

Hierarchical Models

PSYC 573

University of Southern California

March 3, 2022

Therapeutic Touch Example ($N = 28$)

Data Points From One Person

y : whether the guess of
which hand was hovered over
was correct

Person S01

y	s
1	S01
0	S01
0	S01
0	S01
0	S01
0	S01
0	S01
0	S01
0	S01
0	S01

Binomial Model

We can use a Bernoulli model:

$$y_i \sim \text{Bern}(\theta)$$

for $i = 1, \dots, N$

Assuming exchangeability given θ , more succinct to write

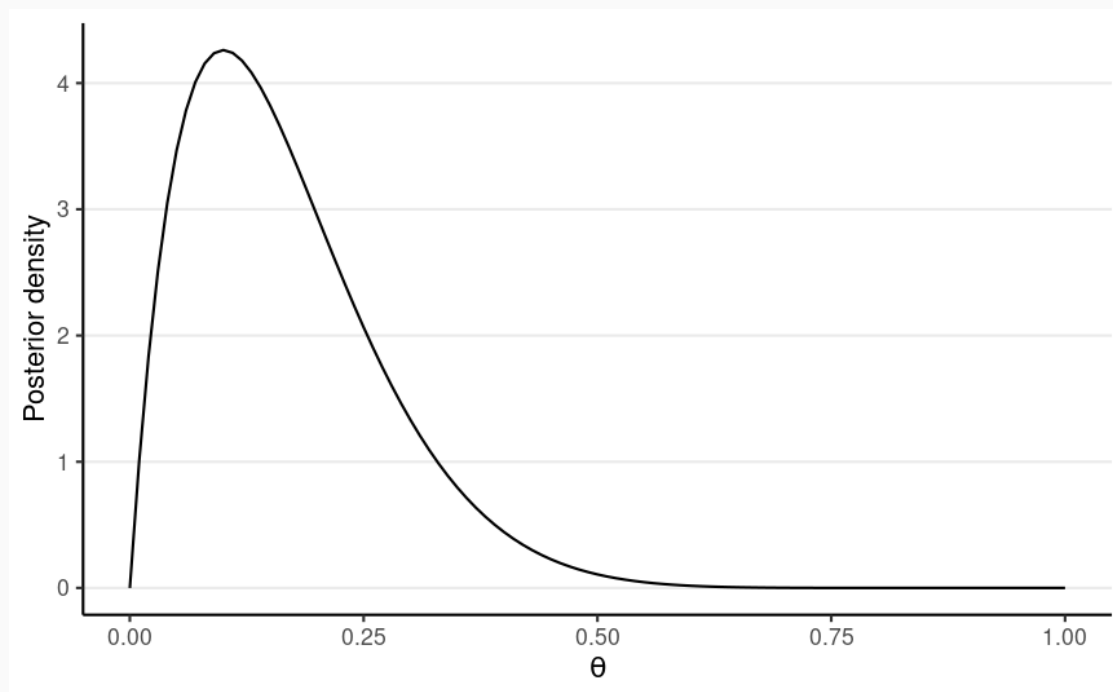
$$z \sim \text{Bin}(N, \theta)$$

for $z = \sum_{i=1}^N y_i$

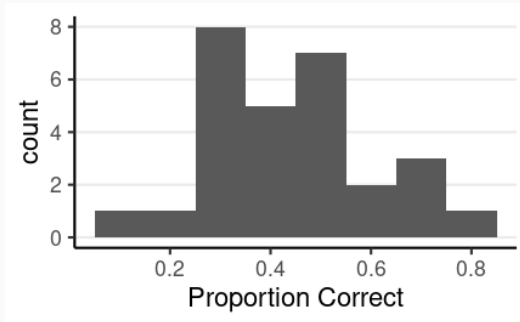
- Bernoulli: Individual trial
- Binomial: total count of "1"s

1 success, 9 failures

Posterior: Beta(2, 10)



Multiple People



We could repeat the binomial model for each of the 28 participants, to obtain posteriors for $\theta_1, \dots, \theta_{28}$

But . . .

