v	vector of contrast coefficients

Value

Returns 1-row matrix with the estimate for the contrast. The matrix has the following columns:

- Estimate - estimate of log-linear contrast
- SE - standard error of log-linear contrast
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- exp(Estimate) - the exponentiated estimate
- exp(LL) - lower limit of the exponentiated confidence interval
- exp(UL) - upper limit of the exponentiated confidence interval
- df - degrees of freedom

References

Bonett DG, Price RM (2020). “Confidence intervals for ratios of means and medians.” *Journal of Educational and Behavioral Statistics*, **45**(6), 750–770. ISSN 1076-9986, doi: [10.3102/1076998620934125](https://doi.org/10.3102/1076998620934125), <https://journals.sagepub.com/doi/10.3102/1076998620934125>.

Examples

```
m1 <- c(45.1, 39.2, 36.3, 34.5)
m2 <- c(30.0, 35.1, 35.3, 36.2)
sd1 <- c(10.7, 10.5, 9.4, 11.5)
sd2 <- c(12.3, 12.0, 10.4, 9.6)
n1 <- c(40, 20, 50, 25)
n2 <- c(40, 20, 48, 26)
v <- c(.5, .5, -.5, -.5)
meta.lc.meanratio2(.05, m1, m2, sd1, sd2, n1, n2, v)

# Should return:
#      Estimate      SE      LL      UL  exp(Estimate)
# Contrast 0.2691627 0.07959269 0.1119191 0.4264064      1.308868
#      exp(LL) exp(UL)      df
# Contrast 1.118422 1.531743 152.8665
```

Description

Computes the estimate, standard error, and confidence interval for an exponentiated log-linear contrast of odds ratios from two or more studies.

Usage

```
meta.lc.odds(alpha, f1, f2, n1, n2, v)
```

Arguments

alpha	alpha level for 1-alpha confidence
f1	vector of group 1 event counts
f2	vector of group 2 event counts
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
v	vector of contrast coefficients

Value

Returns 1-row matrix with the estimate for the contrast. The matrix has the following columns:

- Estimate - estimate of log-linear contrast
- SE - standard error of log-linear contrast
- exp(Estimate) - the exponentiated contrast
- exp(LL) - lower limit of the exponentiated confidence interval
- exp(UL) - upper limit of the exponentiated confidence interval

References

Bonett DG, Price RM (2015). “Varying coefficient meta-analysis methods for odds ratios and risk ratios.” *Psychological Methods*, **20**(3), 394–406. ISSN 1939-1463, doi: [10.1037/met0000032](https://doi.org/10.1037/met0000032), <https://doi.org/10.1037/met0000032>.

Examples

```
n1 <- c(50, 150, 150)
f1 <- c(16, 50, 25)
n2 <- c(50, 150, 150)
f2 <- c(7, 15, 20)
v <- c(1, -1, 0)
meta.lc.odds(.05, f1, f2, n1, n2, v)

# Should return:
#           Estimate      SE exp(Estimate) exp(LL) exp(UL)
# Contrast -0.4596883 0.5895438      0.6314805 0.1988563 2.005305
```

meta.lc.prop.ps	<i>Confidence interval for a linear contrast of proportion differences in paired-samples studies</i>
-----------------	--

Description

Computes the estimate, standard error, and confidence interval for a linear contrast of paired-samples group proportion differences from two or more studies.

Usage

```
meta.lc.prop.ps(alpha, f11, f12, f21, f22, v)
```

Arguments

alpha	alpha level for 1-alpha confidence
f11	vector of frequencies in cell 1,1
f12	vector of frequencies in cell 1,2
f21	vector of frequencies in cell 2,1
f22	vector of frequencies in cell 2,2
v	vector of contrast coefficients

Value

Returns 1-row matrix with the estimate for the contrast. The matrix has the following columns:

- Estimate - estimate of linear contrast
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG, Price RM (2014). "Meta-analysis methods for risk differences." *British Journal of Mathematical and Statistical Psychology*, **67**(3), 371–387. ISSN 00071102, doi: [10.1111/bmsp.12024](https://onlinelibrary.wiley.com/doi/10.1111/bmsp.12024), <https://onlinelibrary.wiley.com/doi/10.1111/bmsp.12024>.

Examples

```
f11 <- c(17, 28, 19)
f12 <- c(43, 56, 49)
f21 <- c(3, 5, 5)
f22 <- c(37, 54, 39)
v <- c(.5, .5, -1)
meta.lc.prop.ps(.05, f11, f12, f21, f22, v)
```

```
# Should return:
#           Estimate      SE      LL      UL
# Contrast -0.01436285  0.06511285 -0.1419817  0.113256
```

meta.lc.prop1	<i>Confidence interval for a linear contrast of proportions.</i>
---------------	--

Description

Computes the estimate, standard error, and an adjusted Wald confidence interval for a linear contrast of proportions from two or more studies.

Usage

```
meta.lc.prop1(alpha, f, n, v)
```

Arguments

alpha	alpha level for 1-alpha confidence
f	vector of sample frequency counts
n	vector of sample sizes
v	vector of contrast coefficients

Value

Returns 1-row matrix with the estimate for the contrast. The matrix has the following columns:

- Estimate - estimate of linear contrast
- SE - standard error
- LL - lower limit of the adjusted Wald confidence interval
- UL - upper limit of the adjusted Wald confidence interval

References

Price RM, Bonett DG (2004). "An improved confidence interval for a linear function of binomial proportions." *Computational Statistics and Data Analysis*, **45**(3), 449–456. ISSN 01679473, doi: [10.1016/S01679473\(03\)000070](https://doi.org/10.1016/S01679473(03)000070), [https://doi.org/10.1016/S0167-9473\(03\)00007-0](https://doi.org/10.1016/S0167-9473(03)00007-0).

Examples

```
f <- c(26, 24, 38)
n <- c(60, 60, 60)
v <- c(-.5, -.5, 1)
meta.lc.prop1(.05, f, n, v)

# Should return:
#           Estimate      SE      LL      UL
# Contrast 0.2119565 0.07602892 0.06294259 0.3609705
```

meta.lc.prop2	<i>Confidence interval for a linear contrast of proportion differences in 2-group studies</i>
---------------	---

Description

Computes the estimate, standard error, and confidence interval for a linear contrast of 2-group proportion differences from two or more studies.

Usage

```
meta.lc.prop2(alpha, f1, f2, n1, n2, v)
```

Arguments

alpha	alpha level for 1-alpha confidence
f1	vector of group 1 event counts
f2	vector of group 2 event counts
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
v	vector of contrast coefficients

Value

Returns 1-row matrix with the estimate for the contrast. The matrix has the following columns:

- Estimate - estimate of the linear contrast
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG, Price RM (2014). “Meta-analysis methods for risk differences.” *British Journal of Mathematical and Statistical Psychology*, **67**(3), 371–387. ISSN 00071102, doi: [10.1111/bmsp.12024](https://onlinelibrary.wiley.com/doi/10.1111/bmsp.12024), <https://onlinelibrary.wiley.com/doi/10.1111/bmsp.12024>.

Examples

```
n1 <- c(50, 150, 150)
n2 <- c(50, 150, 150)
f1 <- c(16, 50, 25)
f2 <- c(7, 15, 20)
v <- c(1, -1, 0)
meta.lc.prop2(.05, f1, f2, n1, n2, v)

# Should return:
#           Estimate      SE      LL      UL
# Contrast -0.05466931 0.09401019 -0.2389259 0.1295873
```

meta.lc.propratio2	<i>Confidence interval for a log-linear contrast of proportion ratios from 2-group studies</i>
--------------------	--

Description

Computes the estimate, standard error, and confidence interval for an exponentiated log-linear contrast of 2-group proportion odds ratios from two or more studies.

Usage

```
meta.lc.propratio2(alpha, f1, f2, n1, n2, v)
```

Arguments

alpha	alpha level for 1-alpha confidence
f1	vector of group 1 event counts
f2	vector of group 2 event counts
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
v	vector of contrast coefficients

Value

Returns 1-row matrix with the estimate for the contrast. The matrix has the following columns:

- Estimate - estimate of log-linear contrast
- SE - standard error of log-linear contrast
- exp(Estimate) - the exponentiated contrast
- exp(LL) - lower limit of the exponentiated confidence interval
- exp(UL) - upper limit of the exponentiated confidence interval

References

Price RM, Bonett DG (2008). “Confidence intervals for a ratio of two independent binomial proportions.” *Statistics in Medicine*, **27**(26), 5497–5508. ISSN 02776715, doi: [10.1002/sim.3376](https://onlinelibrary.wiley.com/doi/10.1002/sim.3376), <https://onlinelibrary.wiley.com/doi/10.1002/sim.3376>.

Examples

```
n1 <- c(50, 150, 150)
f1 <- c(16, 50, 25)
n2 <- c(50, 150, 150)
f2 <- c(7, 15, 20)
v <- c(1, -1, 0)
meta.lc.propratio2(.05, f1, f2, n1, n2, v)

# Should return:
#           Estimate      SE exp(Estimate) exp(LL) exp(UL)
# Contrast -0.3853396 0.4828218      0.6802196 0.2640405 1.752378
```

meta.lc.stdmean.ps *Confidence interval for a linear contrast of standardized mean differences from paired-samples studies*

Description

Computes the estimate, standard error, and confidence interval for a linear contrast of paired-samples standardized mean differences from two or more studies. Equal variances within or across studies is not assumed. This function has an option to use a square root average variance standardizer or a single group standard deviation standardizer.

