

Part 1: Exact Confidence Intervals for Small Sample Random Effects Meta-analysis

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Outline

Introduction

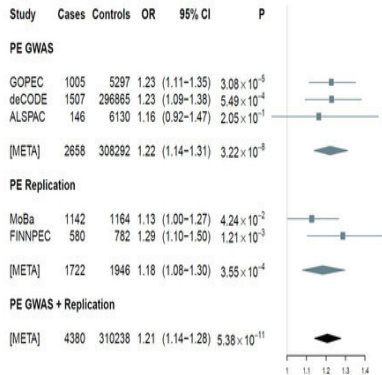
Method

Simulation

Future Work

Meta-analysis is a popular procedure for synthesizing a number of primary studies relating to a single effect into a single summary estimate of size and uncertainty

rs4769613 [hg19: chr13-29138609; risk: C(0.525); other: T; Phet: 0.678]



(from "Variants in the fetal genome near FLT1 are associated with risk of preeclampsia," Nature Genetics, June 2017)

- ▶ The most commonly used model is the random effects model:

$$y_k \stackrel{\text{ind.}}{\sim} \mathcal{N}(\theta_k, \sigma_k^2), k = 1, \dots, K$$

$$\theta_k \stackrel{\text{iid}}{\sim} \mathcal{N}(\Theta, \tau^2)$$

σ_k known

implying

$$y_k \sim \mathcal{N}(\Theta, \sigma_k^2 + \tau^2)$$

- ▶ Goal is inference on Θ
- ▶ τ^2 , accounting for variability between the primary studies, is a nuisance parameter

- ▶ The UMVU estimate of Θ is inverse-variance weighted

$$\frac{\sum_k (\sigma_k^2 + \tau^2)^{-1} y_k}{\sum_k (\sigma_k^2 + \tau^2)^{-1}}$$

with variance

$$\left(\sum_k (\tau^2 + \sigma_k^2)^{-1} \right)^{-1}$$

- ▶ As τ^2 is unknown, typically the DerSimonian-Laird estimator $\hat{\tau}_{DL}^2$ is plugged in

$$\hat{\Theta}_{DL} = \frac{\sum_k (\sigma_k^2 + \hat{\tau}_{DL}^2)^{-1} y_k}{\sum_k (\sigma_k^2 + \hat{\tau}_{DL}^2)^{-1}}$$

► A confidence interval

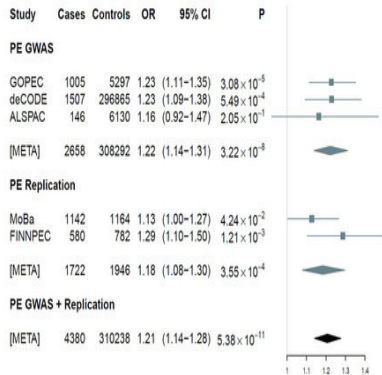
$$\left\{ \hat{\Theta}_{DL} - z_{1-\alpha/2} \left(\sum_{k=1}^K (\hat{\tau}_{DL}^2 + \sigma_k^2)^{-1} \right)^{-1/2}, \hat{\Theta}_{DL} + z_{1-\alpha/2} \left(\sum_{k=1}^K (\hat{\tau}_{DL}^2 + \sigma_k^2)^{-1} \right)^{-1/2} \right\}$$

is obtained from an asymptotic pivot

$$T_0(\Theta; \mathcal{Y}) = (\hat{\Theta}_{DL} - \Theta)^2 \sum_{k=1}^K (\hat{\tau}_{DL}^2 + \sigma_k^2)^{-1} \rightsquigarrow \chi_1^2 \quad (K \rightarrow \infty)$$

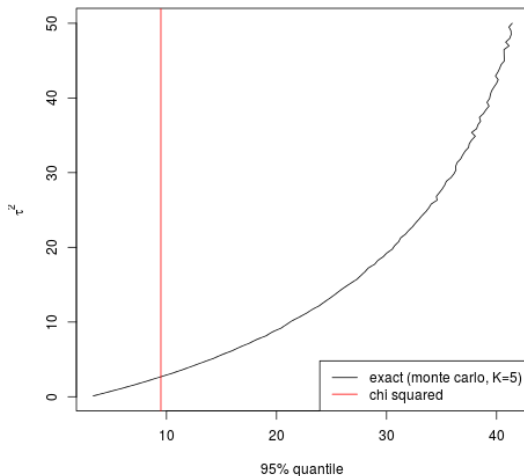
- In many fields, meta-analyses on few (< 6) studies are common
- Even when many primary studies are available, sub-meta-analyses are routinely carried out using few studies

