

Heterogeneity in random-effects meta-analysis

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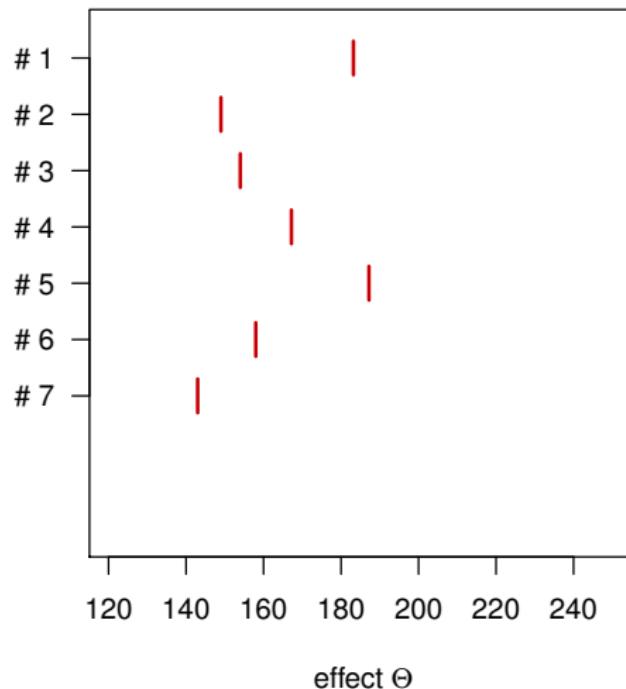


Overview

- Meta analysis
 - the random-effects model
 - frequentist approaches
 - the Bayesian approach
 - example
- Simulation study
 - bias
 - coverage
 - ...
- Conclusions

Meta analysis

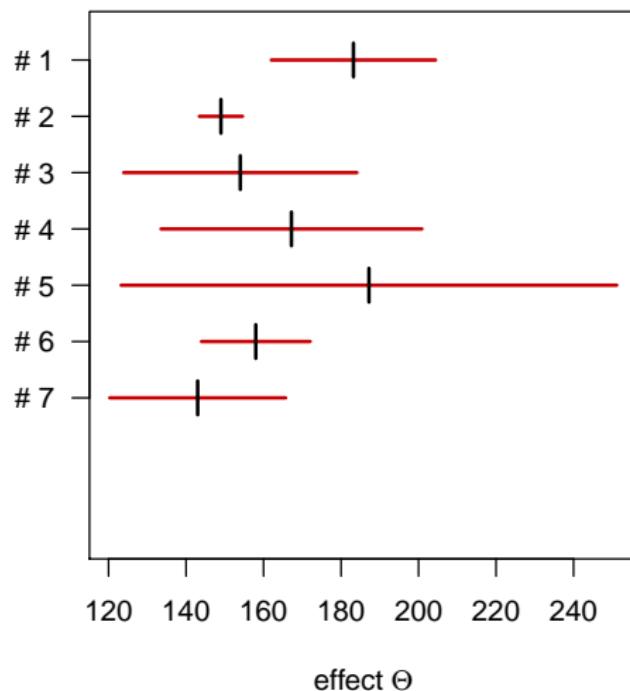
Context



- have:
 - estimates y_i
 - standard errors σ_i
- want:
 - combined estimate $\hat{\Theta}$

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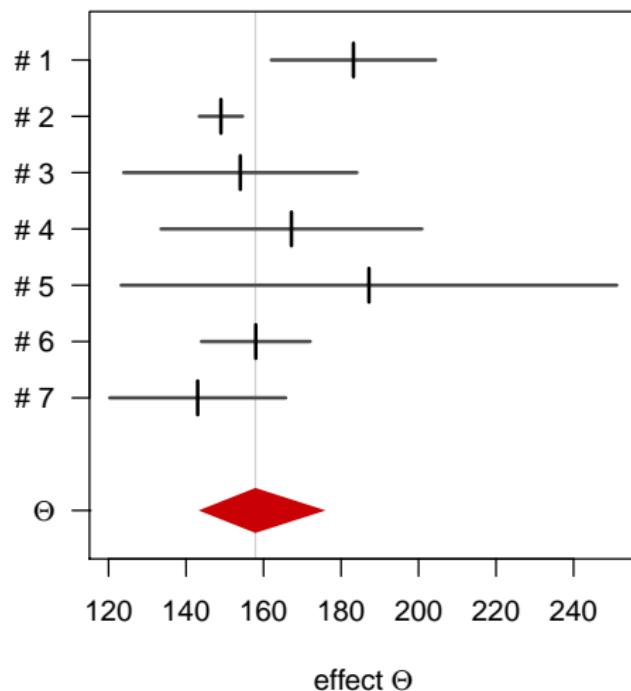
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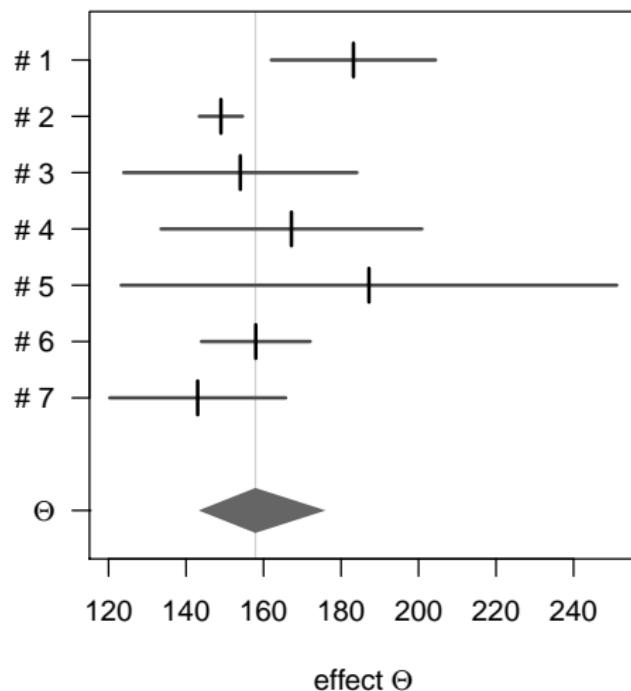
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The random effects model

- assume^{1,2}:

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

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- true parameter value Θ
- heterogeneity τ

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- ingredients:

Data:

- estimates y_i
- standard errors σ_i
- $\Theta \in \mathbb{R}$ of primary interest (“effect”)
- $\tau \in \mathbb{R}^+$ nuisance parameter (“between-trial heterogeneity”)

Parameters:

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Meta analysis

Common approach to inference

- the frequentist approach

- test for $\tau = 0$ vs. $\tau > 0$ (fixed vs. random effects)
- derive estimate $\hat{\tau}$
- derive estimate for Θ *conditional on $\hat{\tau}$ being actual heterogeneity* (plug-in estimate)

³J. Hartung, G. Knapp. On tests of the overall treatment effect in meta analysis with normally distributed responses. *Statistics in Medicine* 20(12):1771–1782, 2001.

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 - derive estimate $\hat{\tau}$
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- Comments:
 - significance tests have low power, $\tau = 0$ hypothesis questionable
 - how to estimate τ ? many approaches, questionable properties
 - conditioning on *fixed* τ value only sensible for great accuracy
 - uncertainty in τ often not accounted for
exception: “Knapp-Hartung” approach³
(leads to Student- t -approximation)
 - usually computationally simple

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Frequentist MA

technically: normal approximation

- usual frequentist procedure:
 - derive heterogeneity estimate $\hat{\tau}$
 - conditional on $\tau = \hat{\tau}$, derive estimate

$$\hat{\Theta} = \frac{1}{\sum w_i} \times \sum w_i y_i \quad \text{where} \quad w_i = \frac{1}{\sigma_i^2 + \hat{\tau}^2}$$

with standard error

$$SE(\hat{\Theta}) = \sqrt{\frac{1}{\sum w_i}}$$

and use Normal approximation for confidence interval

Frequentist MA

technically: Knapp-Hartung adjustment

- “Knapp-Hartung” approach⁴ to considering uncertainty in $\hat{\tau}$:
 - compute

$$q := \frac{1}{k-1} \sum \frac{(y_i - \hat{\Theta})^2}{\sigma_i^2 + \hat{\tau}^2}$$

- use Student-*t* approximation with scale

$$\max\{\sqrt{q}, 1\} \times \text{SE}(\hat{\Theta})$$

and $(k-1)$ degrees of freedom to construct confidence interval
(instead of Normal approximation)

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Meta analysis

The Bayesian approach

- Bayesian approach⁵
 - set up model likelihood
 - specify prior information about unknowns
 - posterior results as \propto prior \times likelihood

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Meta analysis

The Bayesian approach

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 - set up model likelihood
 - specify prior information about unknowns
 - posterior results as \propto prior \times likelihood
- Comments:
 - consideration of prior information
 - consideration of uncertainty
 - straightforward interpretation
 - computationally more expensive, usually done via stochastic integration (MCMC, BUGS)⁶
 - special case of simple random-effects MA may be solved semi-analytically (using `bmeta` R package)

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Bayesian MA

technically: prior, posterior

- have:
 - likelihood $p(\vec{y}, \vec{\sigma} | \Theta, \tau)$
 - prior density $p(\Theta, \tau) = p(\Theta) \times p(\tau)$
- posterior $p(\Theta, \tau | \vec{y}, \vec{\sigma}) \propto p(\vec{y}, \vec{\sigma} | \Theta, \tau) \times p(\Theta, \tau)$
- integrate out marginal posteriors
 - effect $p(\Theta | \vec{y}, \vec{\sigma}) = \int p(\Theta, \tau | \vec{y}, \vec{\sigma}) d\tau$
 - heterogeneity $p(\tau | \vec{y}, \vec{\sigma}) = \int p(\Theta, \tau | \vec{y}, \vec{\sigma}) d\Theta$
- inference: posterior expectations, medians, quantiles, . . .

Meta analysis

Frequentist and Bayesian approaches

- many heterogeneity estimators available
- different prior specifications possible (should depend on context)

⁷ K. Sidik, J.N. Jonkman. A comparison of heterogeneity variance estimators in combining results of studies. *Statistics in Medicine* 26(9):1964–1981, 2007.

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 - DerSimonian-Laird estimator (DL)
 - restricted ML estimator (REML)⁷
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 - half-Normal ($\sigma = 0.5$)
 - half-Normal ($\sigma = 1.0$)
 - Uniform (0.0, 4.0)

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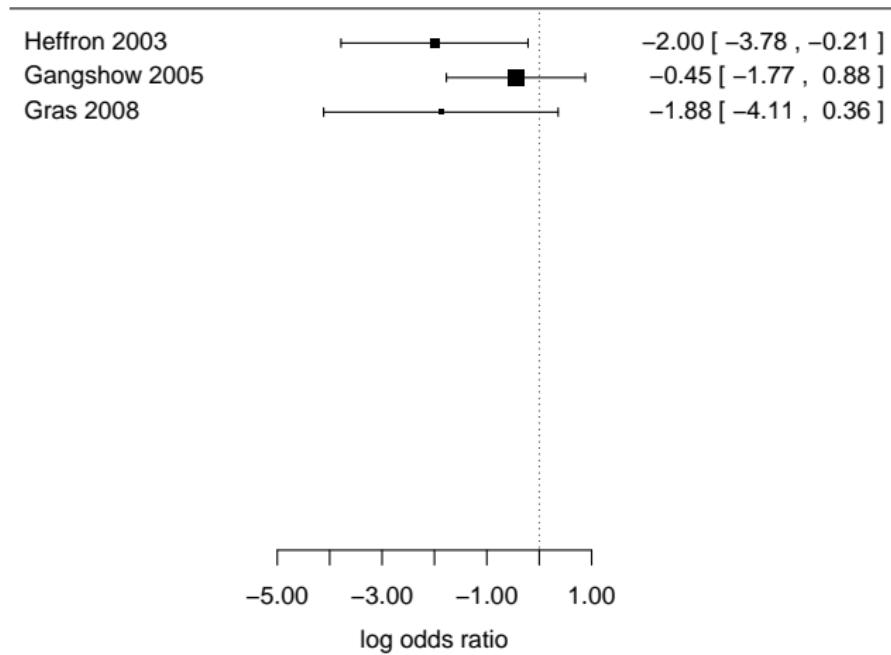
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Example

Crins et al. (2014) data¹⁰

Liver transplant example: steroid-resistant rejection (SRR)