Usage

```
meta.lc.stdmean.ps(alpha, m1, m2, sd1, sd2, cor, n, v, stdzr)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for group 1
m2	vector of sample means for group 2
sd1	vector of sample SDs for group 1
sd2	vector of sample SDs for group 2
cor	vector of sample correlations for paired measurements
n	vector of sample sizes
v	vector of contrast coefficients
stdzr	<ul style="list-style-type: none"> • set to 0 for square root average variance standardizer • set to 1 for group 1 SD standardizer • set to 2 for group 2 SD standardizer

Value

Returns 1-row matrix with the estimate for the contrast. The matrix has the following columns:

- Estimate - estimate of linear contrast
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2009). "Meta-analytic interval estimation for standardized and unstandardized mean differences." *Psychological Methods*, **14**(3), 225–238. ISSN 1939-1463, doi: [10.1037/a0016619](https://doi.org/10.1037/a0016619), <https://doi.org/10.1037/a0016619>.

Examples

```
m1 <- c(53, 60, 53, 57)
m2 <- c(55, 62, 58, 61)
sd1 <- c(4.1, 4.2, 4.5, 4.0)
sd2 <- c(4.2, 4.7, 4.9, 4.8)
cor <- c(.7, .7, .8, .85)
n <- c(30, 50, 30, 70)
v <- c(.5, .5, -.5, -.5)
meta.lc.stdmean.ps(.05, m1, m2, sd1, sd2, cor, n, v, 0)

# Should return:
#           Estimate      SE      LL      UL
# Contrast  0.5127577 0.1392232 0.2398851 0.7856302
```

meta.lc.stdmean2	<i>Confidence interval for a linear contrast of standardized mean differences from 2-group studies</i>
------------------	--

Description

Computes the estimate, standard error, and confidence interval for a linear contrast of 2-group standardized mean differences from two or more studies. Equal variances within or across studies is not assumed. Use the square root average variance standardizer (`stdzr = 0`) for 2-group experimental designs. Use the square root weighted variance standardizer (`stdzr = 3`) for 2-group nonexperimental designs with simple random sampling. The `stdzr = 1` and `stdzr = 2` options can be used with either 2-group experimental or nonexperimental designs.

Usage

```
meta.lc.stdmean2(alpha, m1, m2, sd1, sd2, n1, n2, v, stdzr)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for group 1
m2	vector of sample means for group 2
sd1	vector of sample SDs for group 1
sd2	vector of sample SDs for group 2
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
v	vector of contrast coefficients
stdzr	<ul style="list-style-type: none"> • set to 0 for square root average variance standardizer • set to 1 for group 1 SD standardizer • set to 2 for group 2 SD standardizer • set to 3 for square root weighted variance standardizer

Value

Returns 1-row matrix with the estimate for the contrast. The matrix has the following columns:

- Estimate - estimate of linear contrast
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2009). "Meta-analytic interval estimation for standardized and unstandardized mean differences." *Psychological Methods*, **14**(3), 225–238. ISSN 1939-1463, doi: [10.1037/a0016619](https://doi.org/10.1037/a0016619), <https://doi.org/10.1037/a0016619>.

Examples

```
m1 <- c(45.1, 39.2, 36.3, 34.5)
m2 <- c(30.0, 35.1, 35.3, 36.2)
sd1 <- c(10.7, 10.5, 9.4, 11.5)
sd2 <- c(12.3, 12.0, 10.4, 9.6)
n1 <- c(40, 20, 50, 25)
n2 <- c(40, 20, 48, 26)
v <- c(.5, .5, -.5, -.5)
meta.lc.stdmean2(.05, m1, m2, sd1, sd2, n1, n2, v, 0)

# Should return:
#           Estimate      SE      LL      UL
# Contrast 0.8557914 0.2709192 0.3247995 1.386783
```

meta.lm.agree

Meta-regression analysis for G agreement indicces

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a G-index of agreement. The estimates are OLS estimates with standard errors that accomodate residual heteroscedasticity.

Usage

```
meta.lm.agree(alpha, f11, f12, f21, f22, X)
```

Arguments

alpha	alpha level for 1-alpha confidence
f11	vector of frequencies in cell 1,1
f12	vector of frequencies in cell 1,2
f21	vector of frequencies in cell 2,1
f22	vector of frequencies in cell 2,2
X	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- z - z-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

Examples

```
f11 <- c(40, 20, 25, 30)
f12 <- c(3, 2, 2, 1)
f21 <- c(7, 6, 8, 6)
f22 <- c(26, 25, 13, 25)
x1 <- c(1, 1, 4, 6)
x2 <- c(1, 1, 0, 0)
X <- matrix(cbind(x1, x2), 4, 2)
meta.lm.agree(.05, f11, f12, f21, f22, X)

# Should return:
#   Estimate      SE      z      p      LL      UL
# b0 0.1904762 0.38772858 0.4912617 0.623 -0.56945786 0.9504102
# b1 0.0952381 0.07141957 1.3335013 0.182 -0.04474169 0.2352179
# b2 0.4205147 0.32383556 1.2985438 0.194 -0.21419136 1.0552207
```

 meta.lm.cor

Meta-regression analysis for Pearson or partial correlations

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a Fisher-transformed Pearson or partial correlation. The estimates are OLS estimates with standard errors that accommodate residual heteroscedasticity. The correlations are Fisher-transformed and hence the parameter estimates do not have a simple interpretation. However, the hypothesis test results can be used to decide if a population slope is either positive or negative.

Usage

```
meta.lm.cor(alpha, n, cor, q, X)
```

Arguments

alpha	alpha level for 1-alpha confidence
n	vector of sample sizes
cor	vector of Pearson or partial correlations
q	number of control variables
X	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - Standard error
- z - z-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

Examples

```
n <- c(55, 190, 65, 35)
cor <- c(.40, .65, .60, .45)
q <- 0
x1 <- c(18, 25, 23, 19)
X <- matrix(x1, 4, 1)
meta.lm.cor(.05, n, cor, q, X)

# Should return:
#      Estimate      SE      z      p      LL      UL
# b0 -0.47832153 0.48631509 -0.983563 0.325 -1.431481595 0.47483852
# b1 0.05047154 0.02128496 2.371231 0.018 0.008753794 0.09218929
```

meta.lm.cor.gen

Meta-regression analysis for correlations

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a Fisher-transformed correlation. The correlations can be of different types (e.g., Pearson, partial, Spearman). The estimates are OLS estimates with standard errors that accommodate residual heteroscedasticity. This function uses sample correlations and their standard errors as input. The correlations are Fisher-transformed and hence the parameter estimates do not have a simple interpretation. However, the hypothesis test results can be used to decide if a population slope is either positive or negative.

Usage

```
meta.lm.cor.gen(alpha, cor, se, X)
```

Arguments

alpha	alpha level for 1-alpha confidence
cor	vector of sample correlations
se	number of control variables
X	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- z - z-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

Examples

```
cor <- c(.40, .65, .60, .45)
se <- c(.182, .114, .098, .132)
x1 <- c(18, 25, 23, 19)
X <- matrix(x1, 4, 1)
meta.lm.cor.gen(.05, cor, se, X)

# Should return:
#      Estimate      SE      z      p
# b0 -0.47832153 0.63427931 -0.7541181 0.451
# b1  0.05047154 0.02879859  1.7525699 0.080
```

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a log-complement Cronbach reliability. The estimates are OLS estimates with standard errors that accommodate residual heteroscedasticity. The exponentiated slope estimate for a predictor variable describes a multiplicative change in variable.

Usage

```
meta.lm.cronbach(alpha, n, rel, q, X)
```

Arguments

alpha	alpha level for 1-alpha confidence
n	vector of sample sizes
rel	vector of sample alpha reliabilities
q	number of measurements (e.g., items)
X	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - exponentiated OLS estimate
- SE - standard error
- z - z-value
- p - p-value
- LL - lower limit of the exponentiated confidence interval
- UL - upper limit of the exponentiated confidence interval

References

Bonett DG (2010). "Varying coefficient meta-analytic methods for alpha reliability." *Psychological Methods*, **15**(4), 368–385. ISSN 1939-1463, doi: [10.1037/a0020142](https://doi.org/10.1037/a0020142), <https://doi.org/getdoi.cfm?doi=10.1037/a0020142>.

Examples

```
n <- c(583, 470, 546, 680)
rel <- c(.91, .89, .90, .89)
x1 <- c(1, 0, 0, 0)
X <- matrix(x1, 4, 1)
meta.lm.cronbach(.05, n, rel, 10, X)

# Should return:
#   Estimate      SE      z      p      LL      UL
# b0 -2.2408328 0.03675883 -60.960391 0.000 -2.3128788 -2.16878684
# b1 -0.1689006 0.07204625 -2.344336 0.019 -0.3101087 -0.02769259
```

meta.lm.gen	<i>Meta-regression analysis for any type of effect size</i>
-------------	---

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is any type of effect size. The estimates are OLS estimates with standard errors that accomodate residual heteroscedasticity.

Usage

```
meta.lm.gen(alpha, est, se, X)
```

Arguments

alpha	alpha level for 1-alpha confidence
est	vector of parameter estimates
se	vector of standard errors
X	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- z - z-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

Examples

```
est <- c(4.1, 4.7, 4.9, 5.7, 6.6, 7.3)
se <- c(1.2, 1.5, 1.3, 1.8, 2.0, 2.6)
x1 <- c(10, 20, 30, 40, 50, 60)
x2 <- c(1, 1, 1, 0, 0, 0)
X <- matrix(cbind(x1, x2), 6, 2)
meta.lm.gen(.05, est, se, X)

# Should return:
#      Estimate      SE      z      p      LL      UL
# b0  3.5333333 4.37468253 0.80767766 0.419 -5.0408869 12.1075535
# b1  0.0600000 0.09058835 0.66233679 0.508 -0.1175499 0.2375499
# b2 -0.1666667 2.81139793 -0.05928249 0.953 -5.6769054 5.3435720
```

`meta.lm.mean.ps`*Meta-regression analysis for paired-samples mean differences*

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a paired-samples mean difference. The estimates are OLS estimates with standard errors that accommodate residual heteroscedasticity.

Usage

```
meta.lm.mean.ps(alpha, m1, m2, sd1, sd2, cor, n, X)
```

Arguments

<code>alpha</code>	alpha level for 1-alpha confidence
<code>m1</code>	vector of sample means for group 1
<code>m2</code>	vector of sample means for group 2
<code>sd1</code>	vector of sample SDs for group 1
<code>sd2</code>	vector of sample SDs for group 2
<code>cor</code>	vector of correlations
<code>n</code>	vector of sample sizes
<code>X</code>	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- t - t-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- df - degrees of freedom

References

Bonett DG (2009). "Meta-analytic interval estimation for standardized and unstandardized mean differences." *Psychological Methods*, **14**(3), 225–238. ISSN 1939-1463, doi: [10.1037/a0016619](https://doi.org/10.1037/a0016619), <https://doi.org/10.1037/a0016619>.

Examples

```

n <- c(65, 30, 29, 45, 50)
cor <- c(.87, .92, .85, .90, .88)
m1 <- c(20.1, 20.5, 19.3, 21.5, 19.4)
m2 <- c(10.4, 10.2, 8.5, 10.3, 7.8)
sd1 <- c(9.3, 9.9, 10.1, 10.5, 9.8)
sd2 <- c(7.8, 8.0, 8.4, 8.1, 8.7)
x1 <- c(2, 3, 3, 4, 4)
X <- matrix(x1, 5, 1)
meta.lm.mean.ps(.05, m1, m2, sd1, sd2, cor, n, X)

# Should return:
#   Estimate      SE      t      p      LL      UL  df
# b0      8.00 1.2491990 6.404104 0.000 5.5378833 10.462117 217
# b1      0.85 0.3796019 2.239188 0.026 0.1018213  1.598179 217

```

 meta.lm.mean1

Meta-regression analysis for single means

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a mean from a single group. The estimates are OLS estimates with standard errors that accomodate residual heteroscedasticity.

