

Bayesian hypothesis testing and estimation under the marginalized random-effects meta-analysis model

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Bayesian meta-analysis

- ▶ Meta-analysis literature mainly focused on empirical Bayes and fully Bayesian estimation
- ▶ Bayes factors can be used for Bayesian hypothesis testing
- ▶ A Bayes factor quantifies the evidence for one model relative to a contrasting model

$$B_{12} = \frac{m_1(\mathbf{y})}{m_2(\mathbf{y})}$$

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- ▶ Meta-analysis literature mainly focused on empirical Bayes and fully Bayesian estimation
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$$B_{12} = \frac{m_1(\mathbf{y})}{m_2(\mathbf{y})}$$

- ▶ Existing meta-analytic Bayes factors either focus on a single parameter or are effect size measure dependent [1–4]
- ▶ **Goal:** Proposing a methodology for Bayesian estimation *and* hypothesis testing that can be used for any effect size measure

MAREMA model

- ▶ We use the marginalized random-effects meta-analysis (MAREMA) model,

$$y_i \sim N(\mu, \sigma_i^2 + \tau^2)$$

- ▶ The MAREMA model encompasses three meta-analysis models:
 - ▶ Equal-effect model \rightarrow zero between-study variance
 - ▶ Random-effects model \rightarrow positive between-study variance
 - ▶ Model with a negative between-study variance
- ▶ A negative between-study variance is not uncommon [5] and may be caused by chance or dependencies among the studies

Estimation: Prior distributions

- ▶ A prior distribution is not placed on τ^2 but on the I^2 -statistic
 $\rightarrow I^2 = \tau^2 / (\tau^2 + \tilde{\sigma}^2)$
- ▶ Reparameterizing the MAREMA model using the I^2 -statistic and replacing it with ρ yields

$$y_i \sim N\left(\mu, \sigma_i^2 + \tilde{\sigma}^2 \rho / (1 - \rho)\right)$$

- ▶ The smallest possible value of ρ is a function of the smallest sampling variance (i.e., σ_{min}^2)

$$\rho_{min} = \frac{-\sigma_{min}^2}{-\sigma_{min}^2 + \tilde{\sigma}^2}$$

Estimation: Prior distributions

- ▶ Flat prior distributions are used:

$$\pi(\mu, \rho) = \pi(\mu)\pi(\rho), \text{ with}$$

$$\pi(\mu) \propto \mathbf{1}$$

$$\pi(\rho) = U(\rho_{min}, 1)$$

- ▶ Posterior distributions are obtained using a Gibbs sampler

Estimation: Prior distributions

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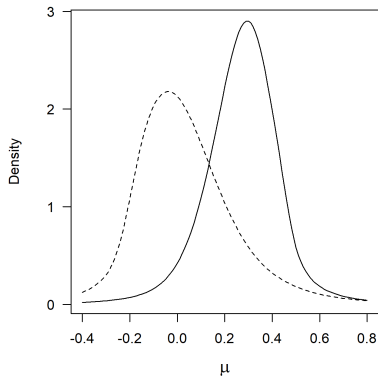
$$\pi(\mu) \propto 1$$

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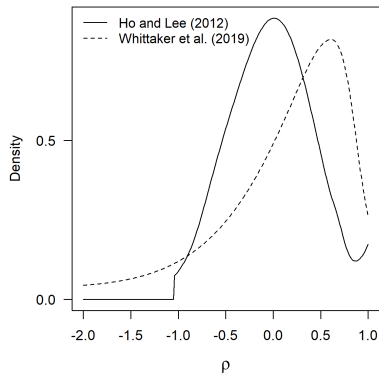
- ▶ Posterior distributions are obtained using a Gibbs sampler
- ▶ Illustrating estimation using two examples:
 - ▶ Ho et al. [6] contains 10 standardized mean differences on the efficacy of EMDR vs. CBT therapy to treat PTSD
 - ▶ Whittaker et al. [7] contains 3 log risk ratio on the difference between using a smartphone app and lower intensity support to quit smoking