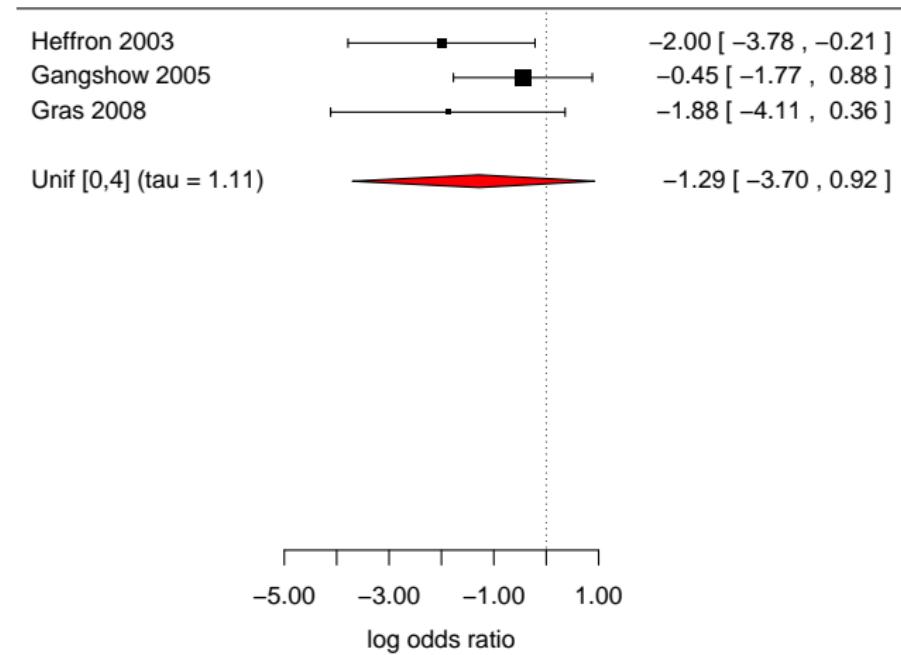
data: 3 estimates
(log ORs)
and standard errors

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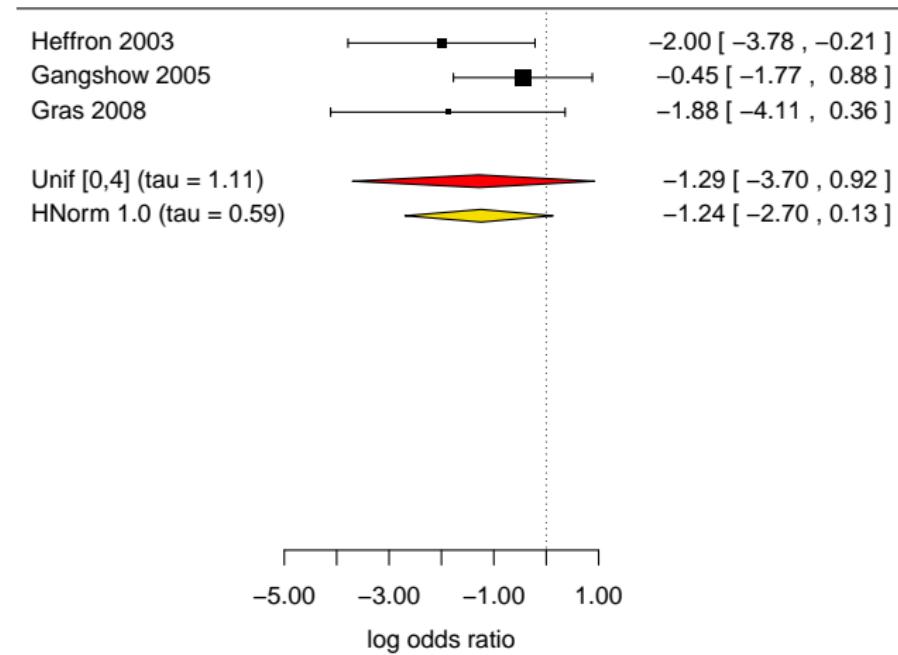
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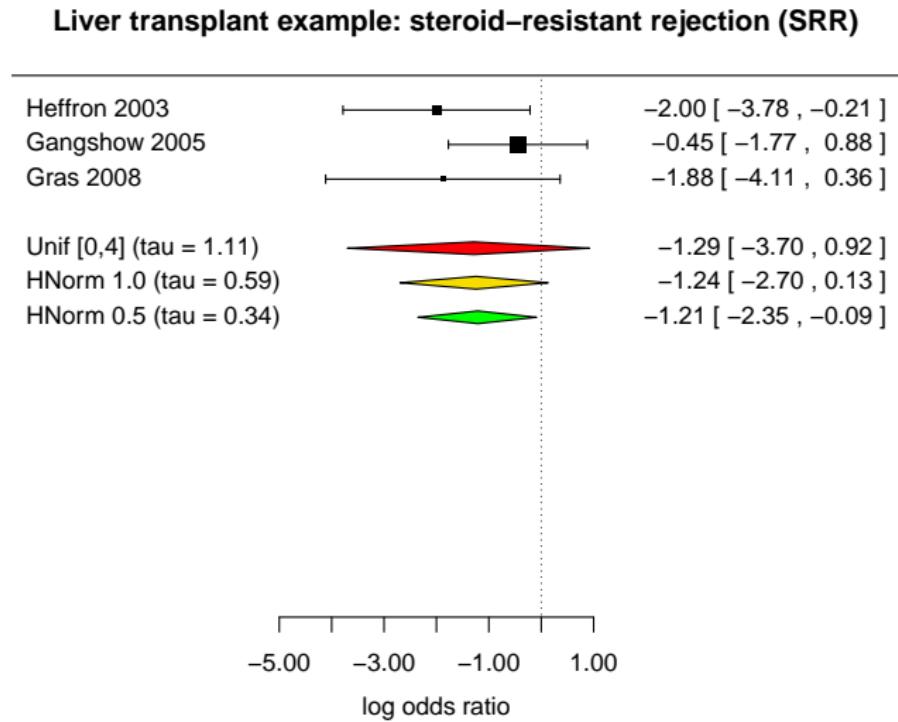
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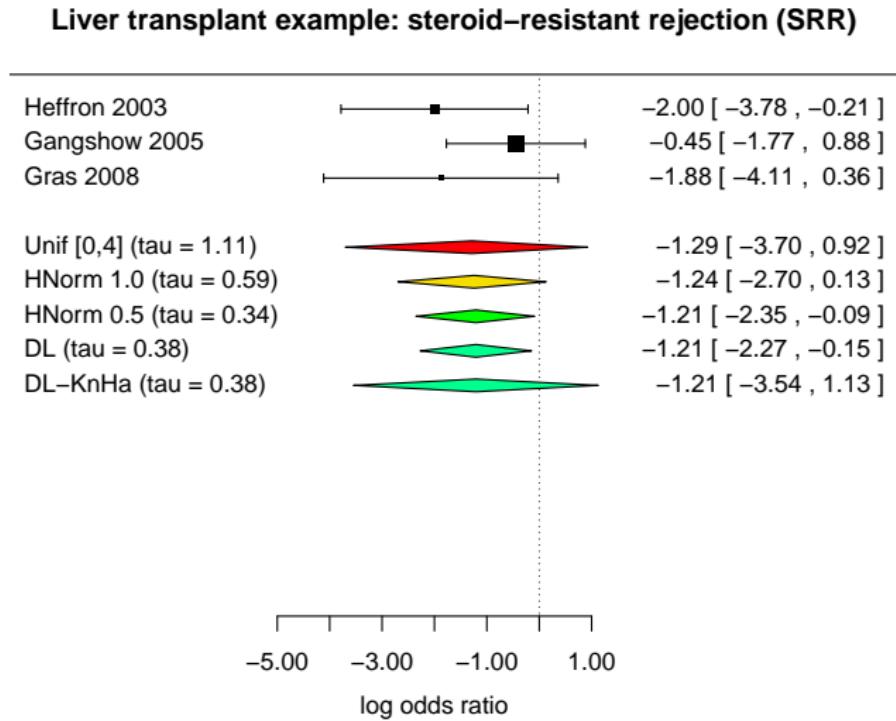


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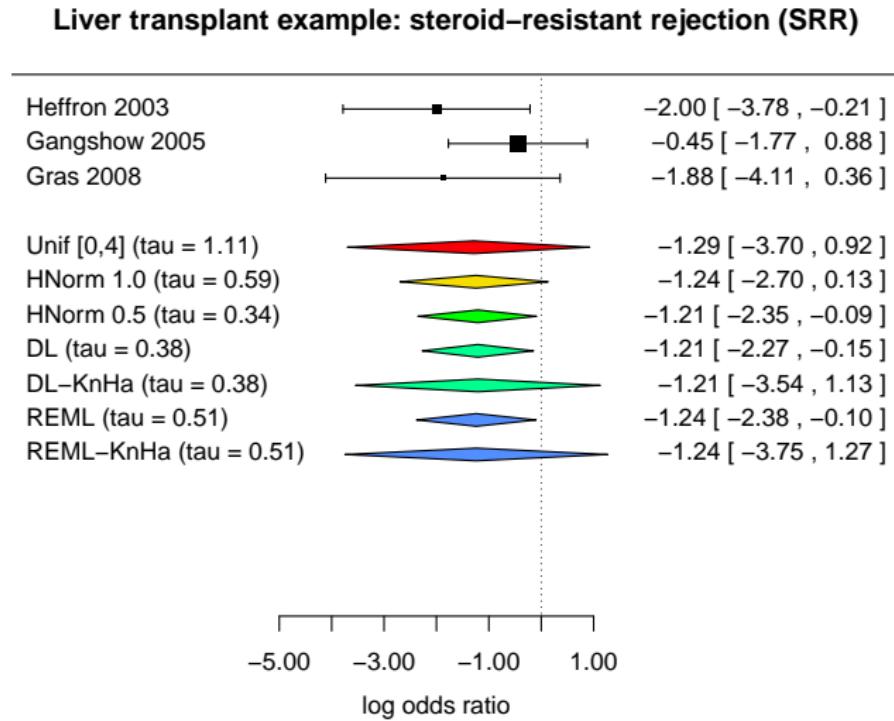


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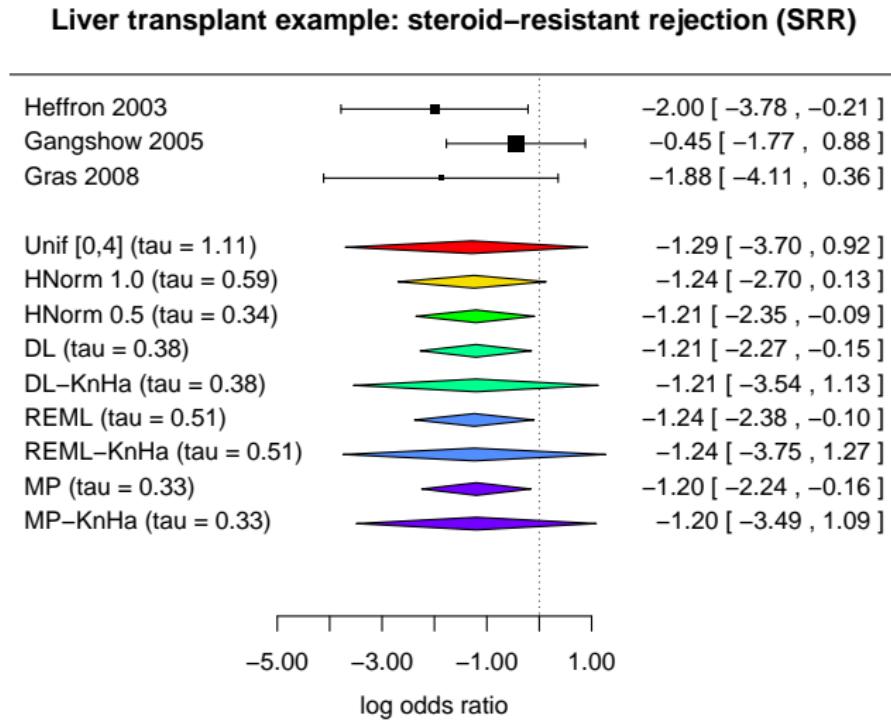


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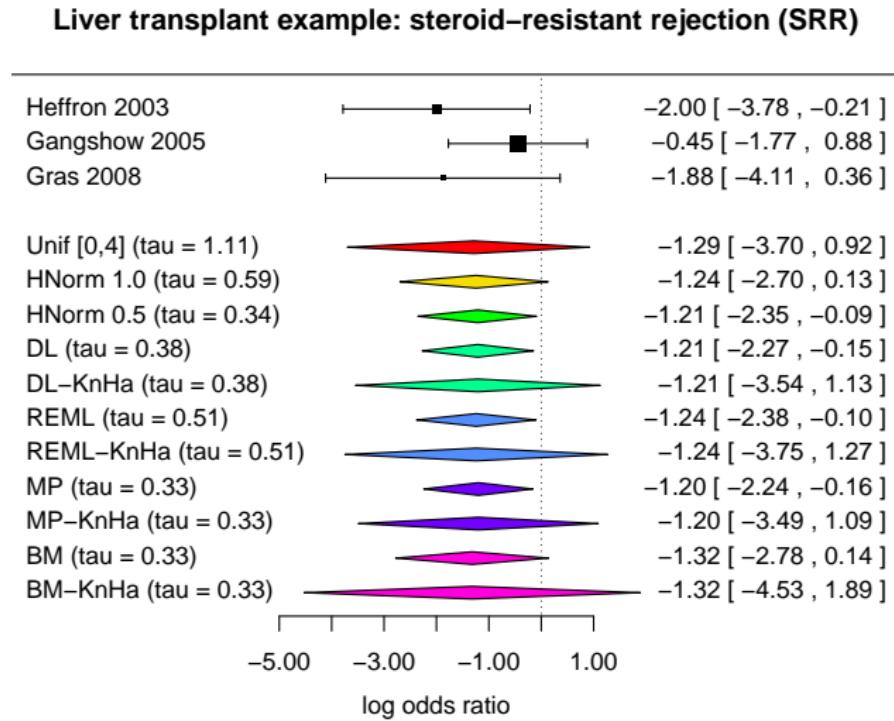


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Crins et al. (2014) data

- different analyses yield different answers
- Bayesian and frequentist analyses answer different questions
- $k = 2$ to 3 studies is a common scenario
(*majority* of meta analyses in Cochrane Database¹¹)

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Crins et al. (2014) data

- different analyses yield different answers
- Bayesian and frequentist analyses answer different questions
- $k = 2$ to 3 studies is a common scenario
(*majority* of meta analyses in Cochrane Database¹¹)
- how does performance compare **in general**, especially for few studies?