Usage

```
meta.lm.mean1(alpha, m, sd, n, X)
```

Arguments

alpha	alpha level for 1-alpha confidence
m	vector of sample means
sd	vector of sample standard deviations
n	vector of sample sizes
X	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- t - t-value

- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- df - degrees of freedom

Examples

```
n <- c(25, 15, 30, 25, 40)
m <- c(20.1, 20.5, 19.3, 21.5, 19.4)
sd <- c(10.4, 10.2, 8.5, 10.3, 7.8)
x1 <- c(1, 1, 0, 0, 0)
x2 <- c(12, 13, 11, 13, 15)
X <- matrix(cbind(x1, x2), 5, 2)
meta.lm.mean1(.05, m, sd, n, X)

# Should return:
#      Estimate      SE      t      p      LL      UL      df
# b0 19.45490196 6.7873381 2.86635227 0.005 6.0288763 32.880928 132
# b1  0.25686275 1.9834765 0.12950128 0.897 -3.6666499  4.180375 132
# b2  0.04705882 0.5064693 0.09291544 0.926 -0.9547876  1.048905 132
```

meta.lm.mean2

Meta-regression analysis for 2-group mean differences

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a 2-group mean difference. The estimates are OLS estimates with standard errors that accommodate residual heteroscedasticity.

Usage

```
meta.lm.mean2(alpha, m1, m2, sd1, sd2, n1, n2, X)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for group 1
m2	vector of sample means for group 2
sd1	vector of sample SDs for group 1
sd2	vector of sample SDs for group 2
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
X	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- t - t-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- df - degrees of freedom

References

Bonett DG (2009). "Meta-analytic interval estimation for standardized and unstandardized mean differences." *Psychological Methods*, **14**(3), 225–238. ISSN 1939-1463, doi: [10.1037/a0016619](https://doi.org/10.1037/a0016619), <https://doi.org/10.1037/a0016619>.

Examples

```
n1 <- c(65, 30, 29, 45, 50)
n2 <- c(67, 32, 31, 20, 52)
m1 <- c(31.1, 32.3, 31.9, 29.7, 33.0)
m2 <- c(34.1, 33.2, 30.6, 28.7, 26.5)
sd1 <- c(7.1, 8.1, 7.8, 6.8, 7.6)
sd2 <- c(7.8, 7.3, 7.5, 7.2, 6.8)
x1 <- c(4, 6, 7, 7, 8)
x2 <- c(1, 0, 0, 0, 1)
X <- matrix(cbind(x1, x2), 5, 2)
meta.lm.mean2(.05, m1, m2, sd1, sd2, n1, n2, X)

# Should return:
#   Estimate      SE      t      p      LL      UL      df
# b0  -15.20  3.4097610 -4.457791 0.000 -21.902415 -8.497585 418
# b1   2.35  0.4821523  4.873979 0.000  1.402255  3.297745 418
# b2   2.85  1.5358109  1.855697 0.064  -0.168875  5.868875 418
```

meta.lm.meanratio.ps *Meta-regression analysis for paired-samples log mean ratios*

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a paired-samples log mean ratio. The estimates are OLS estimates with standard errors that accommodate residual heteroscedasticity.

Usage

```
meta.lm.meanratio.ps(alpha, m1, m2, sd1, sd2, cor, n, X)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for group 1
m2	vector of sample means for group 2
sd1	vector of sample SDs for group 1
sd2	vector of sample SDs for group 2
cor	vector of correlations
n	vector of sample sizes
X	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- z - z-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- exp(Estimate) - the exponentiated estimate
- exp(LL) - lower limit of the exponentiated confidence interval
- exp(UL) - upper limit of the exponentiated confidence interval

Examples

```
n <- c(65, 30, 29, 45, 50)
cor <- c(.87, .92, .85, .90, .88)
m1 <- c(20.1, 20.5, 19.3, 21.5, 19.4)
m2 <- c(10.4, 10.2, 8.5, 10.3, 7.8)
sd1 <- c(9.3, 9.9, 10.1, 10.5, 9.8)
sd2 <- c(7.8, 8.0, 8.4, 8.1, 8.7)
x1 <- c(2, 3, 3, 4, 4)
X <- matrix(x1, 5, 1)
meta.lm.meanratio.ps(.05, m1, m2, sd1, sd2, cor, n, X)

# Should return:
#   Estimate      SE      LL      UL      z      p
# b0 0.50957008 0.13000068 0.254773424 0.7643667 3.919749 0.000
# b1 0.07976238 0.04133414 -0.001251047 0.1607758 1.929697 0.054
#   exp(Estimate) exp(LL) exp(UL)
```

```
# b0      1.664575 1.2901693 2.147634
# b1      1.083030 0.9987497 1.174422
```

meta.lm.meanratio2 *Meta-regression analysis for 2-group log mean ratios*

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a 2-group log mean ratio. The estimates are OLS estimates with standard errors that accommodate residual heteroscedasticity.

Usage

```
meta.lm.meanratio2(alpha, m1, m2, sd1, sd2, n1, n2, X)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for group 1
m2	vector of sample means for group 2
sd1	vector of sample SDs for group 1
sd2	vector of sample SDs for group 2
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
X	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- z - z-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- exp(Estimate) - the exponentiated estimate
- exp(LL) - lower limit of the exponentiated confidence interval
- exp(UL) - upper limit of the exponentiated confidence interval

Examples

```

n1 <- c(65, 30, 29, 45, 50)
n2 <- c(67, 32, 31, 20, 52)
m1 <- c(31.1, 32.3, 31.9, 29.7, 33.0)
m2 <- c(34.1, 33.2, 30.6, 28.7, 26.5)
sd1 <- c(7.1, 8.1, 7.8, 6.8, 7.6)
sd2 <- c(7.8, 7.3, 7.5, 7.2, 6.8)
x1 <- c(4, 6, 7, 7, 8)
X <- matrix(x1, 5, 1)
meta.lm.meanratio2(.05, m1, m2, sd1, sd2, n1, n2, X)

# Should return:
#      Estimate      SE      LL      UL      z p
# b0 -0.40208954 0.09321976 -0.58479692 -0.21938216 -4.313351 0
# b1  0.06831545 0.01484125  0.03922712  0.09740377  4.603078 0
#   exp(Estimate) exp(LL)  exp(UL)
# b0      0.6689208 0.557219 0.8030148
# b1      1.0707030 1.040007 1.1023054

```

 meta.lm.odds

Meta-regression analysis for odds ratios

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a log odds ratio. The estimates are OLS estimates with standard errors that accommodate residual heteroscedasticity. The exponentiated slope estimate for a predictor variable describes a multiplicative change in the odds ratio associated with a 1-unit increase in that predictor variable.

Usage

```
meta.lm.odds(alpha, f1, f2, n1, n2, X)
```

Arguments

alpha	alpha level for 1-alpha confidence
f1	vector of group 1 event counts
f2	vector of group 2 event counts
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
X	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- z - z-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- exp(Estimate) - the exponentiated estimate
- exp(LL) - lower limit of the exponentiated confidence interval
- exp(UL) - upper limit of the exponentiated confidence interval

References

Bonett DG, Price RM (2015). “Varying coefficient meta-analysis methods for odds ratios and risk ratios.” *Psychological Methods*, **20**(3), 394–406. ISSN 1939-1463, doi: [10.1037/met0000032](https://doi.org/10.1037/met0000032), <https://doi.apa.org/getdoi.cfm?doi=10.1037/met0000032>.

Examples

```
n1 <- c(204, 201, 932, 130, 77)
n2 <- c(106, 103, 415, 132, 83)
f1 <- c(24, 40, 93, 14, 5)
f2 <- c(12, 9, 28, 3, 1)
x1 <- c(4, 4, 5, 3, 26)
x2 <- c(1, 1, 1, 0, 0)
X <- matrix(cbind(x1, x2), 5, 2)
meta.lm.odds(.05, f1, f2, n1, n2, X)

# Should return:
#      Estimate      SE          z      p      LL      UL
# b0  1.541895013 0.69815801  2.20851868 0.027  0.1735305 2.91025958
# b1 -0.004417932 0.04840623 -0.09126784 0.927 -0.0992924 0.09045653
# b2 -1.071122269 0.60582695 -1.76803337 0.077 -2.2585213 0.11627674
#   exp(Estimate) exp(LL) exp(UL)
# b0    4.6734381 1.1894969 18.361564
# b1    0.9955918 0.9054779  1.094674
# b2    0.3426238 0.1045049  1.123307
```

`meta.lm.prop.ps`*Meta-regression analysis for paired-samples proportion differences*

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a paired-samples proportion difference. The estimates are OLS estimates with standard errors that accommodate residual heteroscedasticity.

Usage

```
meta.lm.prop.ps(alpha, f11, f12, f21, f22, X)
```

Arguments

<code>alpha</code>	alpha level for 1-alpha confidence
<code>f11</code>	vector of frequencies in cell 1,1
<code>f12</code>	vector of frequencies in cell 1,2
<code>f21</code>	vector of frequencies in cell 2,1
<code>f22</code>	vector of frequencies in cell 2,2
<code>X</code>	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- z - z-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG, Price RM (2014). "Meta-analysis methods for risk differences." *British Journal of Mathematical and Statistical Psychology*, **67**(3), 371–387. ISSN 00071102, doi: [10.1111/bmsp.12024](https://doi.org/10.1111/bmsp.12024), <https://onlinelibrary.wiley.com/doi/10.1111/bmsp.12024>.

Examples

```
f11 <- c(40, 20, 25, 30)
f12 <- c(3, 2, 2, 1)
f21 <- c(7, 6, 8, 6)
f22 <- c(26, 25, 13, 25)
x1 <- c(1, 1, 4, 6)
x2 <- c(1, 1, 0, 0)
X <- matrix(cbind(x1, x2), 4, 2)
meta.lm.prop.ps(.05, f11, f12, f21, f22, X)

# Should return:
#      Estimate      SE      z      p      LL      UL
# b0 -0.21113402 0.21119823 -0.9996960 0.317 -0.62507494 0.20280690
# b1  0.02185567 0.03861947  0.5659236 0.571 -0.05383711 0.09754845
# b2  0.12575138 0.17655623  0.7122455 0.476 -0.22029248 0.47179524
```

 meta.lm.prop1

Meta-regression analysis for single proportions

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a proportion from a single group. The estimates are OLS estimates with standard errors that accomodate residual heteroscedasticity.

Usage