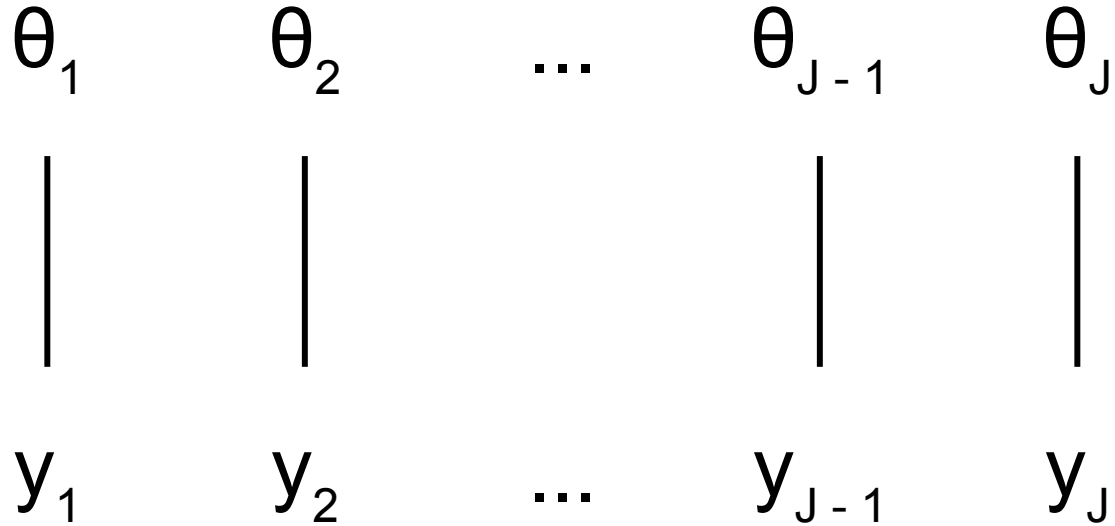
Do we think our belief about θ_1 would inform our belief about θ_2 , etc?

After all, human beings share 99.9% of genetic makeup

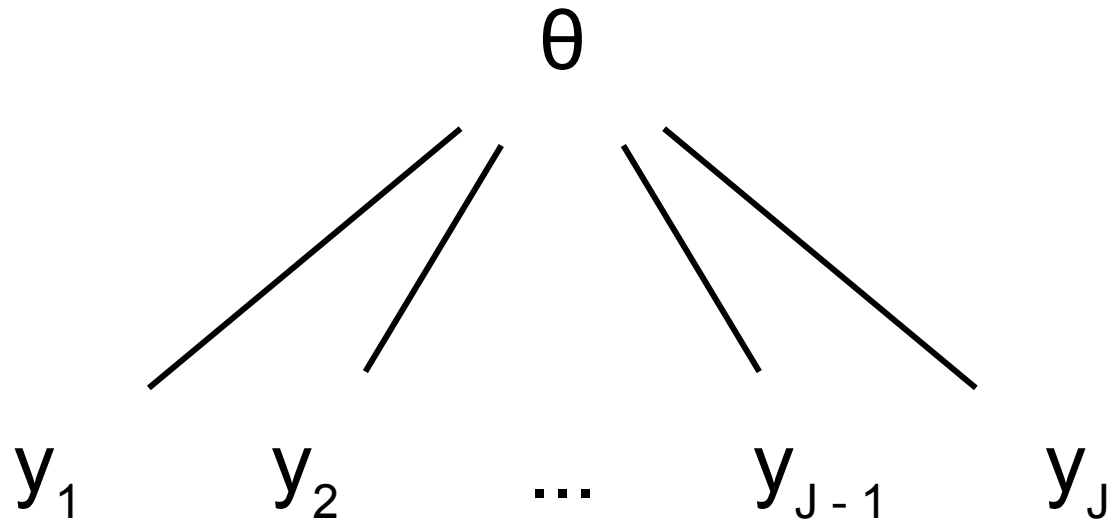
Three Positions of Pooling

- No pooling: each individual is completely different; inference of θ_1 should be independent of θ_2 , etc
- Complete pooling: each individual is exactly the same; just one θ instead of 28 θ_j 's
- **Partial pooling**: each individual has something in common but also is somewhat different