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Simulation study

Setup

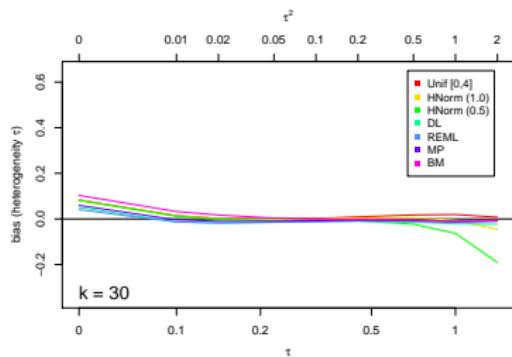
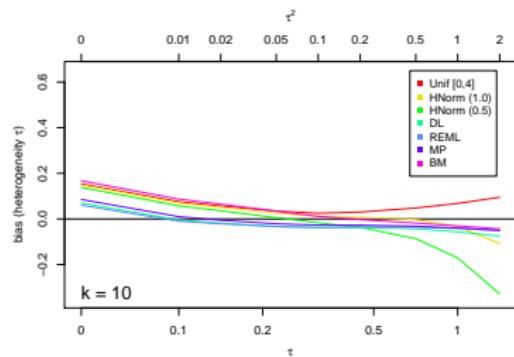
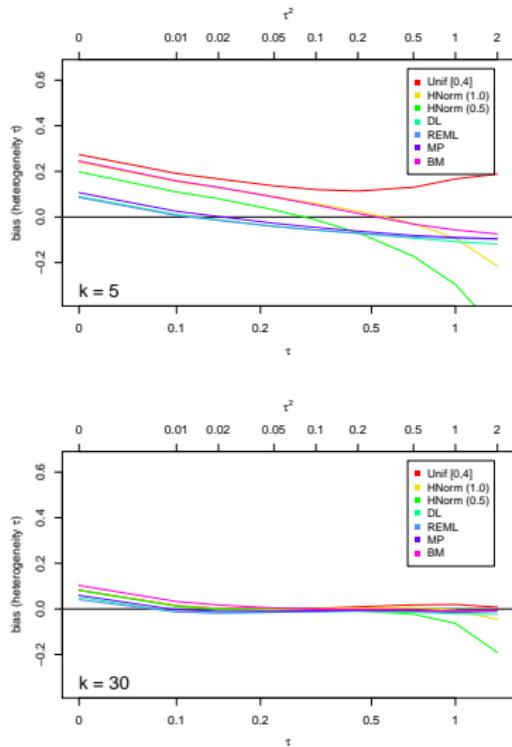
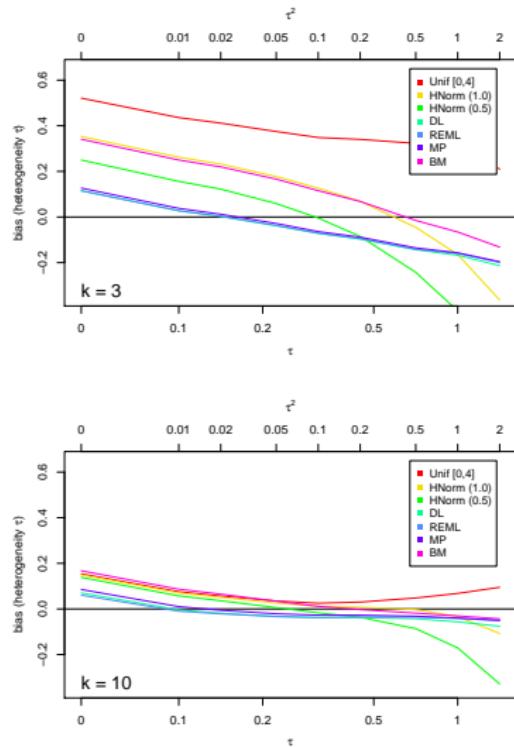
- number of studies: $k \in \{3, 5, 10, 30\}$
- heterogeneity: $\tau^2 \in \{0.00, 0.01, 0.02, 0.05, 0.1, 0.2, 0.5, 1.0, 2.0\}$
 $(I^2 \in \{0.00, 0.06, 0.11, 0.23, 0.37, 0.54, 0.75, 0.85, 0.92\})$
- standard errors σ_i : truncated χ^2 -distribution¹²
- 10'000 repetitions for each combination (k, τ^2)
- compute Bayesian MAs (3 different priors)
- compute frequentist MAs (different τ estimators, Normal and Knapp-Hartung approximation)

¹²S.E. Brockwell, I.R. Gordon. A comparison of statistical methods for meta-analysis. *Statistics in Medicine* 20(6):825–840, 2001.

