# Modeling a latent structure of individual differences in numerical cognition

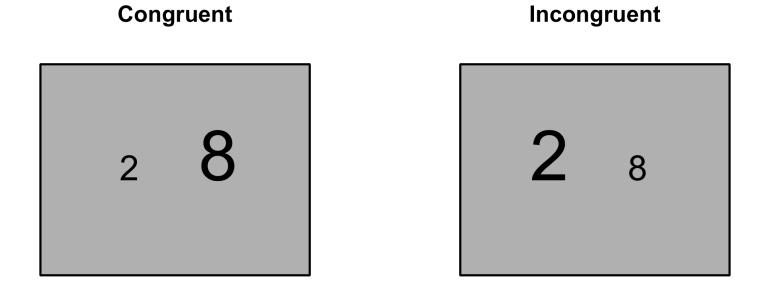
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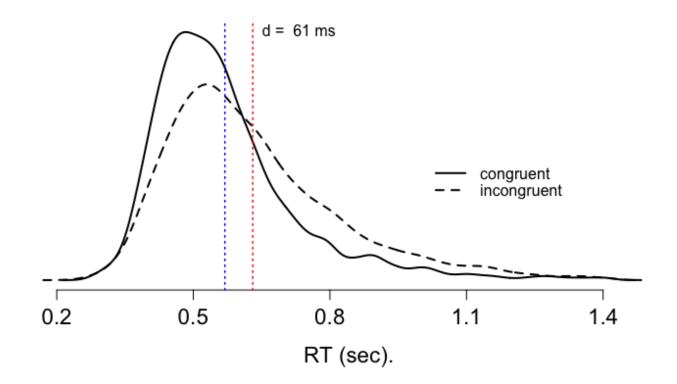
vARMADILLO 2020

# Size congruity effect

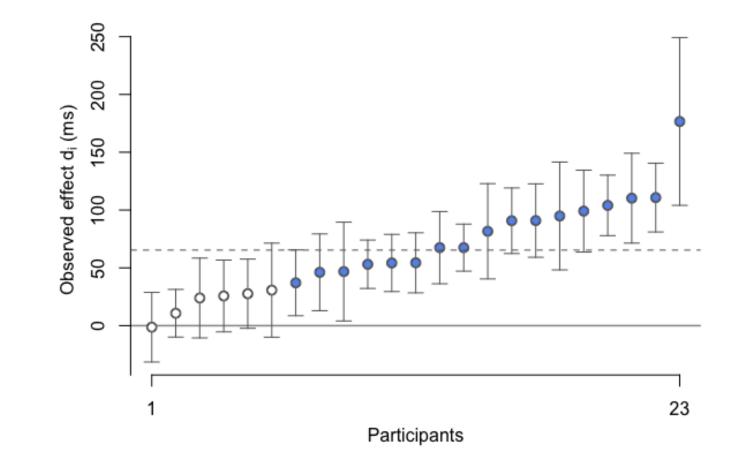
Typical laboratory task: choose the **physically larger** digit



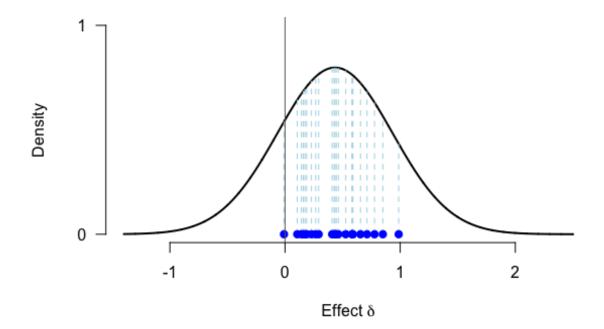
Typical result – mean RT larger for *incongruent trials* 



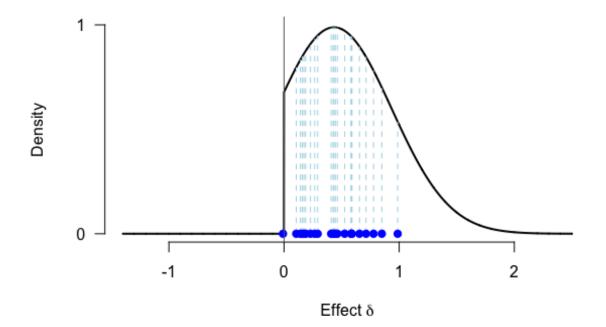
What about individual effects?