```
meta.lm.prop1(alpha, f, n, X)
```

Arguments

alpha	alpha level for 1-alpha confidence
f	vector of sample frequency counts
n	vector of sample sizes
X	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- z - z-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

Examples

```
f <- c(38, 26, 24, 15, 45, 38)
n <- c(80, 60, 70, 50, 180, 200)
x1 <- c(10, 15, 18, 22, 24, 30)
X <- matrix(x1, 6, 1)
meta.lm.prop1(.05, f, n, X)

# Should return:
#      Estimate      SE      z p      LL      UL
# b0  0.63262816 0.06845707  9.241239 0  0.49845477  0.766801546
# b1 -0.01510565 0.00290210 -5.205076 0 -0.02079367 -0.009417641
```

 meta.lm.prop2

Meta-regression analysis for 2-group proportion differences

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a 2-group proportion difference. The estimates are OLS estimates with standard errors that accommodate residual heteroscedasticity.

Usage

```
meta.lm.prop2(alpha, f1, f2, n1, n2, X)
```

Arguments

alpha	alpha level for 1-alpha confidence
f1	vector of group 1 event counts
f2	vector of group 2 event counts
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
X	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- z - z-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG, Price RM (2014). “Meta-analysis methods for risk differences.” *British Journal of Mathematical and Statistical Psychology*, **67**(3), 371–387. ISSN 00071102, doi: [10.1111/bmsp.12024](https://onlinelibrary.wiley.com/doi/10.1111/bmsp.12024), <https://onlinelibrary.wiley.com/doi/10.1111/bmsp.12024>.

Examples

```
f1 <- c(24, 40, 93, 14, 5)
f2 <- c(12, 9, 28, 3, 1)
n1 <- c(204, 201, 932, 130, 77)
n2 <- c(106, 103, 415, 132, 83)
x1 <- c(4, 4, 5, 3, 26)
x2 <- c(1, 1, 1, 0, 0)
X <- matrix(cbind(x1, x2), 5, 2)
meta.lm.prop2(.05, f1, f2, n1, n2, X)

# Should return:
#      Estimate      SE      z      p      LL      UL
# b0  0.089756283  0.034538077  2.5987632  0.009  0.02206290  0.157449671
# b1 -0.001447968  0.001893097 -0.7648672  0.444 -0.00515837  0.002262434
# b2 -0.034670988  0.034125708 -1.0159786  0.310 -0.10155615  0.032214170
```

meta.lm.propratio2 *Meta-regression analysis for proportion ratios*

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a log proportion ratio. The estimates are OLS estimates with standard errors that accommodate residual heteroscedasticity. The exponentiated slope estimate for a predictor variable describes a multiplicative change in the proportion ratio associated with a 1-unit increase in that predictor variable.

Usage

```
meta.lm.propratio2(alpha, f1, f2, n1, n2, X)
```

Arguments

alpha	alpha level for 1-alpha confidence
f1	vector of group 1 event counts
f2	vector of group 2 event counts
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
X	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- z - z-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- exp(Estimate) - the exponentiated estimate
- exp(LL) - lower limit of the exponentiated confidence interval
- exp(UL) - upper limit of the exponentiated confidence interval

References

Price RM, Bonett DG (2008). “Confidence intervals for a ratio of two independent binomial proportions.” *Statistics in Medicine*, **27**(26), 5497–5508. ISSN 02776715, doi: [10.1002/sim.3376](https://onlinelibrary.wiley.com/doi/10.1002/sim.3376), <https://onlinelibrary.wiley.com/doi/10.1002/sim.3376>.

Examples

```
n1 <- c(204, 201, 932, 130, 77)
n2 <- c(106, 103, 415, 132, 83)
f1 <- c(24, 40, 93, 14, 5)
f2 <- c(12, 9, 28, 3, 1)
x1 <- c(4, 4, 5, 3, 26)
x2 <- c(1, 1, 1, 0, 0)
X <- matrix(cbind(x1, x2), 5, 2)
meta.lm.propratio2(.05, f1, f2, n1, n2, X)

# Should return:
#      Estimate      SE      z      p      LL      UL
# b0  1.4924887636 0.69172794 2.15762393 0.031 0.13672691 2.84825062
# b1  0.0005759509 0.04999884 0.01151928 0.991 -0.09741998 0.09857188
# b2 -1.0837844594 0.59448206 -1.82307345 0.068 -2.24894789 0.08137897
#      exp(Estimate) exp(LL) exp(UL)
# b0      4.4481522 1.1465150 17.257565
# b1      1.0005761 0.9071749 1.103594
# b2      0.3383128 0.1055102 1.084782
```

meta.lm.semipart *Meta-regression analysis for semipartial correlations*

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a Fisher-transformed semipartial correlation. The estimates are OLS estimates with standard errors that accommodate residual heteroscedasticity. The correlations are Fisher-transformed and hence the parameter estimates do not have a simple interpretation. However, the hypothesis test results can be used to decide if a population slope is either positive or negative.

Usage

```
meta.lm.semipart(alpha, n, cor, r2, X)
```

Arguments

alpha	alpha level for 1-alpha confidence
n	vector of sample sizes
cor	vector of sample semipartial correlations
r2	vector of squared multiple correlations for full model
X	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- z - z-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

Examples

```
n <- c(128, 97, 210, 217)
cor <- c(.35, .41, .44, .39)
r2 <- c(.29, .33, .36, .39)
x1 <- c(18, 25, 23, 19)
X <- matrix(x1, 4, 1)
meta.lm.semipart(.05, n, cor, r2, X)
```

```
# Should return:
```

```
#      Estimate      SE      z      p      LL      UL
# b0 0.19695988 0.3061757 0.6432905 0.520 -0.40313339 0.79705315
# b1 0.01055584 0.0145696 0.7245114 0.469 -0.01800004 0.03911172
```

meta.lm.spear

Meta-regression analysis for Spearman correlations

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a Fisher-transformed Spearman correlation. The estimates are OLS estimates with standard errors that accommodate residual heteroscedasticity. The correlations are Fisher-transformed and hence the parameter estimates do not have a simple interpretation. However, the hypothesis test results can be used to decide if a population slope is either positive or negative.

Usage

```
meta.lm.spear(alpha, n, cor, X)
```

Arguments

alpha	alpha level for 1-alpha confidence
n	vector of sample sizes
cor	vector of Spearman correlations
X	matrix of predictor values

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- z - z-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

Examples

```

n <- c(150, 200, 300, 200, 350)
cor <- c(.14, .29, .16, .21, .23)
x1 <- c(18, 25, 23, 19, 24)
X <- matrix(x1, 5, 1)
meta.lm.spear(.05, n, cor, X)

# Should return:
#      Estimate      SE      z      p      LL      UL
# b0 -0.08920088 0.26686388 -0.3342561 0.738 -0.612244475 0.43384271
# b1  0.01370866 0.01190212  1.1517825 0.249 -0.009619077 0.03703639

```

meta.lm.stdmean.ps	<i>Meta-regression analysis for paired-samples standardized mean differences</i>
--------------------	--

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a paired-samples standardized mean difference. The estimates are OLS estimates with standard errors that accommodate residual heteroscedasticity.

Usage

```
meta.lm.stdmean.ps(alpha, m1, m2, sd1, sd2, cor, n, X, stdzr)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for group 1
m2	vector of sample means for group 2
sd1	vector of sample SDs for group 1
sd2	vector of sample SDs for group 2
cor	vector of correlations
n	vector of sample sizes
X	matrix of predictor values
stdzr	<ul style="list-style-type: none"> • set to 0 for square root average variance standardizer • set to 1 for group 1 SD standardizer • set to 2 for group 2 SD standardizer

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- t - t-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2009). "Meta-analytic interval estimation for standardized and unstandardized mean differences." *Psychological Methods*, **14**(3), 225–238. ISSN 1939-1463, doi: [10.1037/a0016619](https://doi.org/10.1037/a0016619), [https://doi.org/getdoi.cfm?doi=10.1037/a0016619](https://doi.org/10.1037/a0016619).

Examples

```
n <- c(65, 30, 29, 45, 50)
cor <- c(.87, .92, .85, .90, .88)
m1 <- c(20.1, 20.5, 19.3, 21.5, 19.4)
m2 <- c(10.4, 10.2, 8.5, 10.3, 7.8)
sd1 <- c(9.3, 9.9, 10.1, 10.5, 9.8)
sd2 <- c(7.8, 8.0, 8.4, 8.1, 8.7)
x1 <- c(2, 3, 3, 4, 4)
X <- matrix(x1, 5, 1)
meta.lm.stdmean.ps(.05, m1, m2, sd1, sd2, cor, n, X, 0)

# Should return:
#   Estimate      SE      z      p      LL      UL
# b0 1.01740253 0.25361725 4.0115667 0.000 0.5203218 1.5144832
# b1 0.04977943 0.07755455 0.6418635 0.521 -0.1022247 0.2017836
```

meta.lm.stdmean2

Meta-regression analysis for 2-group standardized mean differences

Description

This function estimates the intercept and slope coefficients in a meta-regression model where the dependent variable is a 2-group standardized mean difference. The estimates are OLS estimates with standard errors that accommodate residual heteroscedasticity.