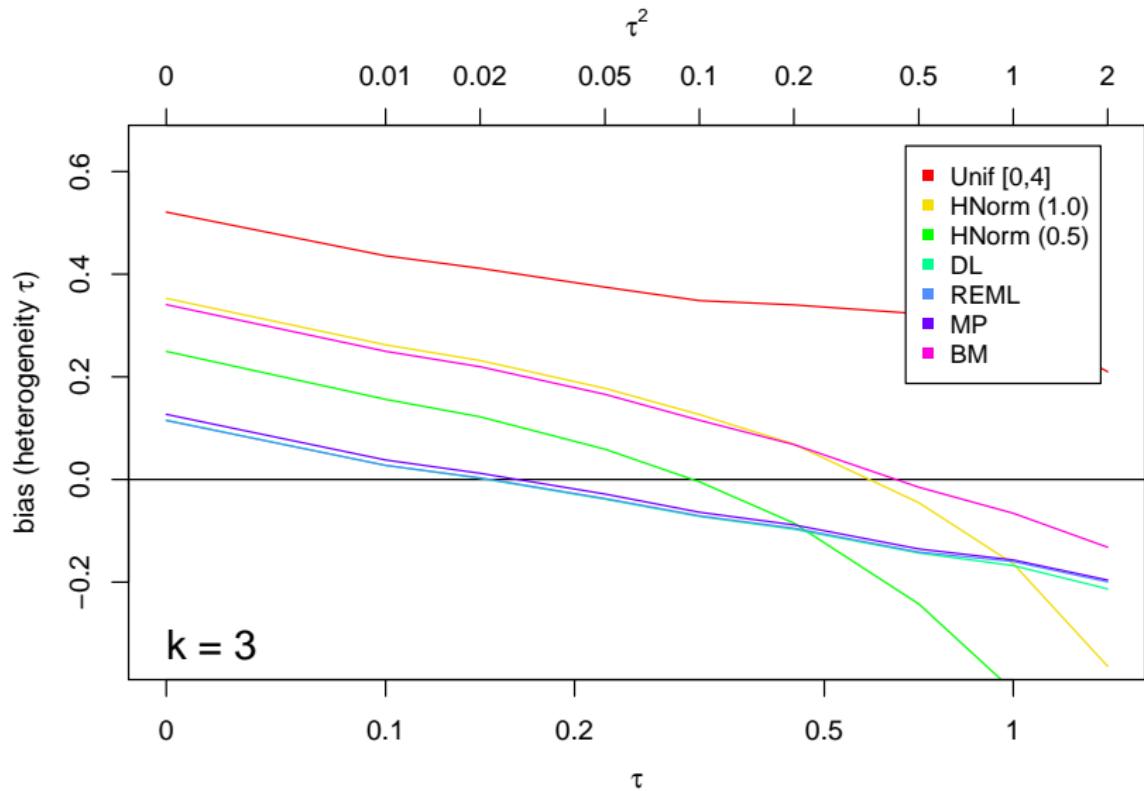
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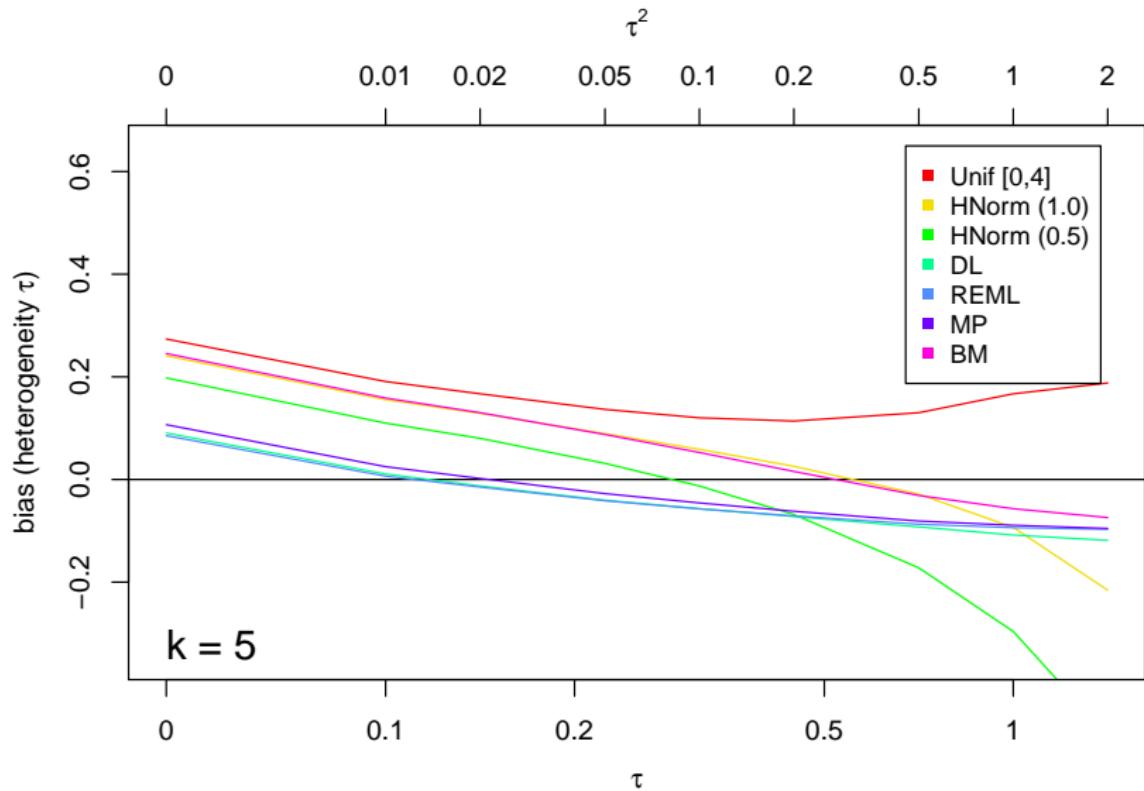
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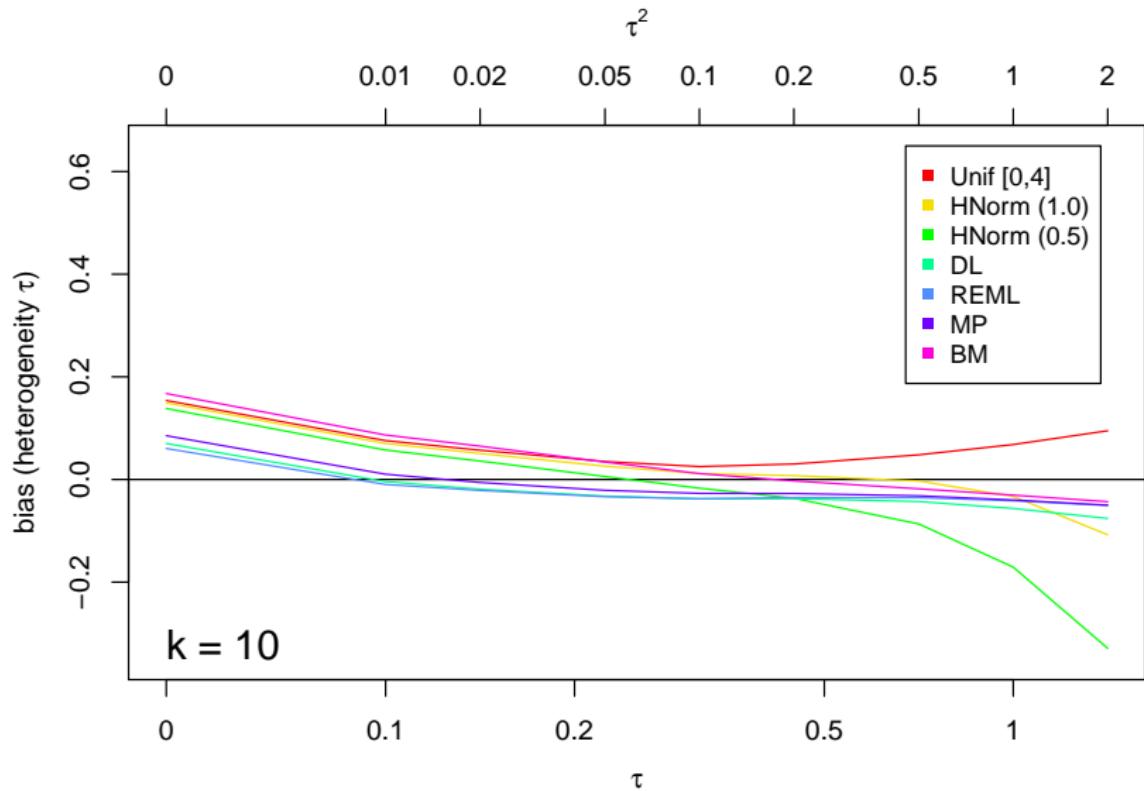
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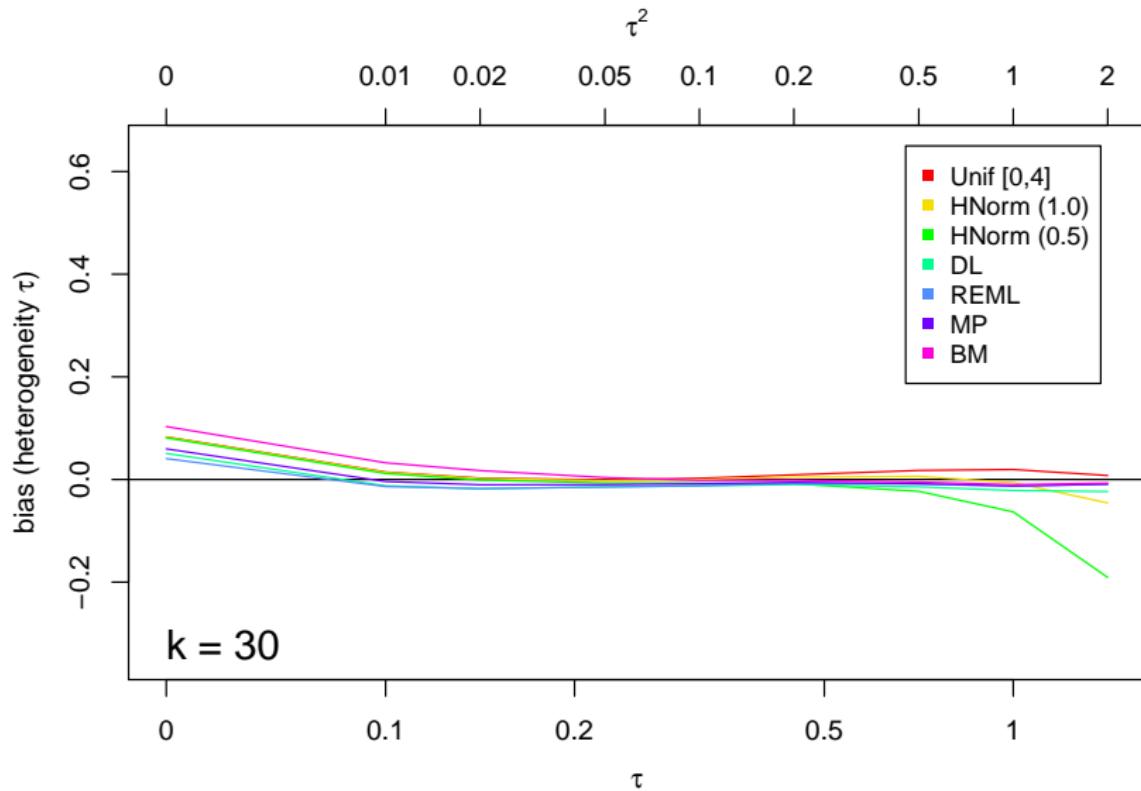
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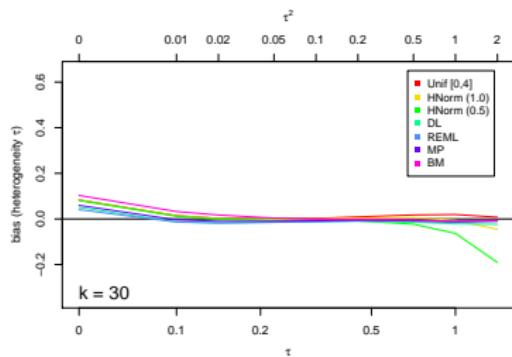
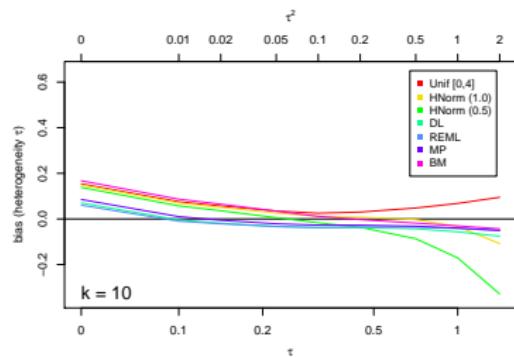
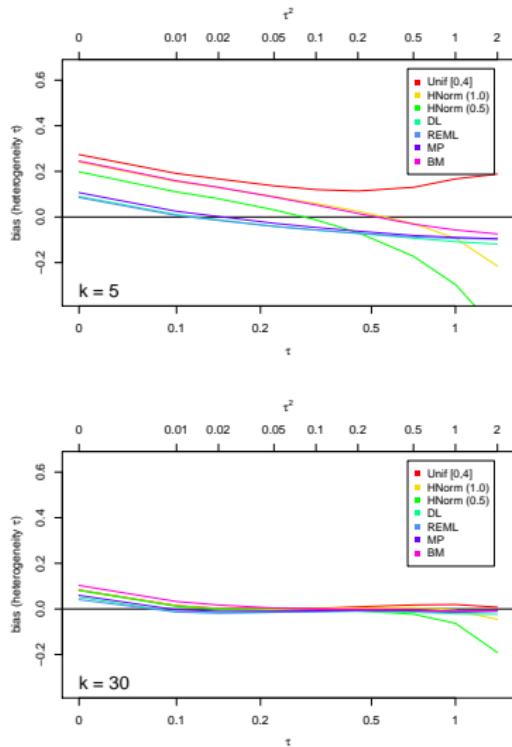
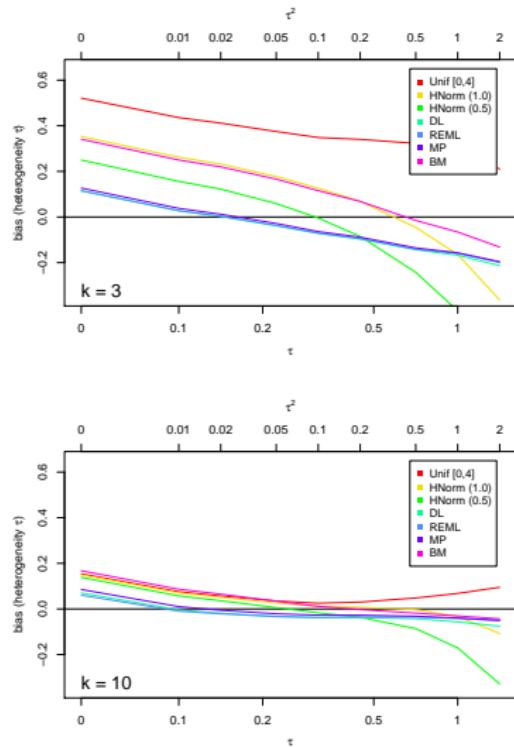


heterogeneity estimation: bias



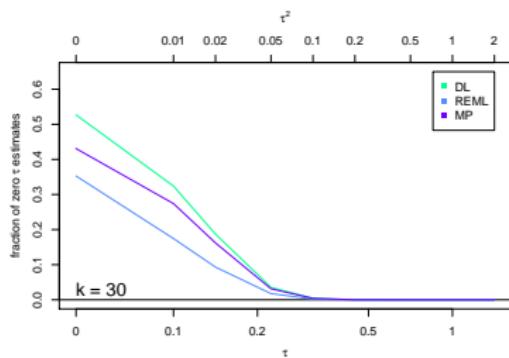
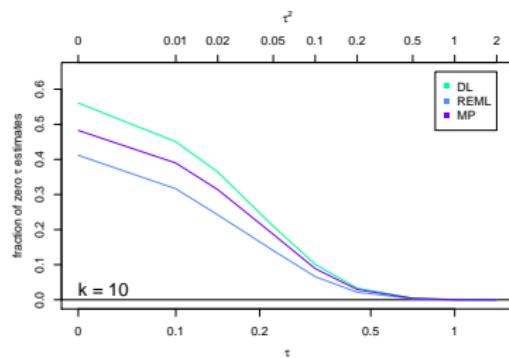
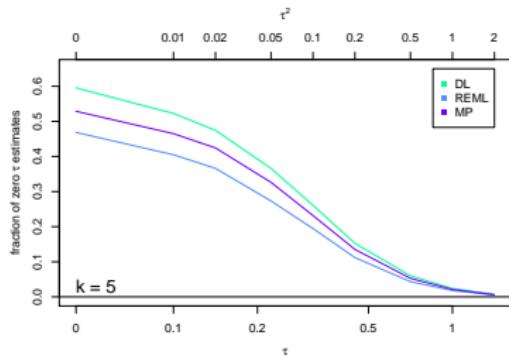
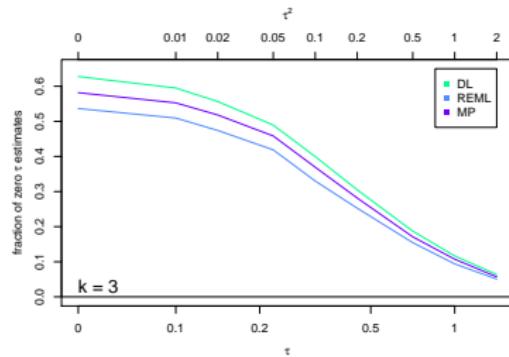
Simulation study