Consider these observed effects  $d_i$  as being drawn from (population) distribution of *true* effects  $\delta$ . What is the structure of this population?



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# A new question

**Does everybody** exhibit the size congruity effect?

- if *yes*, then SCE is obligatory, resistant to strategic control, ...
- if *no*, then SCE is complex, malleable, ...

Importantly, both answers have downstream consequences for processing architecture of numerical cognition

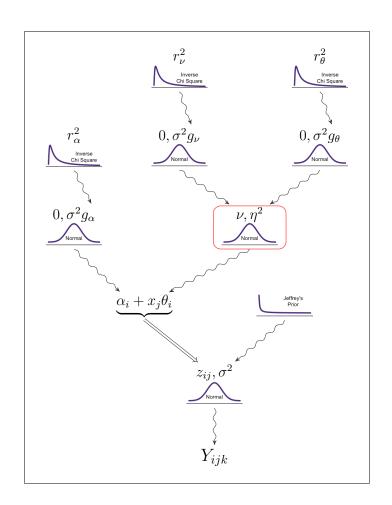
# A new question

**Does everybody** exhibit size congruity effect?

How to answer:

- build competing models of individual difference structures in SCE
- adjudicate the models via Bayesian model comparison

## **Hierarchical structure**



Basic idea (Haaf & Rouder, 2017):

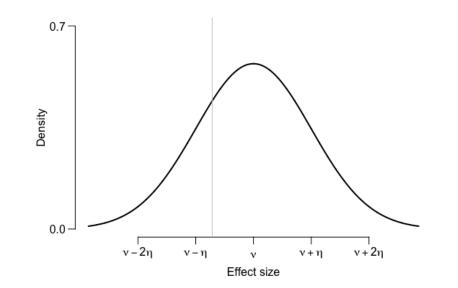
- model RTs as a random-effects linear model with effect parameter θ<sub>i</sub> (i = 1 ..., N)
- assume (as baseline) that  $\theta_i$  is drawn from a normal distribution with mean  $\nu$  and variance  $\eta^2$
- use g-priors (Zellner & Siow, 1980), specifying a priori scale on variance of overall effect and individual variability around effect
- define competing models by
  constraining effect parameter θ<sub>i</sub>

# Four competing models

- 1. Unconstrained model,  $\mathcal{M}_u$
- 2. Positive-effects model,  $\mathcal{M}_+$
- 3. Common-effect model,  $\mathcal{M}_1$
- 4. Null-effect model,  $\mathcal{M}_0$

# **Unconstrained model**

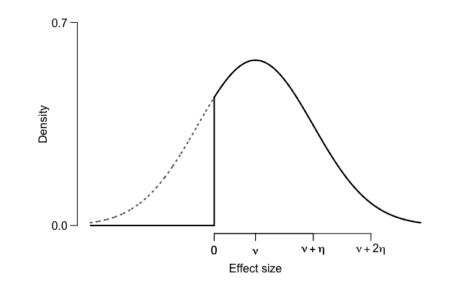
 $\mathcal{M}_u: \theta_i \sim \mathsf{Normal}(\nu, \eta^2)$ 



Qualitative individual differences: some positive, some negative

## **Positive-effects model**

•  $\mathcal{M}_+: \theta_i \sim \text{Normal}_+(\nu, \eta^2)$ 

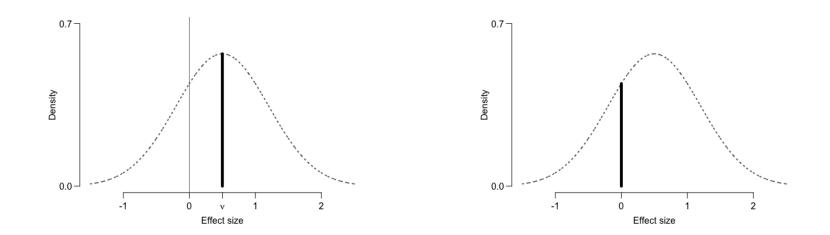


Quantitative individual differences: all positive, vary in magnitude

# **Common-effect** / null models

•  $\mathcal{M}_1: \theta_i = \nu$ 

•  $\mathcal{M}_0: \theta_i = 0$ 



## Model comparison

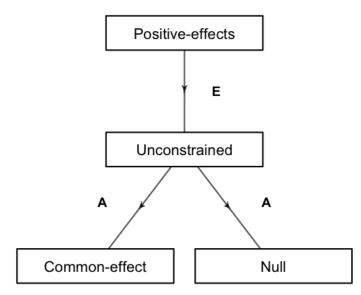
We use the **Bayes factor**, which indexes how well the observed data Y are predicted under one model relative to another:

$$B_{ab} = \frac{p(\boldsymbol{Y} \mid \mathcal{M}_a)}{p(\boldsymbol{Y} \mid \mathcal{M}_b)}$$