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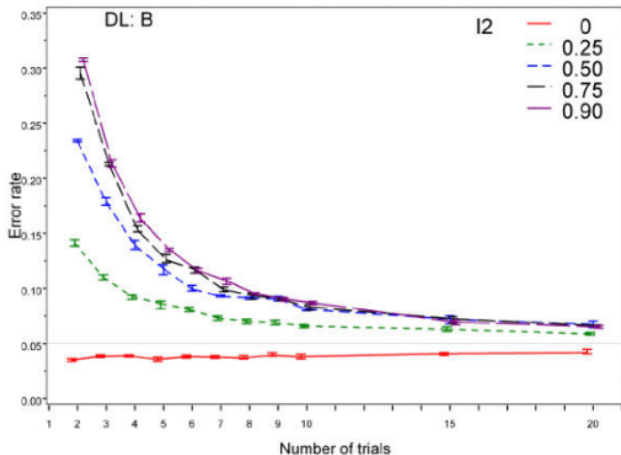


(from "Variants in the fetal genome near FLT1 are associated with risk of preeclampsia," Nature Genetics, June 2017)

- Problem: when the number of studies is few and heterogeneity is present, the pivot is a poor approximation



- ...resulting in poor Type I error control



(from IntHout, Ioannidis and Borm '14)

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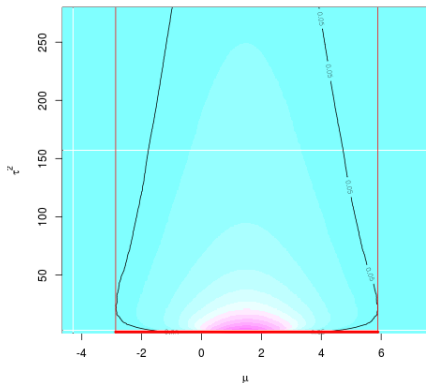
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In the absence of a statistic ancillary to the nuisance parameter, we obtain a CI for Θ at each value of the nuisance parameter and use their union as a conservative CI



Controls the Type I error rate, but at what cost?

- ▶ computational?
- ▶ power?

Computational costs

- ▶ exploit symmetry of the problem
- ▶ $y_k \sim \mathcal{N}(\Theta, \sigma^2 + \tau^2)$ means $y_k - \Theta \sim \mathcal{N}(0, \sigma^2 + \tau^2)$
- ▶ reasonable to require of our testing procedure that testing $H_0 : \Theta = \Theta_0$ given data y_1, \dots, y_K be the same as testing $H_0 : \Theta = 0$ given data $y_1 - \Theta, \dots, y_K - \Theta$
- ▶ *Equivariant* test statistics respect the symmetry of the problem: $T(y_1 - \Theta, \dots, y_K - \Theta) = T(y_1, \dots, y_K) - \Theta$

- ▶ Using an equivariant test statistic we only need to compute the distribution at parameter points $(0, \tau^2)$
 - ▶ can't apply this trick again, not a scale family because of σ_k
- ▶ So the problem is actually 1-dimensional
- ▶ Cost not much of an issue except for statistics that are relatively costly to compute, e.g., MLE
- ▶ Easily parallelized (as is the 2-D problem)

- ▶ Our proposed statistics for testing the simple null $H_0 : (\Theta, \tau^2) = (\Theta_0, \tau_0^2)$:

$$T \{(\Theta_0, \tau_0^2); \mathcal{Y}\} = T_0(\Theta_0; \mathcal{Y}) + c_0 T_{lik} \{(\Theta_0, \tau_0^2); \mathcal{Y}\}$$