heterogeneity estimation: bias

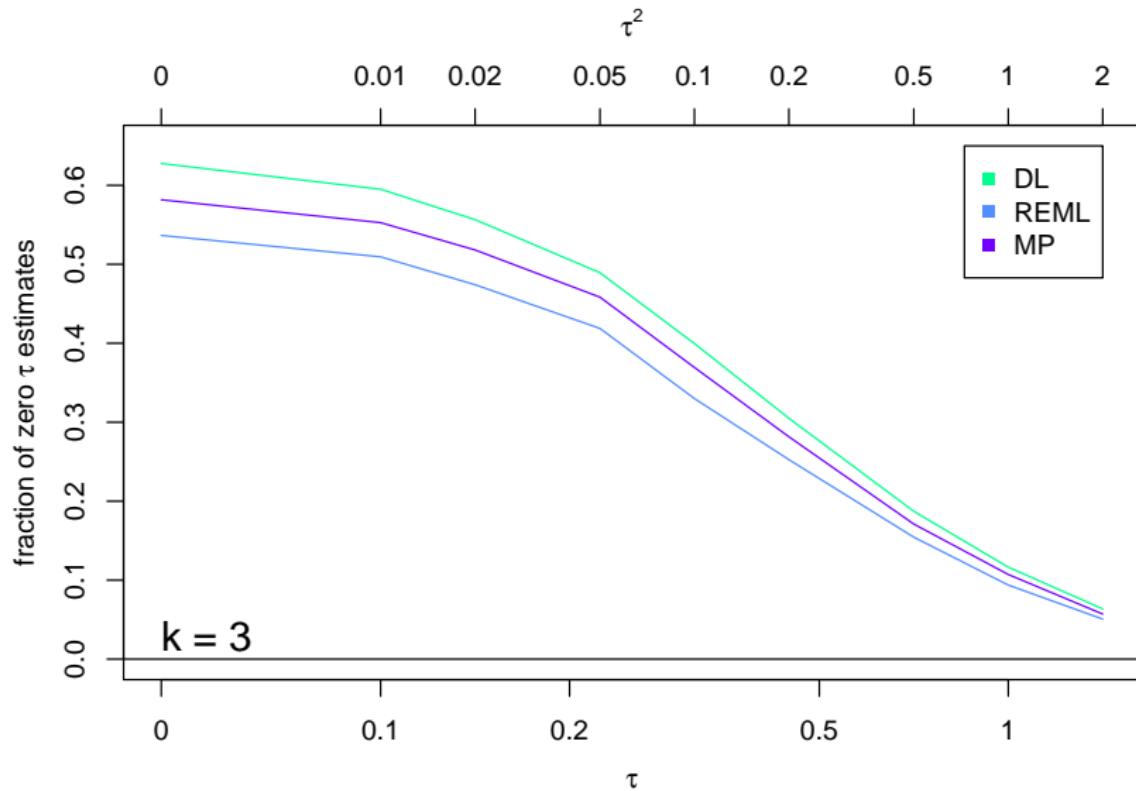


Simulation study

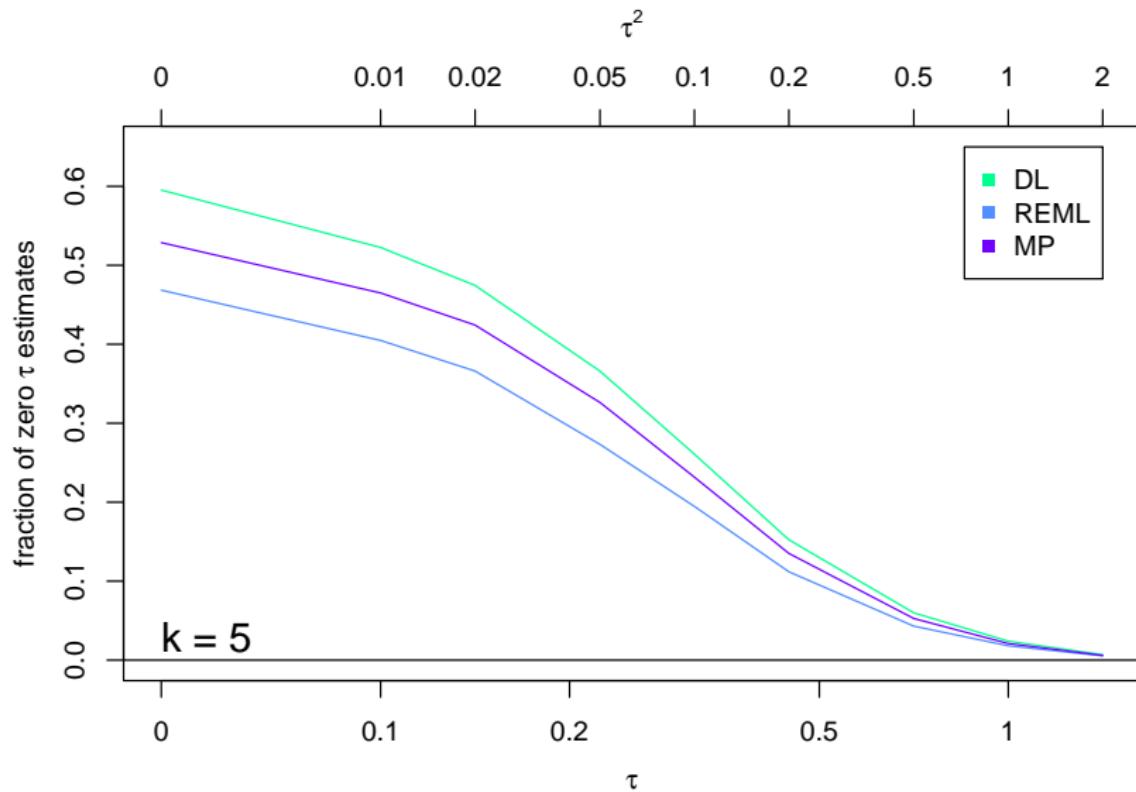
heterogeneity estimation: zero estimates



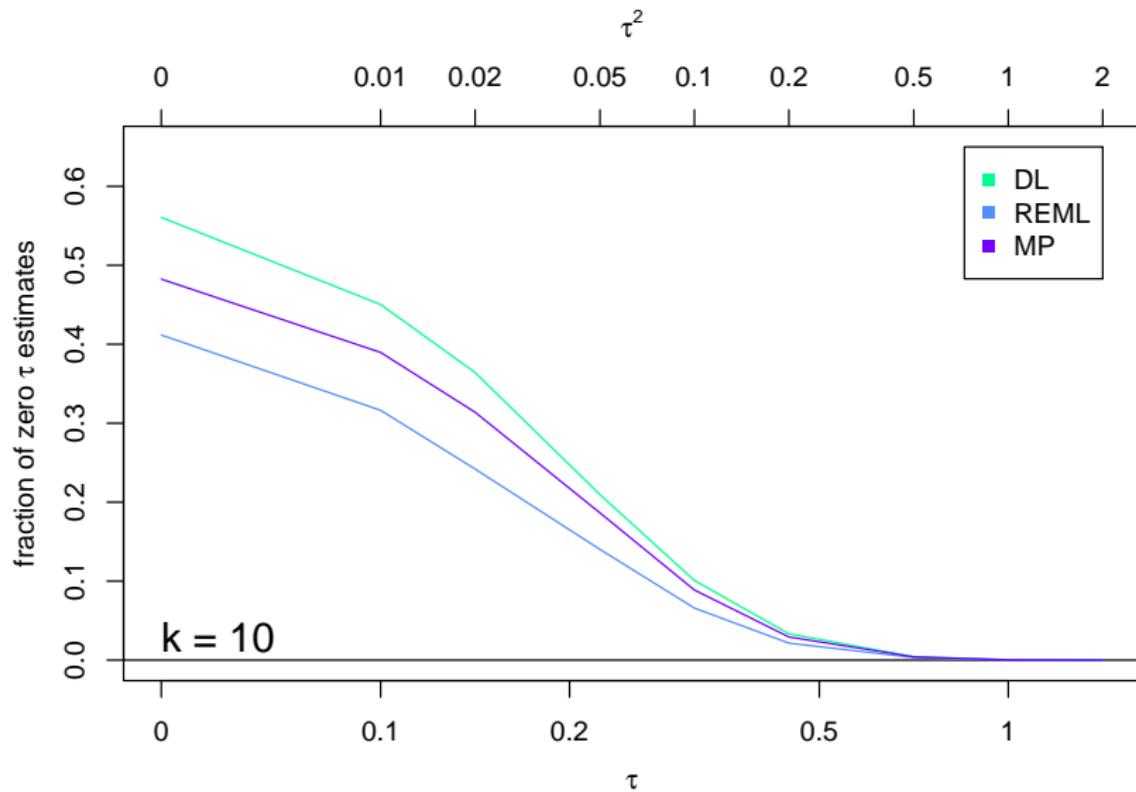
heterogeneity estimation: zero estimates



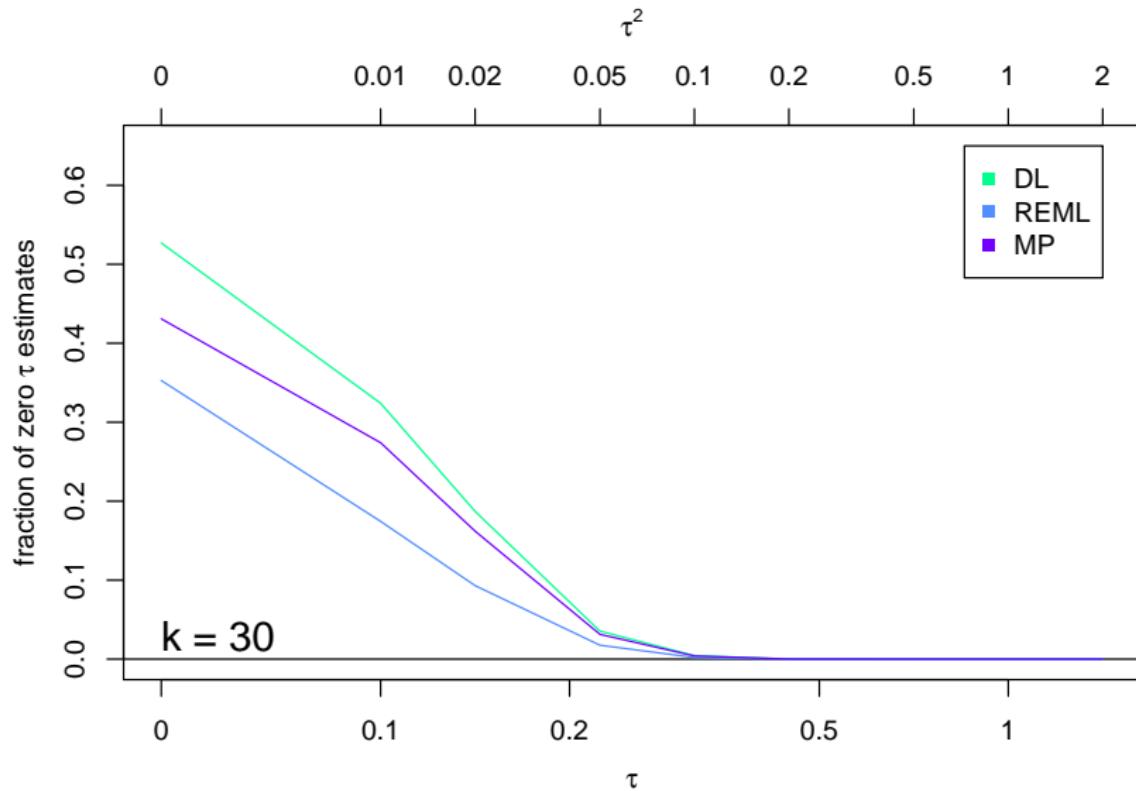
heterogeneity estimation: zero estimates



heterogeneity estimation: zero estimates

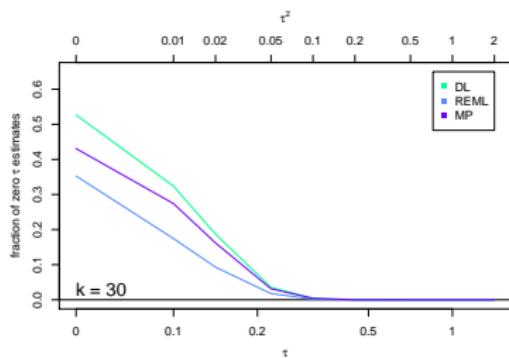
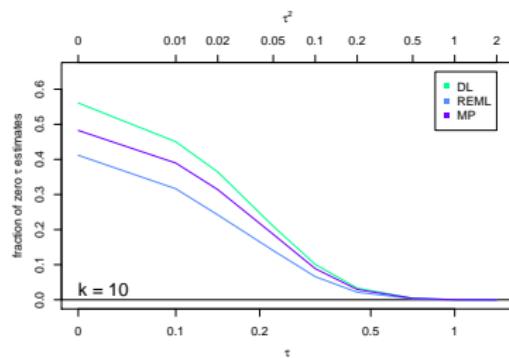
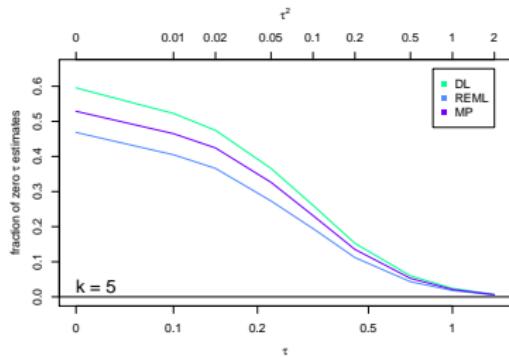
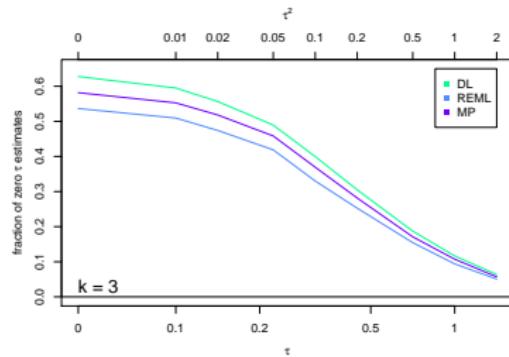


