

Package ‘PINMA’

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Type Package

Title Improved Methods for Constructing Prediction Intervals for Network Meta-Analysis

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Description Parametric bootstrap and Kenward-Roger-type adjustment methods to construct prediction intervals for network meta-analysis are implementable.

Depends R (>= 3.5.0)

Imports stats, MASS, metafor

License GPL-3

Encoding UTF-8

LazyData true

RoxygenNote 7.1.1

R topics documented:

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PINMA-package	<i>The ‘PINMA’ package.</i>
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Description

Improved Methods for Constructing Prediction Intervals for Network Meta-Analysis.

References

Noma, H., Hamura, Y., Sugawara, S. and Furukawa, T. A. (2022+). Improved methods to construct prediction intervals for network meta-analysis. Forthcoming.

data.edit	<i>Transforming arm-level data to contrast-based summary statistics</i>
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Description

Transforming arm-level data to contrast-based summary statistics.

Usage

```
data.edit(study, trt, d, n)
```

Arguments

study	Study ID
trt	Numbered treatment (=1,2,...)
d	Number of events
n	Sample size

Value

Contrast-based summary statistics are generated.

- y: Contrast-based summary estimates.
- S: Vektored within-study covariance matrix.

Examples

```
data(dstr)
attach(dstr)

edat <- data.edit(study, trt, d, n)
```

dstr	<i>Siontis et al. (2018)'s network meta-analysis data</i>
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Description

- study: Study ID
- treat: Treatment
- trt: Numbered treatment (1:CCTA, 2:CMR, 3:exercise ECG, 4:SPECT-MPI, 5:standard care, 6:Stress Echo)
- n: Sample size
- d: Number of events

Usage

```
data(dstr)
```

Format

A arm-based dataset with 29 rows and 5 variables

References

Siontis, G. C., Mavridis, D., Greenwood, J. P., et al. (2018). Outcomes of non-invasive diagnostic modalities for the detection of coronary artery disease: network meta-analysis of diagnostic randomised controlled trials. *BMJ*. **360**: k504.

KR	<i>Kenward-Roger-type adjustment for constructing prediction intervals of network meta-analysis</i>
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Description

Kenward-Roger-type adjustment for constructing prediction intervals of network meta-analysis.

Usage

```
KR(y, S)
```

Arguments

y	Contrast-based summary data of the outcome measure
S	Covariance estimates of y

Value

Results of the Kenward-Roger-type adjustment for inference of multivariate random-effects model and prediction intervals for network meta-analysis.

- Estimates: Restricted maximum likelihood (REML) estimates, their SE, and Wald-type 95% confidence intervals by the Kenward-Roger-type adjustment.
- Between-studies_SD: Between-studies SD estimate.
- 95%PI: 95% prediction intervals by the Kenward-Roger-type adjustment.

References

Noma, H., Hamura, Y., Sugawara, S. and Furukawa, T. A. (2022+). Improved methods to construct prediction intervals for network meta-analysis. Forthcoming.

Examples

```
data(dstr)
attach(dstr)

# Transforming the arm-level data to the contrast-based summaries
edat <- data.edit(study,trt,d,n)

y <- edat$y
S <- edat$S

KR(y,S)    # Results of the NMA analysis (log OR scale)
```

PBS	<i>Parametric bootstrap procedure for constructing prediction intervals of network meta-analysis</i>
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Description

Parametric bootstrap procedure for constructing prediction intervals of network meta-analysis.

Usage

```
PBS(y, S, B=2000)
```

Arguments

y	Contrast-based summary data of the outcome measure
S	Covariance estimates of y
B	Number of bootstrap resampling (default: 2000).

Value

The parametric bootstrap prediction intervals for network meta-analysis.

- Estimates: Restricted maximum likelihood (REML) estimates, their SE, and 95% Wald-type confidence intervals.
- Between-studies_SD: Between-studies SD estimate.
- 95%PI: 95% prediction intervals by the parametric bootstrap.

References

Noma, H., Hamura, Y., Sugawara, S. and Furukawa, T. A. (2022+). Improved methods to construct prediction intervals for network meta-analysis. Forthcoming.

Examples

```
data(dstr)
attach(dstr)

# Transforming the arm-level data to the contrast-based summaries
edat <- data.edit(study,trt,d,n)

y <- edat$y
S <- edat$S

PBS(y,S) # Results of the NMA analysis (log OR scale)
```

t.PI	<i>The ordinary t-approximation for constructing prediction intervals of network meta-analysis</i>
------	--

Description

The ordinary t-approximation for constructing prediction intervals of network meta-analysis.

Usage

```
t.PI(y, S)
```

Arguments

y	Contrast-based summary data of the outcome measure
S	Covariance estimates of y

Value

The ordinary t-approximation prediction intervals for network meta-analysis.

- Estimates: Restricted maximum likelihood (REML) estimates, their SE, and Wald-type 95% confidence intervals.
- Between-studies_SD: Between-studies SD estimate.
- 95%PI: 95% prediction intervals by the ordinary t-approximation.

References

Cooper, H., Hedges, L. V., and Valentine, J. C. (2009). *The Handbook of Research Synthesis and Meta-Analysis*, 2nd edition. New York: Russell Sage Foundation.

Chaimani, A., and Salanti, G. (2015). Visualizing assumptions and results in network meta-analysis: the network graphs package. *Stata Journal* **15**, 905-920.

Examples

```
data(dstr)
attach(dstr)

# Transforming the arm-level data to the contrast-based summaries
edat <- data.edit(study,trt,d,n)

y <- edat$y
S <- edat$S

t.PI(y,S)  # Results of the NMA analysis (log OR scale)
```

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