Usage

```
meta.lm.stdmean2(alpha, m1, m2, sd1, sd2, n1, n2, X, stdzr)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for group 1
m2	vector of sample means for group 2
sd1	vector of sample SDs for group 1
sd2	vector of sample SDs for group 2
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
X	matrix of predictor values
stdzr	<ul style="list-style-type: none"> • set to 0 for square root average variance standardizer • set to 1 for group 1 SD standardizer • set to 2 for group 2 SD standardizer • set to 3 for square root weighted variance standardizer

Value

Returns a matrix. The first row is for the intercept with one additional row per predictor. The matrix has the following columns:

- Estimate - OLS estimate
- SE - standard error
- z - z-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2009). "Meta-analytic interval estimation for standardized and unstandardized mean differences." *Psychological Methods*, **14**(3), 225–238. ISSN 1939-1463, doi: [10.1037/a0016619](https://doi.org/10.1037/a0016619), <https://doi.org/10.1037/a0016619>.

Examples

```
n1 <- c(65, 30, 29, 45, 50)
n2 <- c(67, 32, 31, 20, 52)
m1 <- c(31.1, 32.3, 31.9, 29.7, 33.0)
m2 <- c(34.1, 33.2, 30.6, 28.7, 26.5)
sd1 <- c(7.1, 8.1, 7.8, 6.8, 7.6)
sd2 <- c(7.8, 7.3, 7.5, 7.2, 6.8)
x1 <- c(4, 6, 7, 7, 8)
```

```

X <- matrix(x1, 5, 1)
meta.lm.stdmean2(.05, m1, m2, sd1, sd2, n1, n2, X, 0)

# Should return:
#   Estimate      SE      z p      LL      UL
# b0 -1.6988257 0.4108035 -4.135373 0 -2.5039857 -0.8936657
# b1  0.2871641 0.0649815  4.419167 0  0.1598027  0.4145255

```

meta.sub.cor

Confidence interval for a difference in average Pearson or partial correlations for two sets of studies

Description

Computes the estimate, standard error, and confidence interval for a difference in average Pearson or partial correlations for two mutually exclusive sets of studies. Each set can have one or more studies. All of the correlations must be either Pearson correlations or partial correlations.

Usage

```
meta.sub.cor(alpha, n, cor, q, group)
```

Arguments

alpha	alpha level for 1-alpha confidence
n	vector of sample sizes
cor	vector of sample Pearson correlations
q	number of control variables (set to 0 for Pearson)
group	vector of group indicators: <ul style="list-style-type: none"> • 1 for set A • 2 for set B • 0 to ignore

Value

Returns a matrix with three rows:

- Row 1 - Estimate for Set A
- Row 2 - Estimate for Set B
- Row 3 - Estimate for difference, Set A - Set B

The columns are:

- Estimate - estimate of average correlation or difference
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2008). “Meta-analytic interval estimation for bivariate correlations.” *Psychological Methods*, **13**(3), 173–181. ISSN 1939-1463, doi: [10.1037/a0012868](https://doi.org/10.1037/a0012868), <https://doi.org/getdoi.cfm?doi=10.1037/a0012868>.

Examples

```
n <- c(55, 190, 65, 35)
cor <- c(.40, .65, .60, .45)
group <- c(1, 1, 2, 0)
meta.sub.cor(.05, n, cor, 0, group)

# Should return:
#           Estimate      SE      LL      UL
# Set A:      0.525 0.06195298  0.3932082 0.6356531
# Set B:      0.600 0.08128008  0.4171458 0.7361686
# Set A - Set B: -0.075 0.10219894 -0.2645019 0.1387283
```

meta.sub.cronbach	<i>Confidence interval for a difference in average Cronbach reliabilities for two sets of studies</i>
-------------------	---

Description

Computes the estimate, standard error, and confidence interval for a difference in average Cronbach reliability coefficients for two mutually exclusive sets of studies. Each set can have one or more studies. The number of measurements used to compute the sample reliability coefficient is assumed to be the same for all studies.

Usage

```
meta.sub.cronbach(alpha, n, rel, q, group)
```

Arguments

alpha	alpha level for 1-alpha confidence
n	vector of sample sizes
rel	vector of sample Cronbach reliabilities
q	number of measurements (e.g., items)
group	vector of group indicators: <ul style="list-style-type: none"> • 1 for set A • 2 for set B • 0 to ignore

Value

Returns a matrix with three rows:

- Row 1 - Estimate for Set A
- Row 2 - Estimate for Set B
- Row 3 - Estimate for difference, Set A - Set B

The columns are:

- Estimate - estimate of average correlation or difference
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2010). "Varying coefficient meta-analytic methods for alpha reliability." *Psychological Methods*, **15**(4), 368–385. ISSN 1939-1463, doi: [10.1037/a0020142](https://doi.org/10.1037/a0020142), [https://doi.org/getdoi.cfm?doi=10.1037/a0020142](https://doi.org/10.1037/a0020142).

Examples

```
n <- c(120, 170, 150, 135)
rel <- c(.89, .87, .73, .71)
group <- c(1, 1, 2, 2)
q <- 10
meta.sub.cronbach(.05, n, rel, q, group)

# Should return:
#           Estimate      SE      LL      UL
# Set A:      0.88 0.01068845 0.8581268 0.8999386
# Set B:      0.72 0.02515130 0.6684484 0.7668524
# Set A - Set B: 0.16 0.02732821 0.1082933 0.2152731
```

meta.sub.pbcor

Confidence interval for a difference in average point-biserial correlations for two sets of studies

Description

Computes the estimate, standard error, and confidence interval for a difference in average point-biserial correlations for two mutually exclusive sets of studies. Each set can have one or more studies. Two types of point-biserial correlations can be analyzed. One type uses an unweighted variance and is appropriate for 2-group experimental designs. The other type uses a weighted variance and is appropriate for 2-group nonexperimental designs with simple random sampling. Equal variances within or across studies is not assumed.

Usage

```
meta.sub.pbcor(alpha, m1, m2, sd1, sd2, n1, n2, type, group)
```

Arguments

alpha	alpha level for 1-alpha confidence
m1	vector of sample means for group 1
m2	vector of sample means for group 2
sd1	vector of sample SDs for group 1
sd2	vector of sample SDs for group 2
n1	vector of group 1 sample sizes
n2	vector of group 2 sample sizes
type	<ul style="list-style-type: none"> • set to 1 for weighted variance • set to 2 for unweighted variance
group	vector of group indicators: <ul style="list-style-type: none"> • 1 for set A • 2 for set B • 0 to ignore

Value

Returns a matrix with three rows:

- Row 1 - Estimate for Set A
- Row 2 - Estimate for Set B
- Row 3 - Estimate for difference, Set A - Set B

The columns are:

- Estimate - estimate of average correlation or difference
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2020). "Point-biserial correlation: Interval estimation, hypothesis testing, meta-analysis, and sample size determination." *British Journal of Mathematical and Statistical Psychology*, **73**(S1), 113–144. ISSN 0007-1102, doi: [10.1111/bmsp.12189](https://doi.org/10.1111/bmsp.12189), <https://onlinelibrary.wiley.com/doi/10.1111/bmsp.12189>.

Examples

```

m1 <- c(45.1, 39.2, 36.3, 34.5)
m2 <- c(30.0, 35.1, 35.3, 36.2)
sd1 <- c(10.7, 10.5, 9.4, 11.5)
sd2 <- c(12.3, 12.0, 10.4, 9.6)
n1 <- c(40, 20, 50, 25)
n2 <- c(40, 20, 48, 26)
group <- c(1, 1, 2, 2)
meta.sub.pbcor(.05, m1, m2, sd1, sd2, n1, n2, 2, group)

# Should return:
#           Estimate      SE      LL      UL
# Set A:      0.36338772 0.08552728 0.1854777 0.5182304
# Set B:     -0.01480511 0.08741322 -0.1840491 0.1552914
# Set A - Set B: 0.37819284 0.12229467 0.1320530 0.6075828

```

meta.sub.semipart	<i>Confidence interval for a difference in average semipartial correlations for two sets of studies</i>
-------------------	---

Description

Computes the estimate, standard error, and confidence interval for a difference in average semipartial correlations for two sets of mutually exclusive studies. Each set can have one or more studies.

Usage

```
meta.sub.semipart(alpha, n, cor, r2, group)
```

Arguments

alpha	alpha level for 1-alpha confidence
n	vector of sample sizes
cor	vector of sample semi-partial correlations
r2	vector of squared multiple correlations for a model that includes the IV and all control variables
group	vector of group indicators: <ul style="list-style-type: none"> • 1 for set A • 2 for set B • 0 to ignore

Value

Returns a matrix with three rows:

- Row 1 - Estimate for Set A
- Row 2 - Estimate for Set B
- Row 3 - Estimate for difference, Set A - Set B

The columns are:

- Estimate - estimate of average correlation or difference
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

Examples

```
n <- c(55, 190, 65, 35)
cor <- c(.40, .65, .60, .45)
r2 <- c(.25, .41, .43, .39)
group <- c(1, 1, 2, 0)
meta.sub.semipart(.05, n, cor, r2, group)

# Should return:
#           Estimate      SE      LL      UL
# Set A:      0.525 0.05955276  0.3986844 0.6317669
# Set B:      0.600 0.07931155  0.4221127 0.7333949
# Set A - Set B: -0.075 0.09918091 -0.2587113 0.1324682
```

meta.sub.spear	<i>Confidence interval for a difference in average Spearman correlations for two sets of studies</i>
----------------	--

Description

Computes the estimate, standard error, and confidence interval for a difference in average Spearman correlations for two mutually exclusive sets of studies. Each set can have one or more studies.

Usage

```
meta.sub.spear(alpha, n, cor, group)
```

Arguments

alpha	alpha level for 1-alpha confidence
n	vector of sample sizes
cor	vector of sample Spearman correlations
group	vector of group indicators: <ul style="list-style-type: none"> • 1 for set A • 2 for set B • 0 to ignore

Value

Returns a matrix with three rows:

- Row 1 - Estimate for Set A
- Row 2 - Estimate for Set B
- Row 3 - Estimate for difference, Set A - Set B

The columns are:

- Estimate - estimate of average correlation or difference
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett, D.G. (2008). Meta-analytic interval estimation for bivariate correlations. *Psychological Methods*, 13, 173-189.

Examples

```
n <- c(55, 190, 65, 35)
cor <- c(.40, .65, .60, .45)
group <- c(1, 1, 2, 0)
meta.sub.spear(.05, n, cor, group)

# Should return:
#           Estimate      SE      LL      UL
# Set A:      0.525 0.06483629 0.3865928 0.6402793
# Set B:      0.600 0.08829277 0.3992493 0.7458512
# Set A - Set B: -0.075 0.10954158 -0.2760700 0.1564955
```

replicate.cor	<i>Compares Pearson or partial correlations in original and follow-up studies</i>
---------------	---

Description

This function can be used to compare and combine Pearson or partial correlations from the original study and the follow-up study. The confidence level for the difference is $1 - 2\alpha$.

Usage

```
replicate.cor(alpha, cor1, n1, cor2, n2, s)
```

Arguments

alpha	alpha level for 1-alpha confidence
cor1	sample Pearson correlation between y and x in original study
n1	sample size in original study
cor2	sample Pearson correlation between y and x in follow-up study
n2	sample size in follow-up study
s	number of control variables in each study (0 for Pearson)

Value

A 4-row matrix. The rows are:

- Row 1 summarizes the Original study
- Row 2 summarizes the Follow-up study
- Row 3 estimates the difference between studies
- Row 4 estimates the average effect size between the two studies

The columns are:

- Estimate - Pearson or partial correlation estimate
- SE - standard error
- z - z-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2021). "Design and analysis of replication studies." *Organizational Research Methods*, **24**(3), 513–529. ISSN 1094-4281, doi: [10.1177/1094428120911088](https://doi.org/10.1177/1094428120911088), <https://journals.sagepub.com/doi/10.1177/1094428120911088>.

Examples