 $B_{ab} = 10$  means:

- the observed data are 10 times more likely under  $\mathcal{M}_a$  compared to  $\mathcal{M}_b$
- "10-to-1 evidence for  $\mathcal{M}_a$  over  $\mathcal{M}_b$ "

How do we compute 
$$B_{ab} = \frac{p(\boldsymbol{Y} \mid \mathcal{M}_a)}{p(\boldsymbol{Y} \mid \mathcal{M}_b)}$$
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$$p(\boldsymbol{Y} \mid \mathcal{M}) = \int_{\boldsymbol{\xi} \in \Xi} p(\boldsymbol{Y} \mid \boldsymbol{\xi}) p(\boldsymbol{\xi}) d\boldsymbol{\xi}$$

How do we compute  $B_{ab} = \frac{p(\boldsymbol{Y} \mid \mathcal{M}_a)}{p(\boldsymbol{Y} \mid \mathcal{M}_b)}$ ?

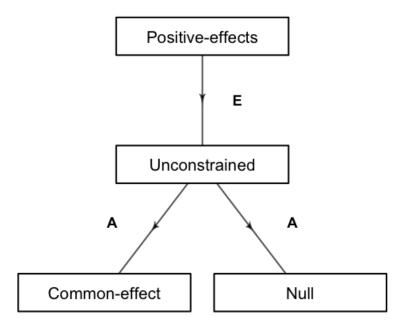
$$p(\boldsymbol{Y} \mid \mathcal{M}) = \int_{\boldsymbol{\xi} \in \Xi} p(\boldsymbol{Y} \mid \boldsymbol{\xi}) p(\boldsymbol{\xi}) d\boldsymbol{\xi}$$

Problem: for our models  $\mathcal{M}$ , the parameter vectors  $\boldsymbol{\xi}$  look like

$$\boldsymbol{\xi} = (\mu, \sigma^2, \nu, \alpha_1, \dots, \alpha_N, \theta_1, \dots, \theta_N, g_\alpha, g_\nu, g_\theta)$$

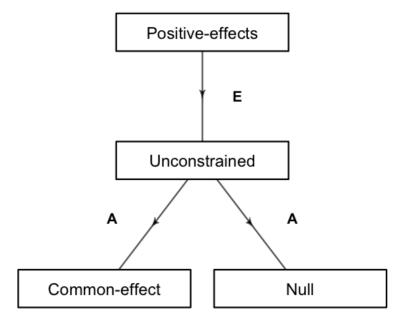
so the integral is carried out in  $\mathbb{R}^{2N+6}$ .

For N = 35, this would be a 76-dimensional integral!



A = analytic approach

- Zellner & Siow (1980); Rouder et al. (2012)
- place *g*-priors on individual intercepts and effect parameters
- everything *except* the *g*-parameters integrates symbolically
- *g*-parameters can be well approximated with MCMC sampling
- techniques coded into BayesFactor package in R



- E = encompassing approach
- Klugkist et al. (2005)

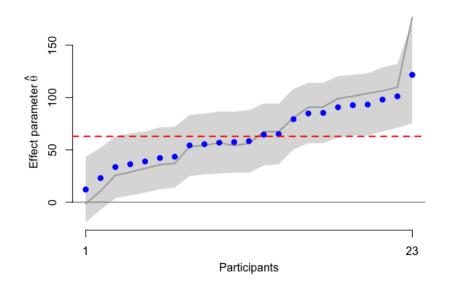
• 
$$B_{+u} = \frac{P(\boldsymbol{\theta} > 0 \mid \boldsymbol{Y}, \mathcal{M}_u)}{P(\boldsymbol{\theta} > 0 \mid \mathcal{M}_u)}$$

 probabilities computed as fraction of MCMC samples from unrestricted model that are **positive** for all individuals (both *a priori* and *a posteriori*)