Application: Posterior distributions

Posterior distribution of μ



Posterior distribution of ρ



Application: Parameter estimates

Ho et al.: [6]

	μ		ρ	
	Estimate	95% CI/CrI	Estimate	95% CI/CrI
MAREMA	0.274 (0.327)	(-0.109;0.638)	-0.026 (-0.016)	(-0.837;0.812)
Frequentist	0.249	(-0.003;0.502)	0.022	(0;0.747)

Whittaker et al.: [7]

	μ		ρ	
	Estimate	95% CI/CrI	Estimate	95% CI/CrI
MAREMA	0.033 (0.043)	(-0.413;0.625)	0.089 (0.597)	(-1.752;0.922)
Frequentist	0.114	(-0.525;0.753)	0.696	(0;0.993)

Bayes factors: Prior distributions

- ▶ In the two examples, we test these hypotheses:

$$H_0 : \mu = 0$$

$$H_1 : \mu < 0$$

$$H_2 : \mu > 0$$

$$H_0 : \rho = 0$$

$$H_1 : \rho < 0$$

$$H_2 : \rho > 0$$

- ▶ A proper prior is needed for Bayes factors, so we cannot use the flat prior for μ
- ▶ We propose a unit-information prior for μ and a uniform prior for ρ under the unconstrained MAREMA model:

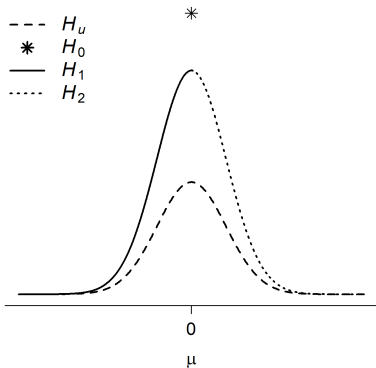
$$\pi_u(\mu, \rho) = \pi_u(\mu|\rho)\pi_u(\rho), \text{ with}$$

$$\pi_u(\mu|\rho) = N(\mu, k(\mathbf{1}'\sum_{\rho}^{-1}\mathbf{1})^{-1})$$

$$\pi(\rho) = U(\rho_{min}, 1)$$

Bayes factors: Prior distributions

Priors specified for μ



Priors specified for ρ

Bayes factors: Computation

- ▶ Marginal likelihoods of the different hypotheses are needed to compute the Bayes factor
- ▶ For example, the marginal likelihood of $H_1 : \mu < 0$ is

$$m_1(\mathbf{y}) = \iint_{\mu < 0} f(\mathbf{y}|\mu, \rho)\pi_1(\mu, \rho)d\mu d\rho$$

- ▶ Marginal likelihoods were approximated using importance sampling or a random walk procedure

Application: Bayes factors

Ho et al.: [6]

	μ			ρ		
	H_0	H_1	H_2	H_0	H_1	H_2
H_0	1.000	4.183	0.265	1.000	3.977	4.979
H_1	0.239	1.000	0.063	0.251	1.000	1.252
H_2	3.779	15.810	1.000	0.201	0.799	1.000
$P(H_q \mathbf{y})$	0.199	0.048	0.753	0.689	0.173	0.138

Note: $H_0 : \mu = 0$; $H_1 : \mu < 0$; $H_2 : \mu > 0$

- ▶ $H_2 : \mu > 0$ is most likely compared to H_0 and H_1
- ▶ Frequentist test: $z = 1.936$, $p = 0.053$

Application: Bayes factors

Ho et al.: [6]

	μ			ρ		
	H_0	H_1	H_2	H_0	H_1	H_2
H_0	1.000	4.183	0.265	1.000	3.977	4.979
H_1	0.239	1.000	0.063	0.251	1.000	1.252
H_2	3.779	15.810	1.000	0.201	0.799	1.000
$P(H_q \mathbf{y})$	0.199	0.048	0.753	0.689	0.173	0.138

