Bivariate meta-analysis with insufficient reporting of the correlation between outcomes on the study level

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Basis: the univariate random-effects meta-analysis model

- o data:
	- effect estimates y_i $(i = 1, \ldots, k)$
	- standard errors s_{i1} (known, fixed)
- normal-normal hierarchical model (NNHM):

$$
y_i|\theta_i, \sigma_i \sim \text{Normal}(\theta_i, s_i^2)
$$

$$
\theta_i|\mu, \tau \sim \text{Normal}(\mu, \tau^2)
$$

• marginally:

$$
y_i|\mu, \tau, \sigma_i \sim \text{Normal}(\mu, s_i^2 + \tau^2)
$$

- parameters:
	- overall mean effect μ
	- heterogeneity τ
	- (study-specific means θ_i)

Bivariate model: motivation

- sometimes two (or more) similar/related effect estimates per study, examples:
	- overall survival / disease-free survival
	- \bullet pain relief / pain free
	- different symptom scales
	- \bullet ...
- both may be reported by all or some studies
- use of additional data may improve estimation and broaden evidence base¹
- usually: (within-study) correlations between endpoints required

 1 e.g.: R.D. Riley, K.R. Abrams, P.C. Lambert, A.J. Sutton, J.R. Thompson. An evaluation of bivariate random-effects meta-analysis for the joint synthesis of two correlated outcomes. Statistics in Medicine, 26(1):78-97, 2007.

Bivariate generalization A: known correlations (1)

- data:
	- bivariate effect estimates $Y_i = (y_{i1}, y_{i2})'$ $(i = 1, ..., k)$
	- pairs of standard errors s_{i1} , s_{i2} (known, fixed)
	- (within-study-) correlations r_i (known, fixed)
- **•** bivariate generalization:

$$
Y_i | \Theta_i, \Sigma_i \sim \text{Normal}\left(\Theta_i = \begin{pmatrix} \theta_{i1} \\ \theta_{i2} \end{pmatrix}, \Sigma_i = \begin{pmatrix} s_{i1}^2 & r_i s_{i1} s_{i2} \\ r_i s_{i1} s_{i2} & s_{i2}^2 \end{pmatrix} \right),
$$

$$
\Theta_i | \mu, T \sim \text{Normal}\left(\mu = \begin{pmatrix} \mu_1 \\ \mu_2 \end{pmatrix}, T = \begin{pmatrix} \tau_1^2 & \varrho_B \tau_1 \tau_2 \\ \varrho_B \tau_1 \tau_2 & \tau_2^2 \end{pmatrix} \right)
$$

• marginally:

$$
Y_i|\mu, \Sigma_i, T \dots \sim \text{Normal}\left(\mu = \left(\begin{array}{c} \mu_1 \\ \mu_2 \end{array}\right), \Lambda_i = \left(\begin{array}{cc} s_{i1}^2 + \tau_1^2 & \lambda_{i;1,2} \\ \lambda_{i;2,1} & s_{i2}^2 + \tau_2^2 \end{array}\right)\right)
$$

where $\Lambda_i = \Sigma_i + T$, and the covariance term is

$$
\lambda_{i;1,2} = \lambda_{i;2,1} = r_i s_{i1} s_{i2} + \varrho_B \tau_1 \tau_2
$$

Bivariate generalization A: known correlations (2)

• marginally:

$$
Y_i|\mu, \Sigma_i, T \ldots \sim \text{Normal}\left(\mu = \left(\begin{array}{c} \mu_1 \\ \mu_2 \end{array}\right), \Lambda_i = \left(\begin{array}{cc} s_{i1}^2 + \tau_1^2 & \lambda_{i;1,2} \\ \lambda_{i;2,1} & s_{i2}^2 + \tau_2^2 \end{array}\right)\right)
$$

where $\Lambda_i = \Sigma_i + T$, and the covariance term is

$$
\lambda_{i;1,2} = \lambda_{i;2,1} = r_i s_i s_i s_i + \varrho_B \tau_1 \tau_2
$$

- parameters:
	- (effects μ_1, μ_2 , heterogeneities τ_2, τ_2 , as in univariate case)
	- between-study correlation ρ_B
- required: within-study correlations r_i
- \bullet (what if the r_i are not provided?)

Bivariate generalization B: common-correlation model

marginal model:

$$
Y_i|\mu, \Sigma_i, T \dots \sim \text{Normal}\left(\mu = \begin{pmatrix} \mu_1 \\ \mu_2 \end{pmatrix}, \Lambda_i = \begin{pmatrix} s_{i1}^2 + \tau_1^2 & \lambda_{i;1,2} \\ \lambda_{i;2,1} & s_{i2}^2 + \tau_2^2 \end{pmatrix}\right)
$$

• treat (within-study) correlation as single, common parameter ϱ_W :

$$
\lambda_{i;1,2} = \lambda_{i;2,1} = \varrho_{\mathsf{W}} s_{i1} s_{i2} + \varrho_{\mathsf{B}} \tau_1 \tau_2
$$