heterogeneity estimation: zero estimates



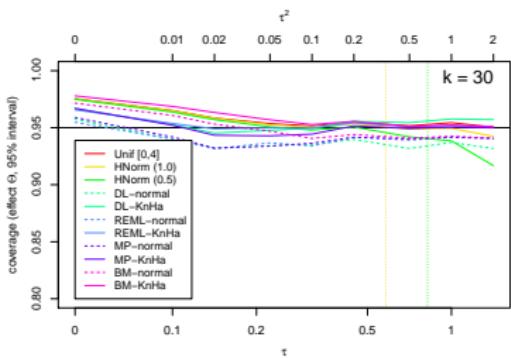
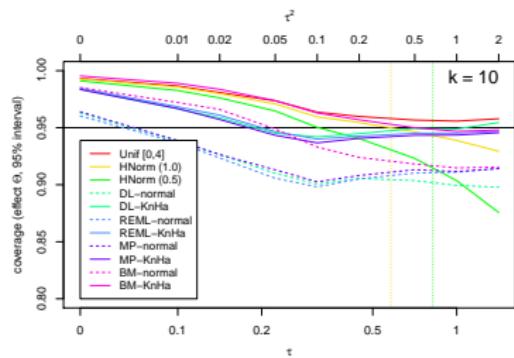
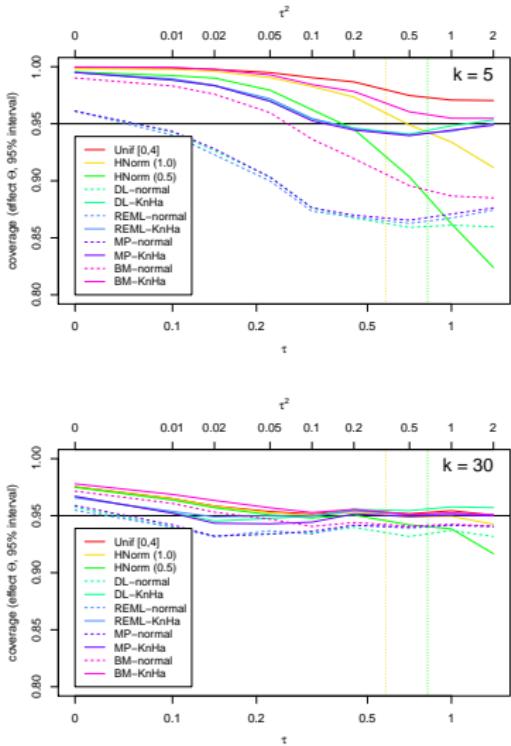
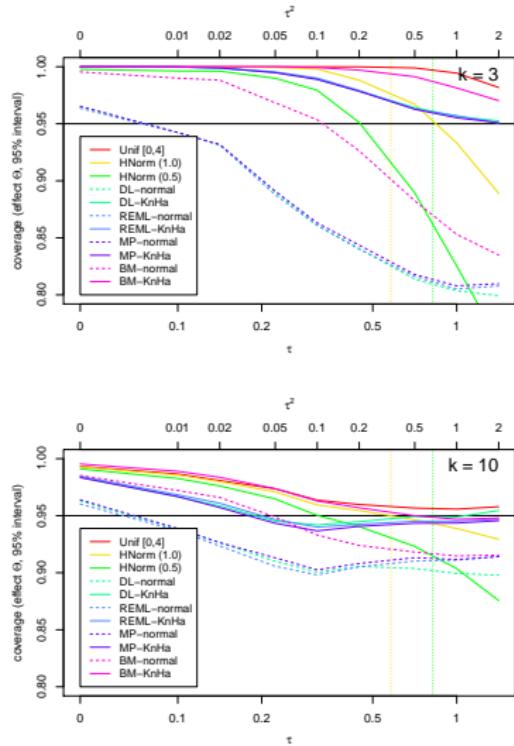
Simulation study