Note: $H_0 : \rho = 0$; $H_1 : \rho < 0$; $H_2 : \rho > 0$

- ▶ $H_0 : \rho = 0$ is most likely compared to H_1 and H_2
- ▶ Frequentist test: $Q(9) = 9.417$, $p = 0.400$

Application: Bayes factors

Whittaker et al.: [7]

	μ			ρ		
	H_0	H_1	H_2	H_0	H_1	H_2
H_0	1.000	2.558	2.115	1.000	10.958	2.901
H_1	0.391	1.000	0.827	0.091	1.000	0.265
H_2	0.473	1.209	1.000	0.345	3.778	1.000
$P(H_q \mathbf{y})$	0.537	0.210	0.254	0.696	0.064	0.240

Note: $H_0 : \mu = 0$; $H_1 : \mu < 0$; $H_2 : \mu > 0$

- ▶ $H_0 : \mu = 0$ is most likely compared to H_1 and H_2 but no strong evidence
- ▶ Frequentist test: $z = 0.349$, $p = 0.727$

Whittaker et al.: [7]

	μ			ρ		
	H_0	H_1	H_2	H_0	H_1	H_2
H_0	1.000	2.558	2.115	1.000	10.958	2.901
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Note: $H_0 : \rho = 0$; $H_1 : \rho < 0$; $H_2 : \rho > 0$

- ▶ $H_0 : \rho = 0$ is most likely compared to H_1 and H_2
- ▶ Frequentist test: $Q(2) = 6.240$, $p = 0.044$

- ▶ The proposed Bayesian estimation and hypothesis testing is novel, because
 - ▶ It is based on the MAREMA model
 - ▶ A prior is placed on ρ (i.e., I^2 -statistic) rather than on τ^2
 - ▶ It does not depend on the effect size measure

- ▶ One-sided and point hypotheses were tested, but combined hypotheses can also be tested $\rightarrow H : \mu > 0 \ \& \ \rho > 0$

- ▶ Informative hypotheses can also be implemented

- ▶ Bayesian estimation and Bayes factors are included in the R package `BFpack` [8]

- ▶ Future research may focus on:
 - ▶ Extending the methodology to meta-regression models
 - ▶ Allowing for multiple outcomes per study and more complicated hierarchical structures
 - ▶ Taking uncertainty in the within-study variance into account
 - ▶ Studying to what extent the methodology gets distorted by publication bias

Thank you for your attention

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Paper:

Van Aert, R. C. M., & Mulder, J. (2021). Bayesian hypothesis testing and estimation under the marginalized random-effects meta-analysis model. *Psychonomic Bulletin & Review*. doi: [10.3758/s13423-021-01918-9](https://doi.org/10.3758/s13423-021-01918-9)

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[8]

Mulder J, Williams DR, Gu X, Tomarken A, Boeing-Messing F, Olsson-Collentine A, et al. BFpack: Flexible Bayes factor testing of scientific theories in R. *Journal of Statistical Software* 2021.

[9]

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- ▶ Bayes factors and Bayesian estimation are included in the R package BFpack [8]
- ▶ BF() function only needs a fitted modeling object → object returned by a random-effects meta-analysis using metafor [9]:

```
res2 <- rma(yi = yi, vi = vi) # RE meta-analysis
BF(res2)
```

```
## Call:
## BF.rma.uni(x = res2)
##
## Bayesian hypothesis test
## Type: exploratory
## Object: rma.uni
## Parameter: between-study heterogeneity & effect size
## Method: Bayes factor using uniform prior for icc & unit information prior for effect
##
## Posterior probabilities:
##      Pr(=0) Pr(<0) Pr(>0)
## I^2  0.696  0.064  0.240
## mu    0.537  0.210  0.254
```