• parameters:

- (effects μ_1 , μ_2 , heterogeneities τ_2 , τ_2 , as in univariate case)
- \bullet between-study correlation ρ_B
- \bullet within-study correlation ρ_w

Bivariate generalization C: random-correlation model

• marginal model:

$$
Y_i|\mu, \Sigma_i, T \dots \sim \text{Normal}\left(\mu = \begin{pmatrix} \mu_1 \\ \mu_2 \end{pmatrix}, \Lambda_i = \begin{pmatrix} s_{i1}^2 + \tau_1^2 & \lambda_{i;1,2} \\ \lambda_{i;2,1} & s_{i2}^2 + \tau_2^2 \end{pmatrix}\right)
$$

treat correlation as <mark>random effec</mark>t $\varrho_{\mathsf{W}i}\!\!$:

$$
\lambda_{i;1,2} = \lambda_{i;2,1} = \varrho_{W_i} s_{i1} s_{i2} + \varrho_B \tau_1 \tau_2
$$

where

$$
\text{atanh}(\varrho_{W_i}) \sim \text{Normal}(\mu_W, \sigma_W^2)
$$

- parameters:
	- (effects μ_1, μ_2 , heterogeneities τ_2, τ_2 , as in univariate case)
	- \bullet between-study correlation ρ_B
	- (mean) within-study correlation tanh(μ_W)
	- within-study correlation heterogeneity σ_w

Bivariate generalization C: random-correlation model

• marginal model:

$$
Y_i|\mu, \Sigma_i, T \dots \sim \text{Normal}\left(\mu = \begin{pmatrix} \mu_1 \\ \mu_2 \end{pmatrix}, \Lambda_i = \begin{pmatrix} s_{i1}^2 + \tau_1^2 & \lambda_{i;1,2} \\ \lambda_{i;2,1} & s_{i2}^2 + \tau_2^2 \end{pmatrix}\right)
$$

treat correlation as <mark>random effec</mark>t $\varrho_{\mathsf{W}i}\!\!$:

$$
\lambda_{i;1,2} = \lambda_{i;2,1} = \varrho_{W_i} s_{i1} s_{i2} + \varrho_B \tau_1 \tau_2
$$

where

$$
\text{atanh}(\varrho_{\text{W}i}) \sim \text{Normal}(\mu_{\text{W}}, \sigma_{\text{W}}^2)^*
$$

- parameters:
	- (effects μ_1, μ_2 , heterogeneities τ_2, τ_2 , as in univariate case)
	- \bullet between-study correlation ρ_B
	- (mean) within-study correlation tanh(μ_W)
	- within-study correlation heterogeneity σ_w

∗ (NB: choice of atanh ("Fisher-z") transform is somewhat ad hoc here)

Bivariate generalization D: alternative model due to Riley, Thompson and Abrams (2008)

• marginal model:

$$
Y_i|\mu, \Sigma_i, T \ldots \sim \text{Normal}\left(\mu = \left(\begin{array}{c} \mu_1 \\ \mu_2 \end{array}\right), \Lambda_i = \left(\begin{array}{cc} s_{i1}^2 + \tau_1^2 & \lambda_{i;1,2} \\ \lambda_{i;2,1} & s_{i2}^2 + \tau_2^2 \end{array}\right)\right)
$$

treat correlations via a $\boldsymbol{\mathrm{single}},$ $\boldsymbol{\mathrm{common}}$ parameter ρ : 2

$$
\lambda_{i;1,2} = \lambda_{i;2,1} = \rho \sqrt{(s_{i1}^2 + \tau_1^2)(s_{i2}^2 + \tau_2^2)}
$$

- parameters:
	- (effects μ_1, μ_2 , heterogeneities τ_2, τ_2 , as in univariate case)
	- \bullet overall correlation $ρ$
- Notes:
	- originally proposed in frequentist context
	- *ρ* mimics $ρ_W$ for "small" $τ₁, τ₂; ρ$ mimics $ρ_B$ for "large" $τ₁, τ₂$
	- shrinkage estimation (of Θ_i) or prediction (of Θ_{k+1}) not possible.

 2 R.D. Riley *et al.* An alternative model for bivariate random-effects meta-analysis when the within-study correlations are unknown. Biostatistics, 9(1):172–186, 2008.

C. R¨over [Bivariate meta-analysis . . .](#page-0-0) August 28, 2024 8 / 16

Bivariate generalization: Four models

$$
Y_i|\mu, \Sigma_i, T \ldots \sim \text{Normal}\left(\mu = \left(\begin{array}{c} \mu_1 \\ \mu_2 \end{array}\right), \Lambda_i = \left(\begin{array}{cc} s_{i1}^2 + \tau_1^2 & \lambda_{i;1,2} \\ \lambda_{i;2,1} & s_{i2}^2 + \tau_2^2 \end{array}\right)\right)
$$

(A) known correlations model:

$$
\lambda_{i;1,2} = \lambda_{i;2,1} = r_i s_{i1} s_{i2} + \varrho_B \tau_1 \tau_2
$$