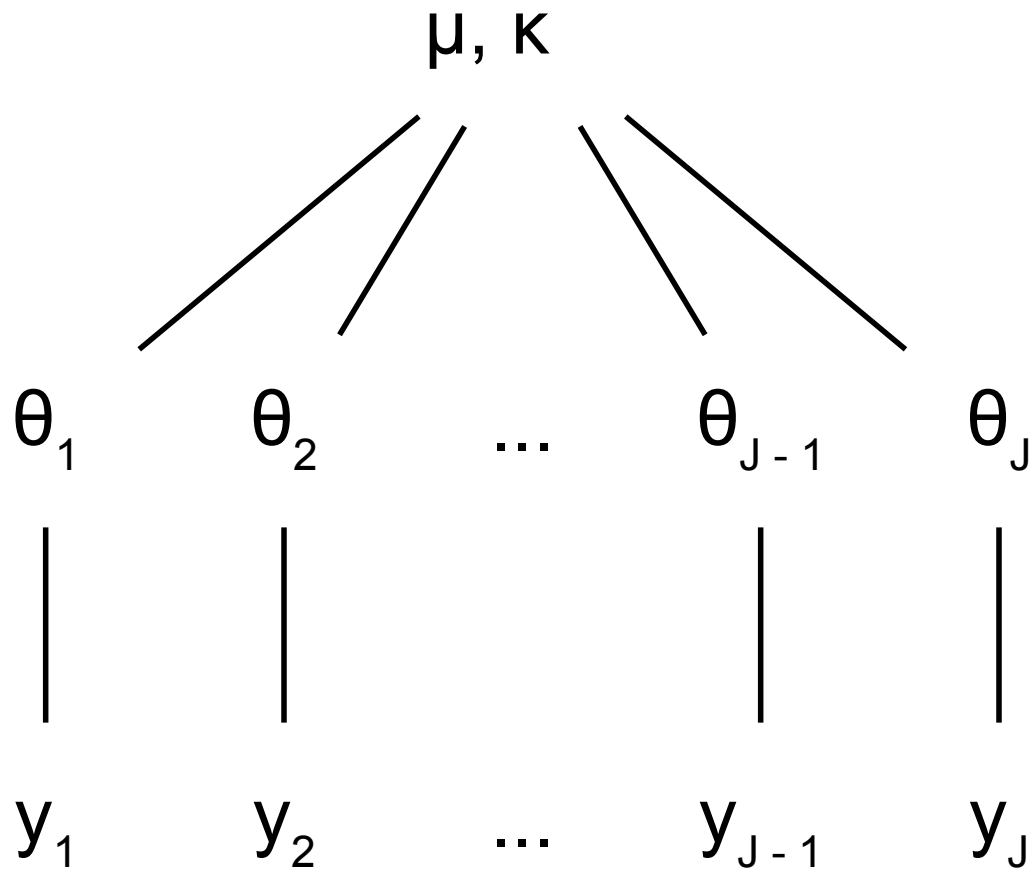
No Pooling



Complete Pooling



Partial Pooling



Partial Pooling in Hierarchical Models

Hierarchical Priors: $\theta_j \sim \text{Beta2}(\mu, \kappa)$

Beta2: *reparameterized* Beta distribution

- mean $\mu = a / (a + b)$
- concentration $\kappa = a + b$

Expresses the prior belief:

Individual θ s follow a common Beta distribution with mean μ and concentration κ

How to Choose κ

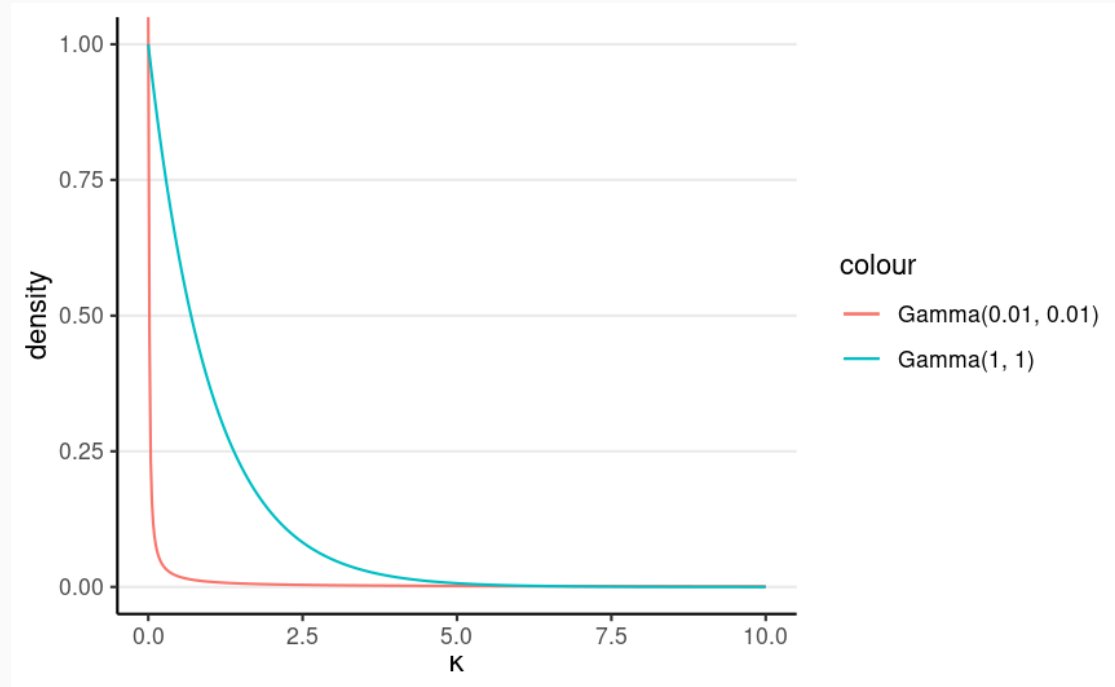
If $\kappa \rightarrow \infty$: everyone is the same; no individual differences (i.e., complete pooling)