```
replicate.cor(.05, .598, 80, .324, 200, 0)

# Should return:
#           Estimate      SE      z      p      LL      UL
# Original:      0.598 0.11396058 6.589418 4.708045e-09 0.4355043 0.7227538
# Follow-up:      0.324 0.07124705 4.819037 2.865955e-06 0.1939787 0.4428347
# Original - Follow-up: 0.274 0.09708614 2.633335 8.455096e-03 0.1065496 0.4265016
# Average:        0.461 0.04854307 7.634998 2.264855e-14 0.3725367 0.5411607
```

 replicate.gen

Compares effect sizes in the original and follow-up studies

Description

This function can be used to compare and combine any effect size (e.g., odds ratio, proportion ratio, proportion difference, slope coefficient, etc.) using the effect size estimate and its standard error from the original study and the follow-up study. The same results can be obtained using the [meta.lc.gen](#) function with appropriate contrast coefficients. The confidence level for the difference is $1 - 2\alpha$.

Usage

```
replicate.gen(alpha, est1, se1, est2, se2)
```

Arguments

alpha	alpha level for 1-alpha confidence
est1	sample effect size in original study
se1	effect size standard error in original study
est2	sample effect size in follow-up study
se2	effect size standard error in follow-up study

Value

A 4-row matrix. The rows are:

- Row 1 summarizes the Original study
- Row 2 summarizes the Follow-up study
- Row 3 estimates the difference between studies
- Row 4 estimates the average effect size between the two studies

Columns are:

- Estimate - effect size estimate

- SE - standard error
- z - z-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2021). “Design and analysis of replication studies.” *Organizational Research Methods*, **24**(3), 513–529. ISSN 1094-4281, doi: [10.1177/1094428120911088](https://doi.org/10.1177/1094428120911088), <https://journals.sagepub.com/doi/10.1177/1094428120911088>.

Examples

```
replicate.gen(.05, .782, .210, .650, .154)

# Should return:
#           Estimate      SE      z      p      LL      UL
# Original:      0.782 0.2100000 3.7238095 1.962390e-04 0.3704076 1.1935924
# Follow-up:     0.650 0.1540000 4.2207792 2.434593e-05 0.3481655 0.9518345
# Original - Follow-up: 0.132 0.2604151 0.5068831 6.122368e-01 -0.2963446 0.5603446
# Average:       0.716 0.1302075 5.4989141 3.821373e-08 0.4607979 0.9712021
```

replicate.mean.ps	<i>Compares paired-samples mean differences in original and follow-up studies</i>
-------------------	---

Description

This function computes confidence intervals for a paired-samples mean difference from an original study and a follow-up study. Confidence intervals for the difference and average effect size also are computed. A Satterthwaite adjustment to the degrees of freedom is used to improve the accuracy of the confidence intervals for the difference and average. The same results can be obtained using the [meta.lc.mean.ps](#) function with appropriate contrast coefficients. The confidence level for the difference is $1 - 2\alpha$.

Usage

```
replicate.mean.ps(
  alpha,
  m11,
  m12,
  sd11,
  sd12,
  cor1,
  n1,
```

```

    m21,
    m22,
    sd21,
    sd22,
    cor2,
    n2
  )

```

Arguments

alpha	alpha level for 1-alpha confidence
m11	sample mean for group 1 in original study
m12	sample mean for group 2 in original study
sd11	sample SD for group 1 in original study
sd12	sample SD for group 2 in original study
cor1	sample correlation of paired observations in original study
n1	sample size in original study
m21	sample mean for group 1 in follow-up study
m22	sample mean for group 2 in follow-up study
sd21	sample SD for group 1 in follow-up study
sd22	sample SD for group 2 in follow-up study
cor2	sample correlation of paired observations in follow-up study
n2	sample size in follow-up study

Value

A 4-row matrix. The rows are:

- Row 1 summarizes the Original study
- Row 2 summarizes the Follow-up study
- Row 3 estimates the difference between studies
- Row 4 estimates the average effect size between the two studies

The columns are:

- Estimate - effect size estimate
- SE - standard error
- df - degrees of freedom
- t - t-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2021). “Design and analysis of replication studies.” *Organizational Research Methods*, **24**(3), 513–529. ISSN 1094-4281, doi: [10.1177/1094428120911088](https://doi.org/10.1177/1094428120911088), <https://journals.sagepub.com/doi/10.1177/1094428120911088>.

Examples

```
replicate.mean.ps(.05, 86.22, 70.93, 14.89, 12.32, .765, 20,
                 84.81, 77.24, 15.68, 16.95, .702, 75)

# Should return:
#           Estimate      SE      t      p
# Original:      15.29 2.154344  7.097288 9.457592e-07
# Follow-up:      7.57 1.460664  5.182575 1.831197e-06
# Original - Follow-up: 7.72 2.602832  2.966000 5.166213e-03
# Average:       11.43 1.301416  8.782740 1.010232e-10
#           LL      UL      df
# Original:     10.780906 19.79909 19.00000
# Follow-up:     4.659564 10.48044 74.00000
# Original - Follow-up: 3.332885 12.10712 38.40002
# Average:      8.796322 14.06368 38.40002
```

replicate.mean2	<i>Compares 2-group mean differences in original and follow-up studies</i>
-----------------	--

Description

This function computes confidence intervals for a 2-group mean difference from an original study and a follow-up study. Confidence intervals for the difference and average effect size also are computed. A Satterthwaite adjustment to the degrees of freedom is used to improve the accuracy of the confidence intervals. The same results can be obtained using the [meta.lc.mean2](#) function with appropriate contrast coefficients. The confidence level for the difference is $1 - 2\alpha$.

Usage

```
replicate.mean2(
  alpha,
  m11,
  m12,
  sd11,
  sd12,
  n11,
  n12,
  m21,
  m22,
  sd21,
```

```
sd22,
n21,
n22
)
```

Arguments

alpha	alpha level for 1-alpha confidence
m11	sample mean for group 1 in original study
m12	sample mean for group 2 in original study
sd11	sample SD for group 1 in original study
sd12	sample SD for group 2 in original study
n11	sample size for group 1 in original study
n12	sample size for group 2 in original study
m21	sample mean for group 1 in follow-up study
m22	sample mean for group 2 in follow-up study
sd21	sample SD for group 1 in follow-up study
sd22	sample SD for group 2 in follow-up study
n21	sample size for group 1 in follow-up study
n22	sample size for group 2 in follow-up study

Value

A 4-row matrix. The rows are:

- Row 1 summarizes the Original study
- Row 2 summarizes the Follow-up study
- Row 3 estimates the difference between studies
- Row 4 estimates the average effect size between the two studies

The columns are:

- Estimate - effect size estimate
- SE - standard error
- t - t-value
- p - p-value
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval
- df - degrees of freedom

References

Bonett DG (2021). "Design and analysis of replication studies." *Organizational Research Methods*, **24**(3), 513–529. ISSN 1094-4281, doi: [10.1177/1094428120911088](https://doi.org/10.1177/1094428120911088), <https://journals.sagepub.com/doi/10.1177/1094428120911088>.

Examples

```

replicate.mean2(.05, 21.9, 16.1, 3.82, 3.21, 40, 40,
                25.2, 19.1, 3.98, 3.79, 75, 75)

# Should return:
#           Estimate      SE      t      p
# Original:      5.80 0.7889312  7.3517180 1.927969e-10
# Follow-up:      6.10 0.6346075  9.6122408 0.000000e+00
# Original - Follow-up: -0.30 1.0124916 -0.2962988 7.673654e-01
# Average:        5.95 0.5062458 11.7531843 0.000000e+00
#           LL      UL      df
# Original:      4.228624 7.371376 75.75255
# Follow-up:      4.845913 7.354087 147.64728
# Original - Follow-up: -1.974571 1.374571 169.16137
# Average:        4.950627 6.949373 169.16137

```

replicate.plot

Plot to compare estimates from an original and follow-up study

Description

Generates a basic plot using ggplot2 to visualize the estimates from an original and follow-up study

Usage

```

replicate.plot(
  result,
  focus = c("Both", "Difference", "Average"),
  reference_line = NULL,
  diamond_height = 0.2,
  difference_axis_ticks = 5,
  ggtheme = ggplot2::theme_classic()
)

```

Arguments

result • a result matrix from any of the replicate functions in vcmeta

focus • Optional specification of the focus of the plot; defaults to 'Both'

- Both - a bit busy; plots each estimate, difference, and average
- Difference - plot each estimate and difference between them
- Average - plot each estimate and the average effect size

reference_line Optional x-value for a reference line. Only applies if focus is 'Difference' or 'Both'. Defaults to NULL, in which case a reference line is not drawn.

- diamond_height • Optional height of the diamond representing average effect size. Only applies if focus is 'Average' or 'Both'. Defaults to 0.2
- difference_axis_ticks • Optional requested number of ticks on the difference axis. Only applies if focus is 'Difference' or 'Both'. Defaults to 5.
- ggtheme • optional ggplot2 theme object; defaults to theme_classic()

Value

Returns a ggplot object. If stored, can be further customized via the ggplot API

Examples

```
# Compare Damisch et al., 2010 to Calin-Jageman & Caldwell 2014
# Damisch et al., 2010, Exp 1, German participants made 10 mini-golf putts
# Half were told they had a 'lucky' golf ball; half were not
# Found a large but uncertain improvement in shots made in the luck condition
# Calin-Jageman & Caldwell, 2014, Exp 1, was a pre-registered replication with
# input from Damisch, though with English-speaking participants
#
# Here we compare the effect sizes, in original units, for the two studies
# Use the replicate.mean2 function because the design is 2-group between-subj

library(ggplot2)
damisch_v_calinjageman_raw <- replicate.mean2(
  alpha = 0.05,
  m11 = 6.42,
  m12 = 4.75,
  sd11 = 1.88,
  sd12 = 2.15,
  n11 = 14,
  n12 = 14,
  m21 = 4.73,
  m22 = 4.62,
  sd21 = 1.958,
  sd22 = 2.12,
  n21 = 66,
  n22 = 58
)

# View the comparison:
damisch_v_calinjageman_raw

# Now plot the comparison, focusing on the difference
replicate.plot(damisch_v_calinjageman_raw, focus = "Difference")

# Plot the comparison, focusing on the average
replicate.plot(damisch_v_calinjageman_raw,
  focus = "Average",
  reference_line = 0,
```