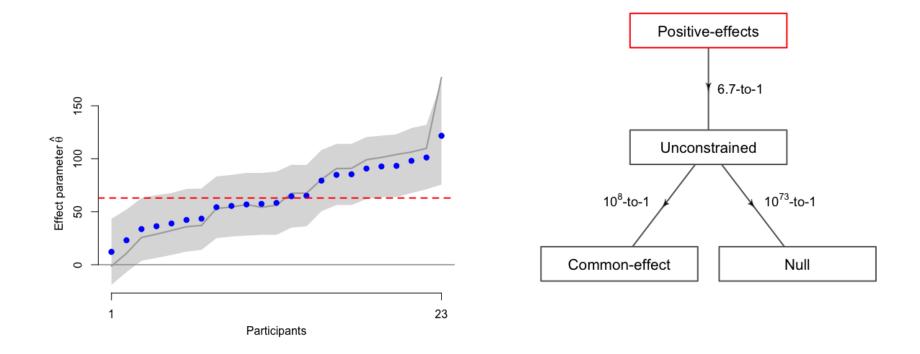
## Datasets

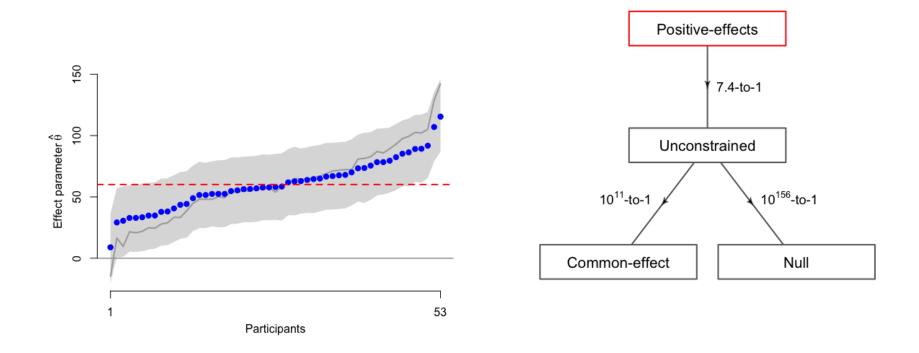
We modeled three datasets, each using the same basic size-congruity task.

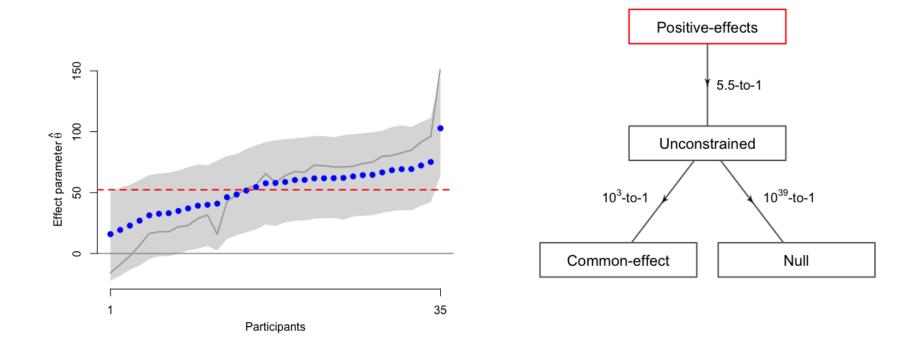
Dataset	N	# obs	Mean RT	SD	Error rate
1	23	8,832	614 ms	349 ms	5.6%
2	53	20,352	593 ms	299 ms	4.2%
3	35	6,720	674 ms	354 ms	4.5%



- Red line = estimated effect  $\theta$  from  $\mathcal{M}_1$
- Blue dots = individual effect estimates  $\theta_i$
- Gray line = estimates from mean differences  $d_i$
- Gray area = 95% credible intervals







# Summary points

**Does everybody** exhibit the size congruity effect?

- if *yes*, then SCE is obligatory, resistant to strategic control, ...
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 what does this say about *early vs. late* interaction debate (e.g., Faulkenberry et al., 2016; Sobel et al., 2016; 2017; Faulkenberry, Vick, & Bowman, 2018; Bowman, 2020) Some other benefits:

- Bayes factors easy to interpret
- hierarchical structure removes trial noise from individual estimates
- common effect (CE) model provides important self-check:
  - if CE model is best, is our design adequate to capture individual differences
- Might be good approach to disentangle competing theories of mental arithmetic
  - Does everyone exhibit size-by-format interaction?
  - Does everyone reflect fast counting in small addition problems?

#### Thank you!

- slides available at https://tomfaulkenberry.github.io/talks.html
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