heterogeneity estimation: zero estimates

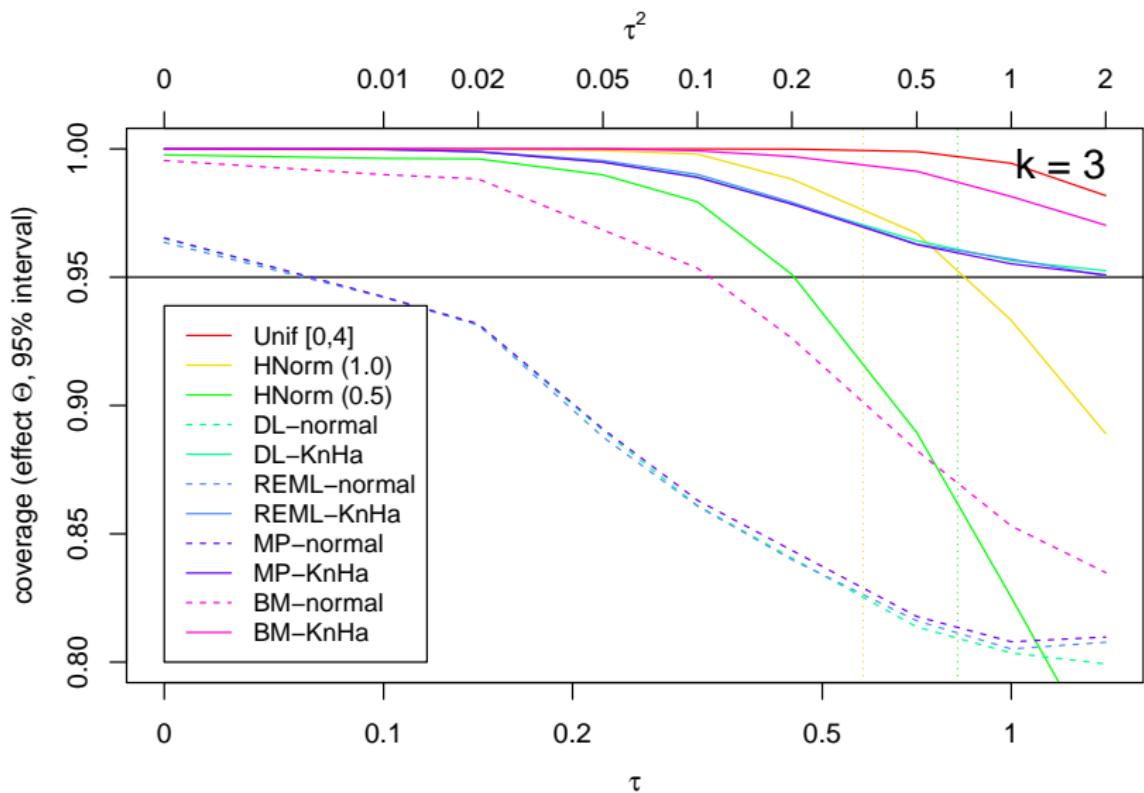


Simulation study

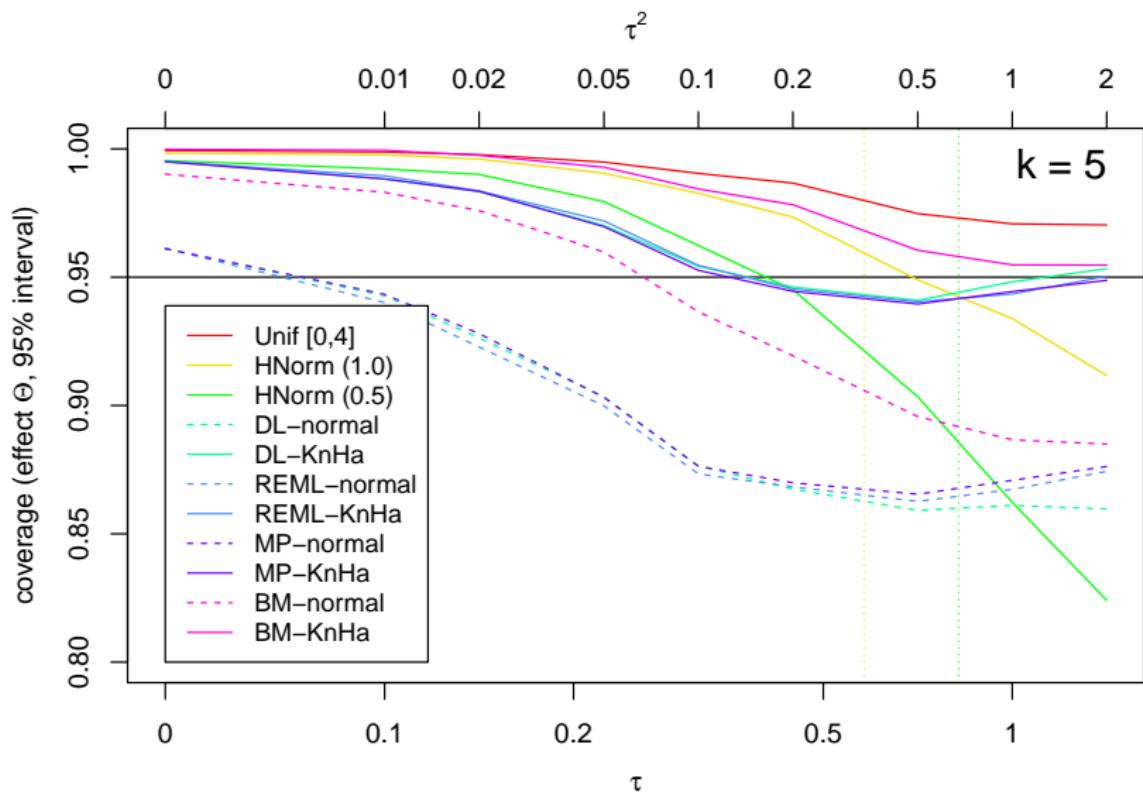
effect estimation: 95% CI coverage



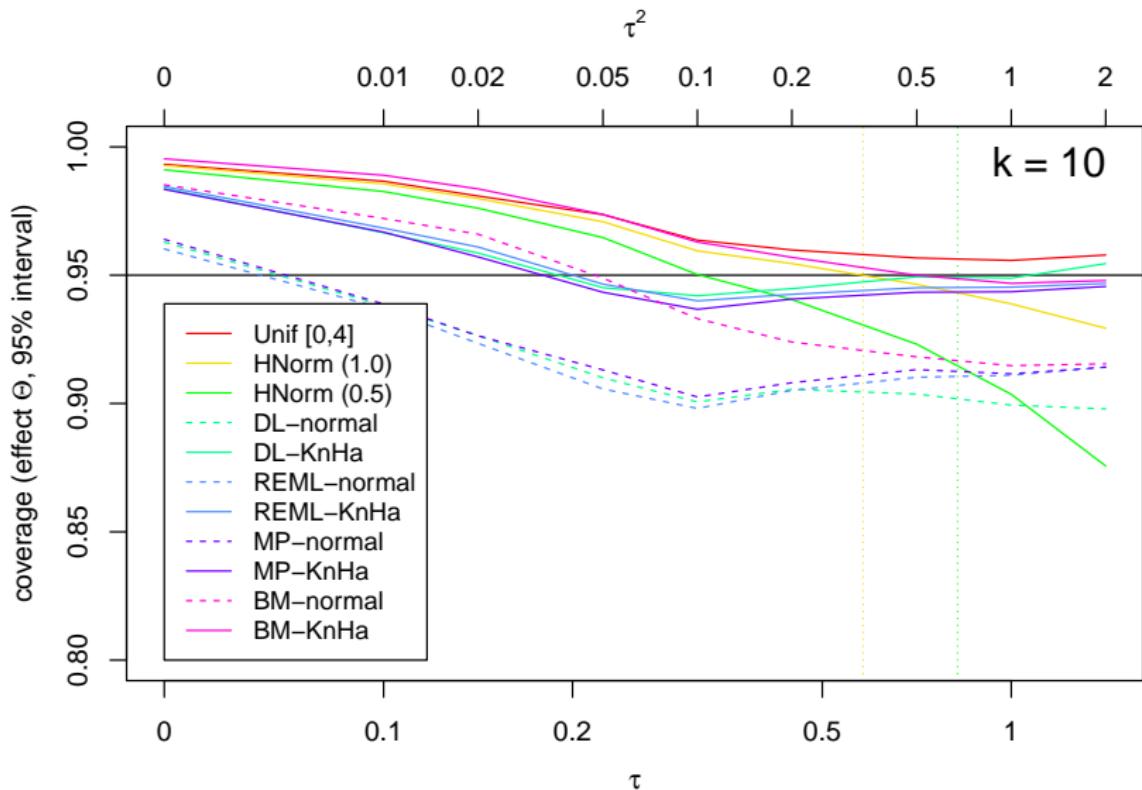
effect estimation: 95% CI coverage



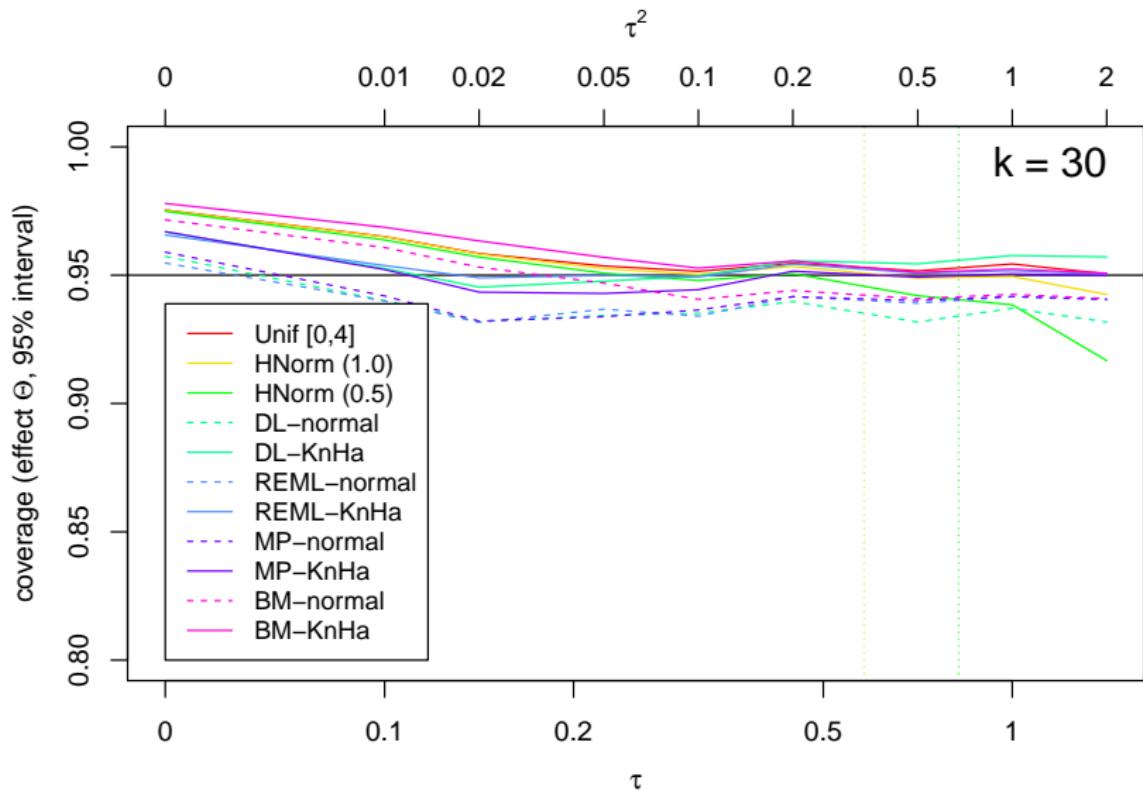
effect estimation: 95% CI coverage



effect estimation: 95% CI coverage

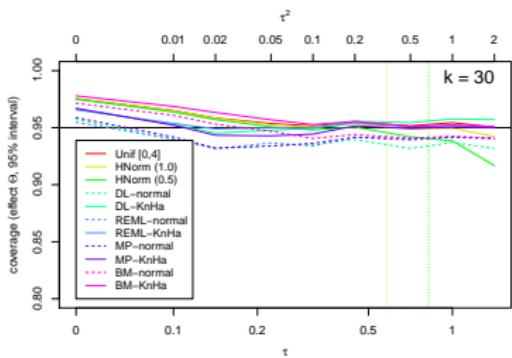
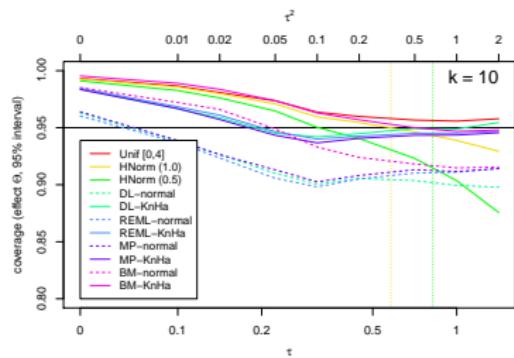
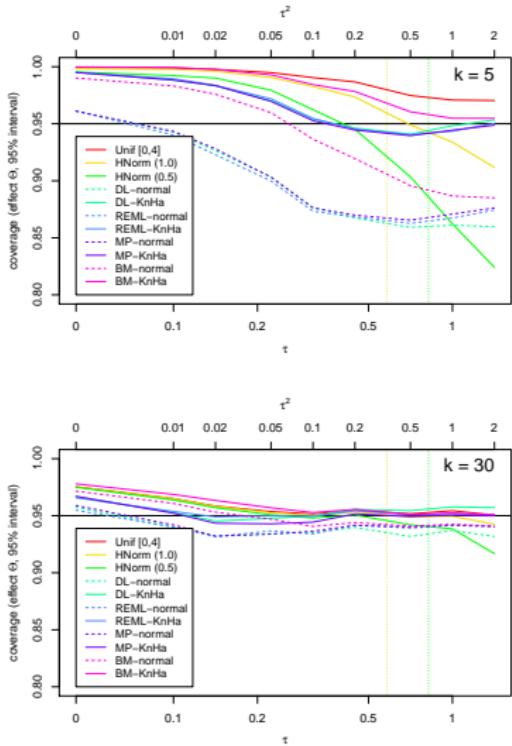
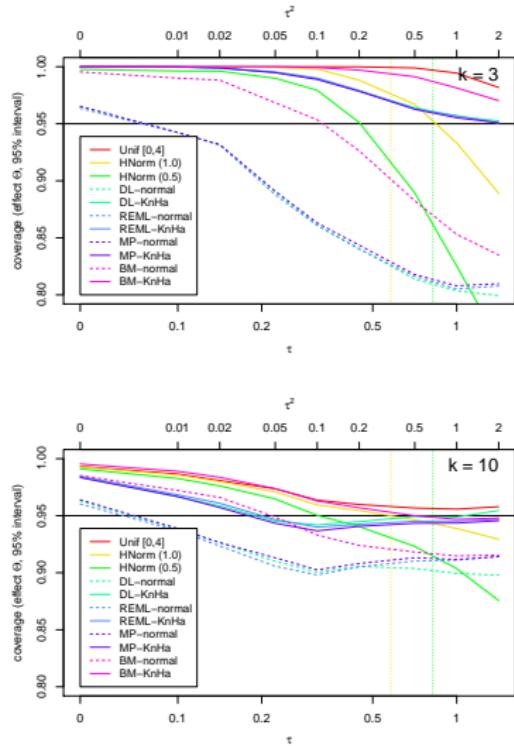


effect estimation: 95% CI coverage



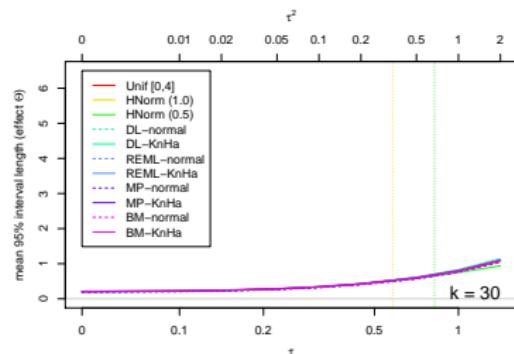
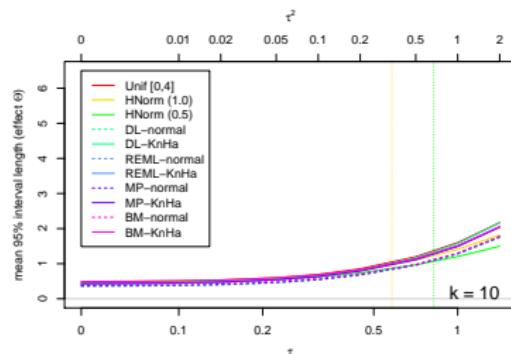
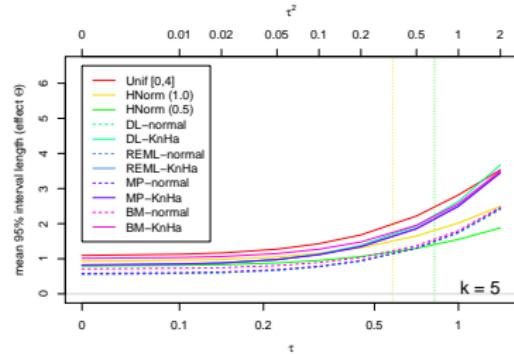
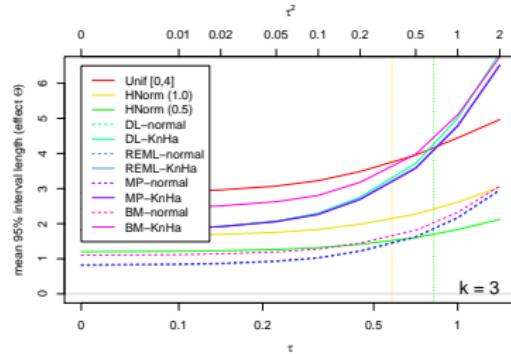
Simulation study