```

    diamond_height = 0.1
  )

# Kind of busy, but plot the comparison with both difference and average
# In this case, store the plot for manipulation
myplot <- replicate.plot(
  damisch_v_calinjageman_raw,
  focus = "Both",
  reference_line = 0
)

# View the stored plot
myplot

# Change x-labels and study labels
myplot <- myplot + xlab("Difference in Putts Made, Lucky - Control")
myplot <- myplot + scale_y_discrete(
  labels = c(
    "Average",
    "Difference",
    "Calin-Jageman & Caldwell, 2014",
    "Damisch et al., 2010"
  )
)

# View the updated plot
myplot

```

replicate.stdmean.ps *Compares paired-samples standardized mean differences in original and follow-up studies*

Description

This function computes confidence intervals for a paired-samples standardized mean difference from an original study and a follow-up study. Confidence intervals for the difference and average effect size also are computed. The same results can be obtained using the [meta.lc.stdmean.ps](#) function with appropriate contrast coefficients. The confidence level for the difference is $1 - 2\alpha$.

Usage

```

replicate.stdmean.ps(
  alpha,
  m11,
  m12,
  sd11,
  sd12,

```

```

cor1,
n1,
m21,
m22,
sd21,
sd22,
cor2,
n2
)

```

Arguments

alpha	alpha level for 1-alpha confidence
m11	sample mean for group 1 in original study
m12	sample mean for group 2 in original study
sd11	sample SD for group 1 in original study
sd12	sample SD for group 2 in original study
cor1	sample correlation of paired observations in original study
n1	sample size in original study
m21	sample mean for group 1 in follow-up study
m22	sample mean for group 2 in follow-up study
sd21	sample SD for group 1 in follow-up study
sd22	sample SD for group 2 in follow-up study
cor2	sample correlation of paired observations in follow-up study
n2	sample size in follow-up study

Value

A 4-row matrix. The rows are:

- Row 1 summarizes the Original study
- Row 2 summarizes the Follow-up study
- Row 3 estimates the difference between studies
- Row 4 estimates the average effect size between the two studies

The columns are:

- Estimate - effect size estimate
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2021). “Design and analysis of replication studies.” *Organizational Research Methods*, 24(3), 513–529. ISSN 1094-4281, doi: [10.1177/1094428120911088](https://doi.org/10.1177/1094428120911088), <https://journals.sagepub.com/doi/10.1177/1094428120911088>.

Examples

```
replicate.stdmean.ps(
  alpha = .05,
  m11 = 86.22, m12 = 70.93, sd11 = 14.89, sd12 = 12.32, cor1 = .765, n1 = 20,
  m21 = 84.81, m22 = 77.24, sd21 = 15.68, sd22 = 16.95, cor2 = .702, n2 = 75
)

# Should return:
#           Estimate      SE      LL      UL
# Original:      1.0890300 0.22915553 0.6697353 1.5680085
# Follow-up:      0.4604958 0.09590506 0.2756687 0.6516096
# Original - Follow-up: 0.6552328 0.24841505 0.2466264 1.0638392
# Average:        0.7747629 0.12420752 0.5313206 1.0182052
```

replicate.stdmean2 *Compares 2-group standardized mean differences in original and follow-up studies*

Description

This function computes confidence intervals for a 2-group standardized mean difference from an original study and a follow-up study. Confidence intervals for the difference and average effect size also are computed. The same results can be obtained using the [meta.lc.stdmean2](#) function with appropriate contrast coefficients. The confidence level for the difference is $1 - 2\alpha$.

Usage

```
replicate.stdmean2(
  alpha,
  m11,
  m12,
  sd11,
  sd12,
  n11,
  n12,
  m21,
  m22,
  sd21,
  sd22,
  n21,
```

n22
)

Arguments

alpha	alpha level for 1-alpha confidence
m11	sample mean for group 1 in original study
m12	sample mean for group 2 in original study
sd11	sample SD for group 1 in original study
sd12	sample SD for group 2 in original study
n11	sample size for group 1 in original study
n12	sample size for group 2 in original study
m21	sample mean for group 1 in follow-up study
m22	sample mean for group 2 in follow-up study
sd21	sample SD for group 1 in follow-up study
sd22	sample SD for group 2 in follow-up study
n21	sample size for group 1 in follow-up study
n22	sample size for group 2 in follow-up study

Value

A 4-row matrix. The rows are:

- Row 1 summarizes the Original study
- Row 2 summarizes the Follow-up study
- Row 3 estimates the difference between studies
- Row 4 estimates the average effect size between the two studies

The columns are:

- Estimate - effect size estimate
- SE - standard error
- LL - lower limit of the confidence interval
- UL - upper limit of the confidence interval

References

Bonett DG (2021). "Design and analysis of replication studies." *Organizational Research Methods*, **24**(3), 513–529. ISSN 1094-4281, doi: [10.1177/1094428120911088](https://doi.org/10.1177/1094428120911088), <https://journals.sagepub.com/doi/10.1177/1094428120911088>.

Examples

```

replicate.stdmean2(.05, 21.9, 16.1, 3.82, 3.21, 40, 40,
                  25.2, 19.1, 3.98, 3.79, 75, 75)

# Should return:
#
#           Estimate      SE      LL      UL
# Original:      1.62803662 0.2594668 1.1353486 2.1524396
# Follow-up:      1.56170447 0.1870576 1.2030461 1.9362986
# Original - Follow-up: 0.07422178 0.3198649 -0.4519092 0.6003527
# Average:        1.59487055 0.1599325 1.2814087 1.9083324

```

se.cor

*Computes the standard error for a Pearson or partial correlation***Description**

This function can be used to compute the standard error of a Pearson or partial correlation using the sample correlation, sample size, and number of control variables. The effect size estimate and standard error output from this function can be used as input in the [meta.ave.gen](#), [meta.lc.gen](#), and [meta.lm.gen](#) functions in applications where a combination of different types of correlations are used in the meta-analysis.

Usage

```
se.cor(cor, q, n)
```

Arguments

cor	sample Pearson or partial correlation
q	number of control variables (0 for Pearson)
n	sample size

Value

Returns a one-row matrix:

- Estimate - estimate of Pearson or partial correlation
- SE - standard error

References

Bonett DG (2008). "Meta-analytic interval estimation for bivariate correlations." *Psychological Methods*, **13**(3), 173–181. ISSN 1939-1463, doi: [10.1037/a0012868](https://doi.org/10.1037/a0012868), <https://doi.apa.org/getdoi.cfm?doi=10.1037/a0012868>.

Examples

```
se.cor(.40, 0, 55)

# Should return:
#           Estimate      SE
# Correlation:    0.4 0.116487
```

se.mean.ps

Computes the standard error for a paired-samples mean difference

Description

This function can be used to compute the standard error of a paired-samples mean difference using the two sample means, sample effect size estimate and standard error output from this function can be used as input in the [meta.ave.gen](#), [meta.lc.gen](#), and [meta.lm.gen](#) functions in applications where compatible mean differences from a combination of 2-group and paired-samples experiments are used in the meta-analysis.

Usage

```
se.mean.ps(m1, m2, sd1, sd2, cor, n)
```

Arguments

m1	sample mean for measurement 1
m2	sample mean for measurement 2
sd1	sample standard deviation for measurement 1
sd2	sample standard deviation for measurement 2
cor	sample correlation for measurements 1 and 2
n	sample size

Value

Returns a one-row matrix:

- Estimate - estimate of mean difference
- SE - standard error

References

Snedecor GW, Cochran WG (1980). *Statistical methods*, 7th edition. ISU University Pres, Ames, Iowa.

Examples

```
se.mean.ps(23.9, 25.1, 1.76, 2.01, .78, 25)

# Should return:
#               Estimate      SE
# Mean difference:   -1.2 0.2544833
```

se.mean2

Computes the standard error for a 2-group mean difference

Description

This function can be used to compute the standard error of a 2-group mean difference using the two sample means, sample standard deviations, and sample sizes. The effect size estimate and standard error output from this function can be used as input in the [meta.ave.gen](#), [meta.lc.gen](#), and [meta.lm.gen](#) functions in applications where compatible mean differences from a combination of 2-group and paired-samples experiments are used in the meta-analysis.

Usage

```
se.mean2(m1, m2, sd1, sd2, n1, n2)
```

Arguments

m1	sample mean for group 1
m2	sample mean for group 2
sd1	sample standard deviation for group 1
sd2	sample standard deviation for group 2
n1	group 1 sample size
n2	group 2 sample size

Value

Returns a one-row matrix:

- Estimate - estimate of mean difference
- SE - standard error

References

Snedecor GW, Cochran WG (1980). *Statistical methods*, 7th edition. ISU University Pres, Ames, Iowa.

Examples

```
se.mean2(21.9, 16.1, 3.82, 3.21, 40, 40)
```

```
#           Estimate      SE
# Mean difference:      5.8 0.7889312
```

```
se.meanratio.ps      Computes the standard error for a paired-samples log mean ratio
```

Description

This function can be used to compute the standard error of a paired-samples mean ratio using the two sample means, sample effect size estimate and standard error output from this function can be used as input in the [meta.ave.gen](#), [meta.lc.gen](#), and [meta.lm.gen](#) functions in application where compatible mean ratios from a combination of 2-group and paired-samples experiments are used in the meta-analysis.

Usage

```
se.meanratio.ps(m1, m2, sd1, sd2, cor, n)
```

Arguments

m1	sample mean for measurement 1
m2	sample mean for measurement 2
sd1	sample standard deviation for measurement 1
sd2	sample standard deviation for measurement 2
cor	sample correlation for measurements 1 and 2
n	sample size

Value

Returns a one-row matrix:

- Estimate - estimate of log mean ratio
- SE - standard error

References

Bonett DG, Price RM (2020). “Confidence intervals for ratios of means and medians.” *Journal of Educational and Behavioral Statistics*, **45**(6), 750–770. ISSN 1076-9986, doi: [10.3102/1076998620934125](https://doi.org/10.3102/1076998620934125), <https://journals.sagepub.com/doi/10.3102/1076998620934125>.

Examples

```
se.meanratio.ps(21.9, 16.1, 3.82, 3.21, .748, 40)

# Should return:
#           Estimate      SE
# Log mean ratio: 0.3076674 0.02130161
```

se.meanratio2

Computes the standard error for a 2-group log mean ratio

Description

This function can be used to compute the standard error of a 2-group mean ratio using the two sample means, sample standard deviations, and sample sizes. The effect size estimate and standard error output from this function can be used as input in the [meta.ave.gen](#), [meta.lc.gen](#), and [meta.lm.gen](#) functions in application where compatible mean ratios from a combination of 2-group and paired-samples experiments are used in the meta-analysis.

Usage