(B) common effect model:

$$
\lambda_{i;1,2} = \lambda_{i;2,1} = \varrho_{\mathsf{W}} s_{i1} s_{i2} + \varrho_{\mathsf{B}} \tau_1 \tau_2
$$

(C) random effects model:

$$
\lambda_{i;1,2} = \lambda_{i;2,1} = \varrho_{W_i} s_{i1} s_{i2} + \varrho_B \tau_1 \tau_2
$$

(D) RTA model

$$
\lambda_{i;1,2} = \lambda_{i;2,1} = \rho \sqrt{(s_{i1}^2 + \tau_1^2)(s_{i2}^2 + \tau_2^2)}
$$

Bivariate generalization: prior specification

- to consistently generalize from univariate case, use separation approach³:
	- specify priors for μ_1 , μ_2 , τ_1 , τ_2 "as usual"
	- specify priors for additional correlation parameters
- "vague" priors for effects (μ_1, μ_2)
- weakly informative priors for heterogeneities (τ_1,τ_2) 4
- priors for correlation parameters: Uniform $[-1, 1]$ or *arcsine prior*⁵

 3 D.L. Burke *et al.* Bayesian bivariate meta-analysis of correlated effects: Impact of the prior distributions on the between-study correlation, borrowing of strength, and joint inferences. Statistical Methods in Medical Research, 27(2):428–450, 2018.

⁴C. Röver, R. Bender, S. Dias, C.H. Schmid, H. Schmidli, S. Sturtz, S. Weber, T. Friede. On weakly informative prior distributions for the heterogeneity parameter in Bayesian random-effects meta-analysis. Research Synthesis Methods, 12(4):448–474, 2021.

 5 B.K. Fosdick, A.E. Raftery. Estimating the correlation in bivariate normal data with known variances and small sample sizes. The American Statistician, 66(1):34–41, 2012.

- four models implemented in R (using JAGS)
- corresponding univariate models $(\lambda_{i:1,2} = \lambda_{i:2,1} = 0.0)$: using bayesmeta package
- estimation of overall means, heterogeneity, correlations (ρ_B , ρ_W , μ_W , σ_W , ρ , ...), study-specific effects $(\theta_i,$ shrinkage estimates)
- demonstrate / compare performance in example

Blood pressure data

- \bullet blood pressure data set 6
- two correlated endpoints: drug effects (mean difference, mmHg) on systolic and diastolic blood pressure (SBP, DBP)
- ²¹ studies
- based on external evidence, within-study correlation (r_i) had also been fixed at $r_i = 0.71$.

 6 C. Geeganage and P.M.W. Bath. Vasoactive drugs for acute stroke. Cochrane Database of Systematic Reviews, 7:1465–1858, 2010.

Y. Wei and J.P.T. Higgins. Estimating within-study covariances in multivariate meta-analysis with multiple outcomes. Statistics in Medicine, 32(7):1191–1205, 2013.

Effect estimates

 \bullet overall effect estimates very similar

Effect estimates

- o overall effect estimates very similar
- some precision gain once correlations are considered; \bullet CI widths compared to univariate analyses:

Correlation estimates

- **o** different correlation parameters shown in common plots
- broad agreement between models
- RE variance (C): σ_W posterior close to prior

Shrinkage estimation

- consideration of correlations particularly useful if only one of two endpoints is given
- allows (e.g.) prediction of 2nd endpoint (θ_{i2}) given the 1st (y_{i1} , s_{i1})
- consider (constructed) case of one missing endpoint: no DBP data for recent "PRISTINE" study

- vague prediction based on 20 remaining DBP estimates alone (univariate MA)
- more precise predictions (θ_{i2}) based on 20 DBP + 21 SBP estimates (PRISTINE's SBP estimate (y_{i1}) was also above average)

Discussion

- model setup:
	- fixed-effect model sensible / pragmatic? (RE variance seems poorly constrained, may require lots of data)
	- atanh ("Fisher-z") transform for 2nd stage sensible? (Motivation? Alternatives?)
	- noticeable advantages also for "simple" RTA model
- prior choice
	- \bullet besides "usual" parameters $-$ correlation priors required
	- "arcsine" prior mimics Jeffreys prior, emphasizes larger (absolute) correlations
	- uniform prior as a "conservative" alternative
	- alternatives, if, e.g., only positive correlations are expected?
- advantages: precision gain (even for overall means), opportunity to jointly analyze similar/different endpoints
- when can we expect advantages for overall mean or shrinkage estimates? (large number of studies, high correlation or large heterogeneity required?)

+++ additional slides +++

systolic (SBP)

 \bullet plain \bullet shrinkage

−30 −10 0 10 20 MD (mmHg)

diastolic (DBP)

 \bullet plain \bullet shrinkage

−20 −10 0 10 MD (mmHg)

Blood pressure data

• posteriors for effects ($\mu_{\rm SBP}$, $\mu_{\rm DBP}$) and heterogeneities ($\tau_{\rm SBP}$, $\tau_{\rm DBP}$)