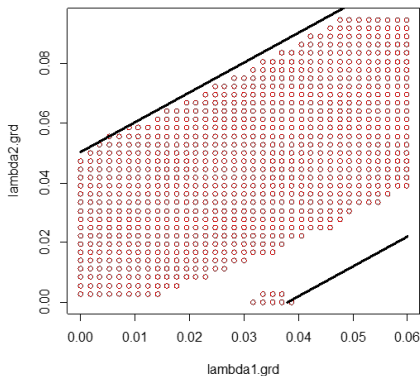
where

$$\begin{aligned} T_0(\Theta_0; \mathcal{Y}) &= (\hat{\Theta}_{DL} - \Theta_0)^2 \sum_{k=1}^K (\hat{\tau}_{DL}^2 + \sigma_k^2)^{-1} \\ T_{lik} \{(\Theta_0, \tau_0^2); \mathcal{X}\} &= -\frac{1}{2} \sum_{k=1}^K \left[\frac{(Y_k - \hat{\Theta}_{DL})^2}{\hat{\tau}_{DL}^2 + \sigma_k^2} + \log \{2\pi(\hat{\tau}_{DL}^2 + \sigma_k^2)\} \right] + \\ &\quad \sum_{k=1}^K \frac{1}{2} \left[\frac{(Y_k - \Theta_0)^2}{\tau_0^2 + \sigma_k^2} + \log \{2\pi(\tau_0^2 + \sigma_k^2)\} \right] \end{aligned}$$

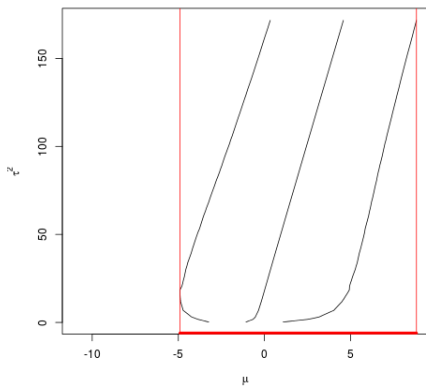
- ▶ Weighted combination of DL statistic and an approximate likelihood ratio statistic
- ▶ Plug DL estimate of Θ into likelihood ratio statistic to avoid computing the MLE
- ▶ These are equivariant

Power, CI length

- Two sources of poor power performance when projecting to form a conservative CI



disconnected regions (from related work by L. Tian)



shear (artist's interpretation)

- ▶ The proposed estimators are quadratic in Θ , so the region is connected
- ▶ The deviation from the vertical of the centers of horizontal sections of the region has variance $O(\tau^{-4})$

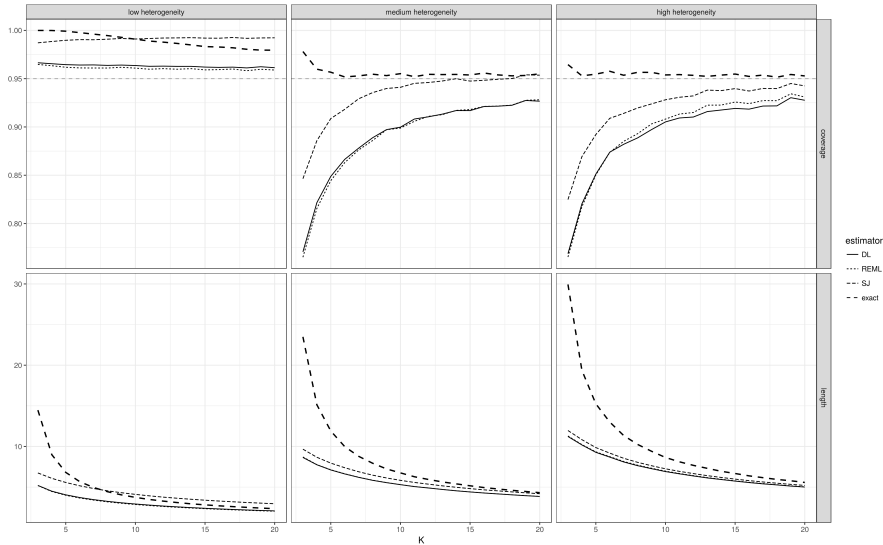
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- ▶ look for similar efficiencies with nonnormal primary study effects data, e.g., rare event proportions
- ▶ examining robustness against nonnormality