```
se.meanratio2(m1, m2, sd1, sd2, n1, n2)
```

Arguments

m1	sample mean for group 1
m2	sample mean for group 2
sd1	sample standard deviation for group 1
sd2	sample standard deviation for group 2
n1	group 1 sample size
n2	group 2 sample size

Value

Returns a one-row matrix:

- Estimate - estimate of log mean ratio
- SE - standard error

References

Bonett DG, Price RM (2020). "Confidence intervals for ratios of means and medians." *Journal of Educational and Behavioral Statistics*, **45**(6), 750–770. ISSN 1076-9986, doi: [10.3102/1076998620934125](https://doi.org/10.3102/1076998620934125), <https://journals.sagepub.com/doi/10.3102/1076998620934125>.

Examples

```
se.meanratio2(21.9, 16.1, 3.82, 3.21, 40, 40)

# Should return:
#           Estimate      SE
# Log mean ratio: 0.3076674 0.041886
```

se.ods	<i>Computes the standard error for a log odds ratio</i>
--------	---

Description

This function computes a log odds ratio and its standard error using the frequency counts and sample sizes in a 2-group design. These frequency counts and sample sizes can be obtained from a 2x2 contingency table. This function is useful in a meta-analysis of odds ratios where some studies report the sample odds ratio and its standard error and other studies only report the frequency counts (or a 2x2 contingency table). The log odds ratio and standard error output from this function can be used as input in the [meta.ave.gen](#), [meta.lc.gen](#), and [meta.lm.gen](#) functions.

Usage

```
se.ods(f1, n1, f2, n2)
```

Arguments

f1	number of participants who have the outcome of interest in group 1
n1	group 1 sample size
f2	number of participants who have the outcome of interest in group 2
n2	group 2 sample size

Value

Returns a one-row matrix:

- Estimate - estimate of log odds ratio
- SE - standard error

References

Bonett DG, Price RM (2015). “Varying coefficient meta-analysis methods for odds ratios and risk ratios.” *Psychological Methods*, **20**(3), 394–406. ISSN 1939-1463, doi: [10.1037/met0000032](https://doi.org/10.1037/met0000032), <https://doi.apa.org/getdoi.cfm?doi=10.1037/met0000032>.

Examples

```
se.odds(36, 50, 21, 50)

# Should return:
#           Estimate      SE
# Log odds ratio: 1.239501 0.4204435
```

se.pbcor

Computes the standard error for a point-biserial correlation

Description

The function computes a point-biserial correlation and its standard error for two types of point-biserial correlations in 2-group designs using the sample means, sample standard deviations, and samples sizes. One type of point-biserial correlation uses an unweighted average of variances and is appropriate for 2-group experimental designs. The other type of point-biserial correlation uses an weighted average of variances and is appropriate for 2-group nonexperimental designs with simple random sampling. This function is useful in a meta-analysis of compatible point-biserial correlations where some studies used a 2-group experimental design and other studies used a 2-group nonexperimental design. The effect size estimate and standard error output from this function can be used as input in the [meta.ave.gen](#), [meta.lc.gen](#), and [meta.lm.gen](#) functions.

Usage

```
se.pbcor(m1, m2, sd1, sd2, n1, n2, type)
```

Arguments

m1	sample mean for group 1
m2	sample mean for group 2
sd1	sample standard deviation for group 1
sd2	sample standard deviation for group 2
n1	group 1 sample size
n2	group 2 sample size
type	<ul style="list-style-type: none"> • set to 1 for weighted variance average • set to 2 for unweighted variance average

Value

Returns a one-row matrix:

- Estimate - estimate of point-biserial correlation
- SE - standard error

References

Bonett DG (2020). "Point-biserial correlation: Interval estimation, hypothesis testing, meta-analysis, and sample size determination." *British Journal of Mathematical and Statistical Psychology*, **73**(S1), 113–144. ISSN 0007-1102, doi: [10.1111/bmsp.12189](https://doi.org/10.1111/bmsp.12189), <https://onlinelibrary.wiley.com/doi/10.1111/bmsp.12189>.

Examples

```
se.pbcor(21.9, 16.1, 3.82, 3.21, 40, 40, 1)

# Should return:
#               Estimate      SE
# Point-biserial correlation: 0.6349786 0.05981325
```

<code>se.semipartial</code>	<i>Computes the standard error for a semipartial correlation</i>
-----------------------------	--

Description

This function can be used to compute the standard error of a semipartial correlation using the sample correlation, sample size, and squared multiple correlation for the full model. The effect size estimate and standard error output from this function can be used as input in the [meta.ave.gen](#), [meta.lc.gen](#), and [meta.lm.gen](#) functions in applications where a combination of different types of correlations are used in the meta-analysis.

Usage

```
se.semipartial(cor, r2, n)
```

Arguments

<code>cor</code>	sample semipartial correlation
<code>r2</code>	squared multiple correlation for full model
<code>n</code>	sample size

Value

Returns a one-row matrix:

- Estimate - estimate of semipartial correlation
- SE - standard error

Examples

```
se.semipartial(.40, .25, 60)

# Should return:
#               Estimate      SE
# Semipartial correlation:    0.4 0.1063262
```

se.slope	<i>Computes a slope and standard error</i>
----------	--

Description

This function can be used to compute a slope and its standard error for a simple linear regression model using the sample Pearson correlation and the standard deviations of response and predictor variables. This function is useful in a meta-analysis of slopes of a simple linear regression model where some studies report the Pearson correlation but not the slope.

Usage

```
se.slope(cor, sdy, sdx, n)
```

Arguments

cor	sample Pearson correlation
sdy	sample standard deviation of the response variable
sdx	sample standard deviation of the predictor variable
n	sample size

Value

Returns a one-row matrix:

- Estimate - estimate of slope
- SE - standard error

References

Snedecor GW, Cochran WG (1980). *Statistical methods*, 7th edition. ISU University Pres, Ames, Iowa.

Examples

```
se.slope(.392, 4.54, 2.89, 60)

# Should return:
#           Estimate      SE
# Slope: 0.6158062 0.1897647
```

```
se.spear
```

Computes the standard error for a Spearman correlation

Description

This function can be used to compute the standard error of a Spearman correlation using the sample correlation and sample size. The standard error from this function can be used as input in the [meta.ave.gen](#), [meta.lc.gen](#), and [meta.lm.gen](#) functions in applications where a combination of different types of correlations are used in the meta-analysis.

Usage

```
se.spear(cor, n)
```

Arguments

cor	sample Spearman correlation
n	sample size

Value

Returns a one-row matrix:

- Estimate - estimate of Spearman correlation
- SE - standard error

References

Bonett DG, Wright TA (2000). "Sample size requirements for estimating Pearson, Kendall and Spearman correlations." *Psychometrika*, **65**(1), 23–28. ISSN 0033-3123, doi: [10.1007/BF02294183](https://doi.org/10.1007/BF02294183), <https://link.springer.com/10.1007/BF02294183>.

Examples

```
se.spear(.40, 55)

# Should return:
#           Estimate      SE
# Spearman correlation: 0.4 0.1210569
```

se.stdmean.ps	<i>Computes the standard error for a paired-samples standardized mean difference</i>
---------------	--

Description

The effect size estimate and standard error output from this function can be used as input in the [meta.ave.gen](#), [meta.lc.gen](#), and [meta.lm.gen](#) functions in applications where compatible standardized mean differences from a combination of 2-group and paired-samples experiments are used in the meta-analysis.

Usage

```
se.stdmean.ps(m1, m2, sd1, sd2, cor, n, stdzr)
```

Arguments

m1	sample mean for measurement 1
m2	sample mean for measurement 2
sd1	sample standard deviation for measurement 1
sd2	sample standard deviation for measurement 2
cor	sample correlation for measurements 1 and 2
n	sample size
stdzr	<ul style="list-style-type: none"> • set to 0 for square root average variance standardizer • set to 1 for group 1 SD standardizer • set to 2 for group 2 SD standardizer • set to 3 for square root weighted variance standardizer

Value

Returns a one-row matrix:

- Estimate - estimate of standardized mean difference
- SE - standard error

References

Bonett DG (2009). "Meta-analytic interval estimation for standardized and unstandardized mean differences." *Psychological Methods*, **14**(3), 225–238. ISSN 1939-1463, doi: [10.1037/a0016619](https://doi.org/10.1037/a0016619), <https://doi.apa.org/getdoi.cfm?doi=10.1037/a0016619>.

Examples

```
se.stdmean.ps(23.9, 25.1, 1.76, 2.01, .78, 25, 0)

# Should return:
#               Estimate      SE
# Standardizedd mean difference: -0.6352097 0.1602852
```

```
se.stdmean2           Computes the standard error for a 2-group standardized mean differ-
                       ence
```

Description

Use the square root average variance standardizer (stdzr = 0) for 2-group experimental designs. Use the square root weighted variance standardizer (stdzr = 3) for 2-group nonexperimental designs with simple random sampling. The single-group standardizers (stdzr = 1 and stdzr = 2) can be used with either 2-group experimental or nonexperimental designs. The effect size estimate and standard error output from this function can be used as input in the [meta.ave.gen](#), [meta.lc.gen](#), and [meta.lm.gen](#) functions in applications where compatible standardized mean differences from a combination of 2-group and paired-samples experiments are used in the meta-analysis.

Usage

```
se.stdmean2(m1, m2, sd1, sd2, n1, n2, stdzr)
```

Arguments

m1	sample mean for group 1
m2	sample mean for group 2
sd1	sample standard deviation for group 1
sd2	sample standard deviation for group 2
n1	group 1 sample size
n2	group 2 sample size
stdzr	<ul style="list-style-type: none"> • set to 0 for square root average variance standardizer • set to 1 for group 1 SD standardizer • set to 2 for group 2 SD standardizer • set to 3 for square root weighted variance standardizer

Value

Returns a one-row matrix:

- Estimate - estimate of standardized mean difference
- SE - standard error

References

Bonett DG (2009). "Meta-analytic interval estimation for standardized and unstandardized mean differences." *Psychological Methods*, **14**(3), 225–238. ISSN 1939-1463, doi: [10.1037/a0016619](https://doi.org/10.1037/a0016619), <https://doi.org/10.1037/a0016619>.

Examples

```
se.stdmean2(21.9, 16.1, 3.82, 3.21, 40, 40, 0)

# Should return:
#               Estimate      SE
# Standardized mean difference: 1.643894 0.2629049
```

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