effect estimation: 95% CI coverage

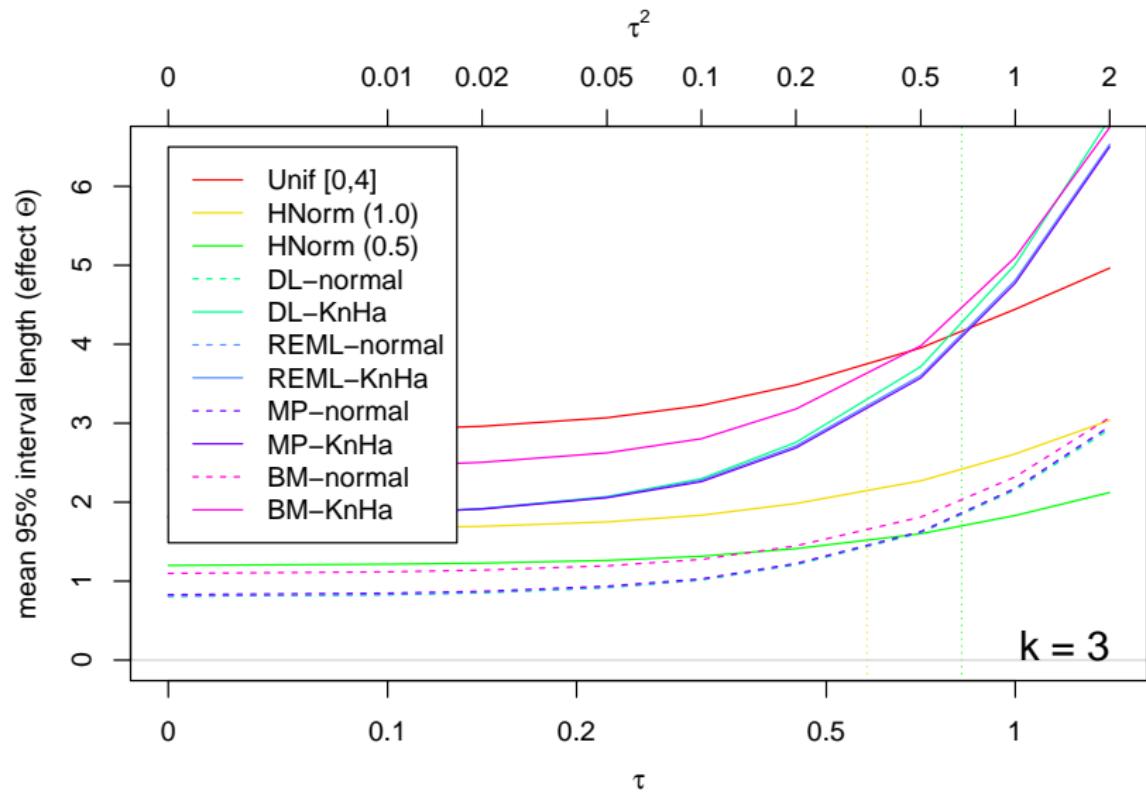


Simulation study

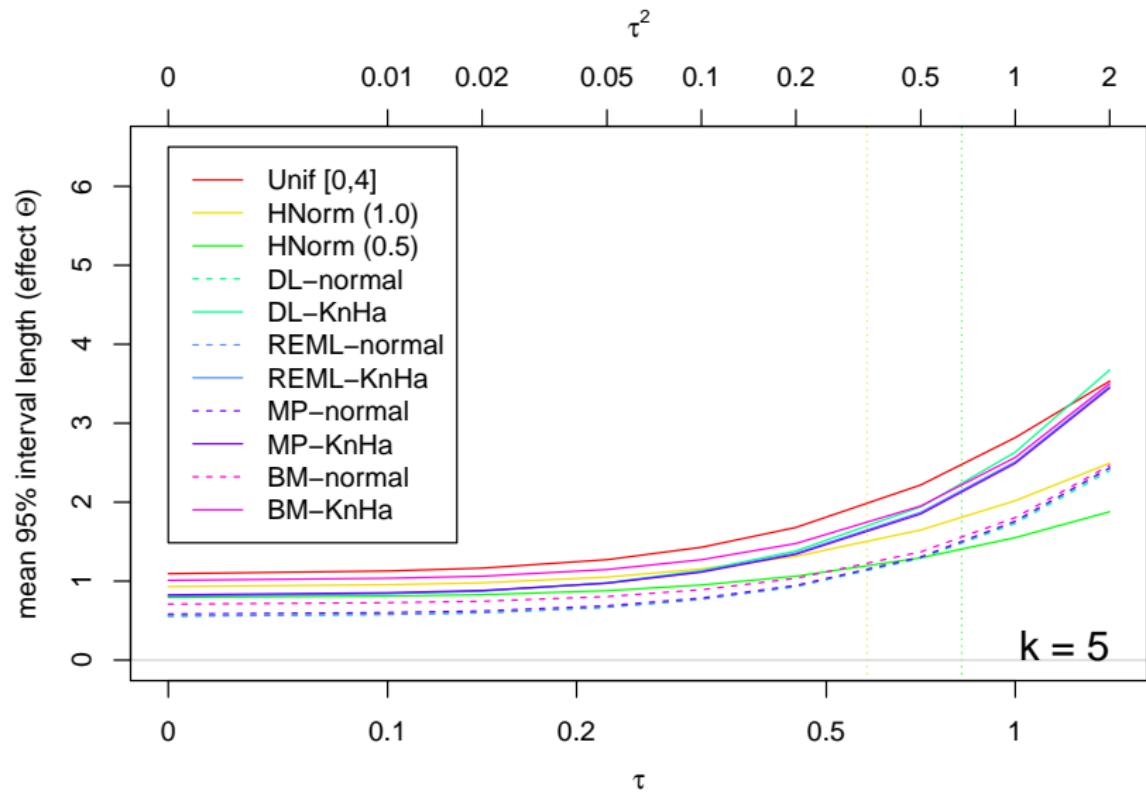
effect estimation: 95% CI length



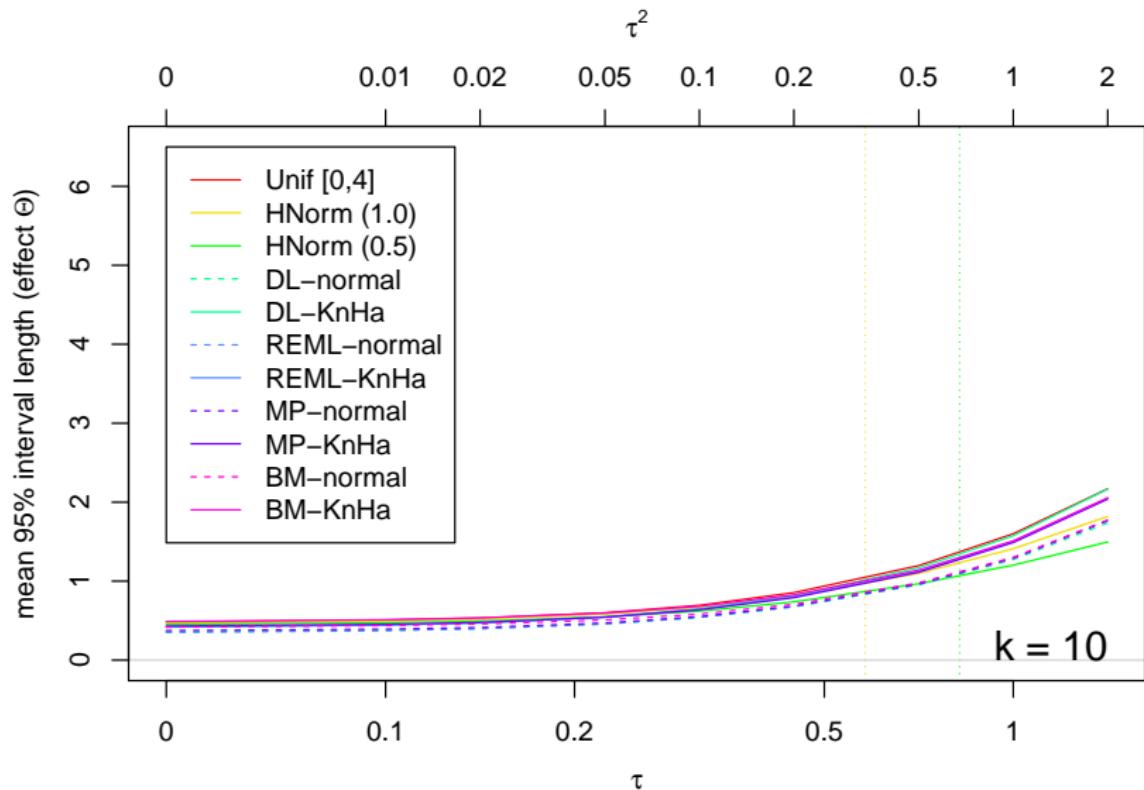
effect estimation: 95% CI length



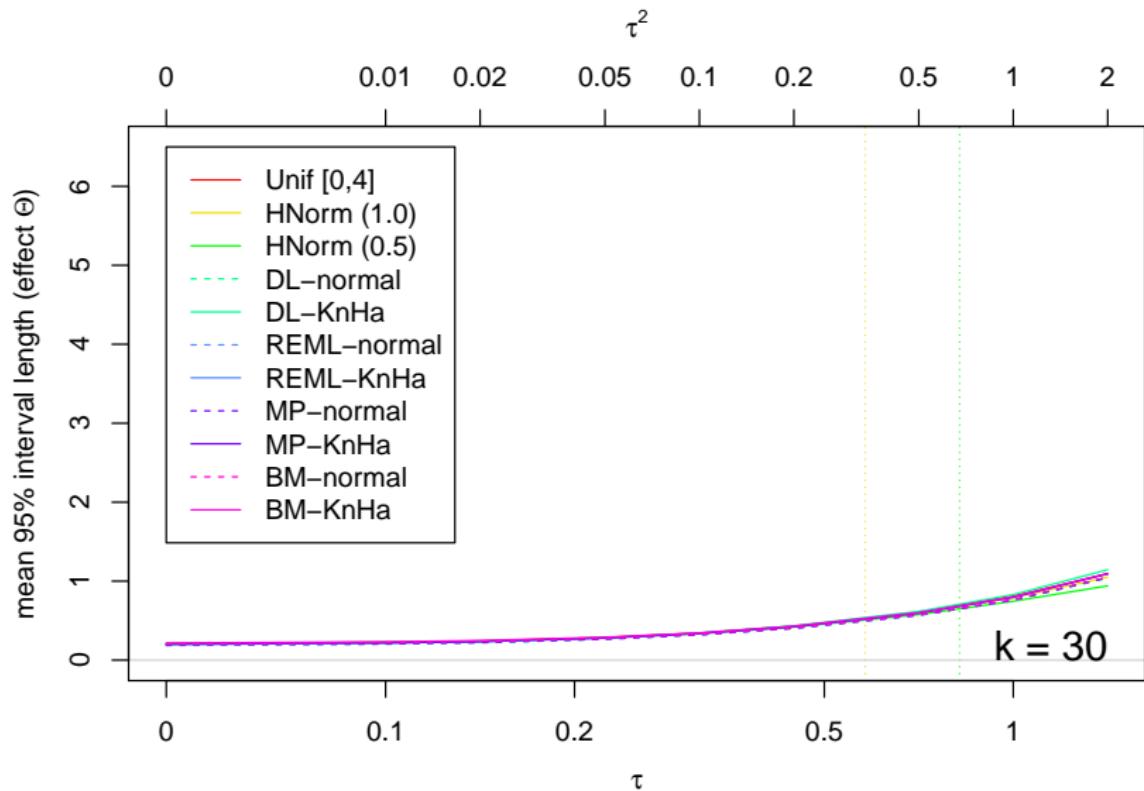
effect estimation: 95% CI length



effect estimation: 95% CI length

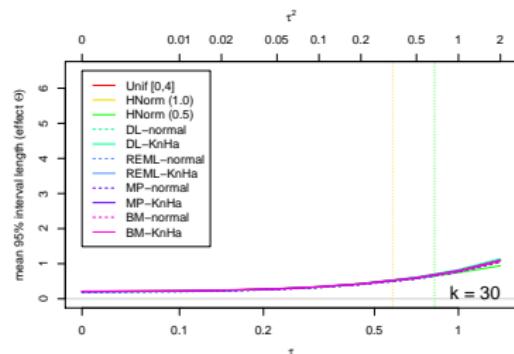
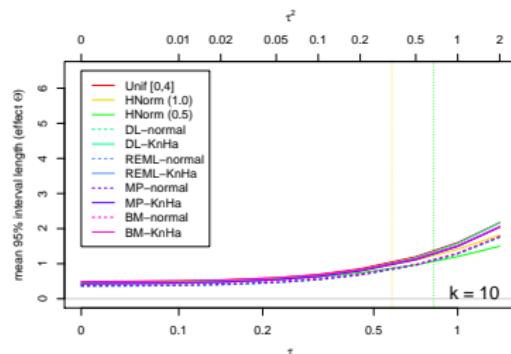
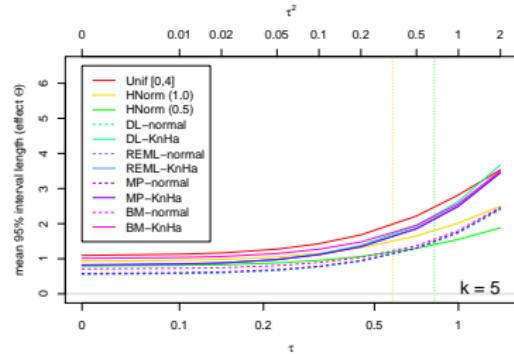
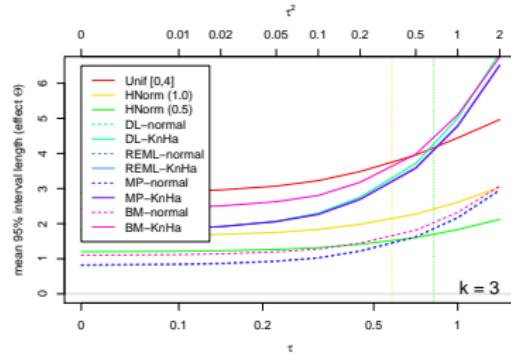


effect estimation: 95% CI length



Simulation study

effect estimation: 95% CI length



Conclusions

- small differences between different frequentist methods
- differences most pronounced in common case of few studies
- consideration of estimation uncertainty:
undercoverage with normal approximation,
application of Knapp-Hartung adjustment crucial for nominal level
- surprisingly many zero τ estimates
- Bayesian methods behave as expected:
conservative / anticonservative for “small” / “large” τ
 (“Mean coverage” (calibration) accurate *by construction*)
- Bayesian methods allow to utilize external information
(effect and heterogeneity, e.g.¹³)
- `bmeta` R package to appear on CRAN soon
- **ACKNOWLEDGMENT:**
funded by the EU through InSPiRe (FP HEALTH 2013 - 602144)

¹³R.M. Turner et al. Predictive distributions for between-study heterogeneity and simple methods for their application in Bayesian meta-analysis. *Statistics in Medicine* 34(6):984–998, 2015.

+++ additional slides +++

Implementation

bmeta R package under development

```
> cochrano1 <- bmeta(Cochran1954[, "mean"], sqrt(Cochran1954[, "se2"]))
> cochrano2 <- bmeta(Cochran1954[, "mean"], sqrt(Cochran1954[, "se2"]),
+                      mu.prior.mean=150, mu.prior.sd=100,
+                      tau.prior=function(x){return(dexp(x, rate=0.05))})
>
> cochrano1$summary
      tau        mu    mu.pred
mode 10.303255 156.504954 154.16345
median 12.888735 157.896520 157.33321
mean 14.844457 158.547999 158.54800
sd   9.950631   8.358115 19.70028
95% lower 0.000000 143.180913 119.77459
95% upper 32.665117 176.106158 200.12309
>
> # compute posterior quantiles:
> cochrano1$qposterior(mu.p=c(0.005, 0.995))
[1] 135.0429 187.3122
>
> # plot posterior density:
> x <- seq(from=130, to=190, length=100)
> plot(x, cochrano2$dposterior(mu=x), type="l")
> lines(x, cochrano1$dposterior(mu=x))
```