If $\kappa = 0$: everybody is different; nothing is shared (i.e., no pooling)

We can fix a κ value based on our belief of how individuals are similar or different

A more Bayesian approach is to treat κ as an unknown, and use Bayesian inference to update our belief about κ

Generic prior by Kruschke (2015): $\kappa \sim \text{Gamma}(0.01, 0.01)$



Sometimes you may want a stronger prior like Gamma(1, 1), if it is unrealistic to do no pooling

Full Model

Model

Stan code

Model:

$$z_j \sim \text{Bin}(N_j, \theta_j)$$

$$\theta_j \sim \text{Beta2}(\mu, \kappa)$$

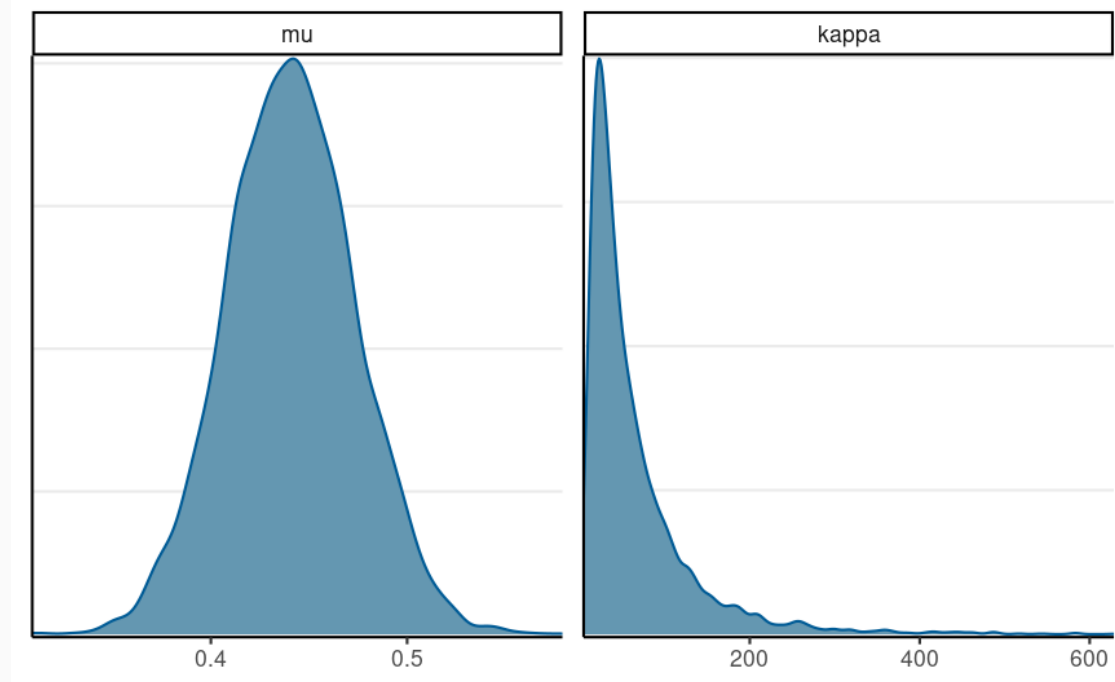
Prior:

$$\mu \sim \text{Beta}(1.5, 1.5)$$

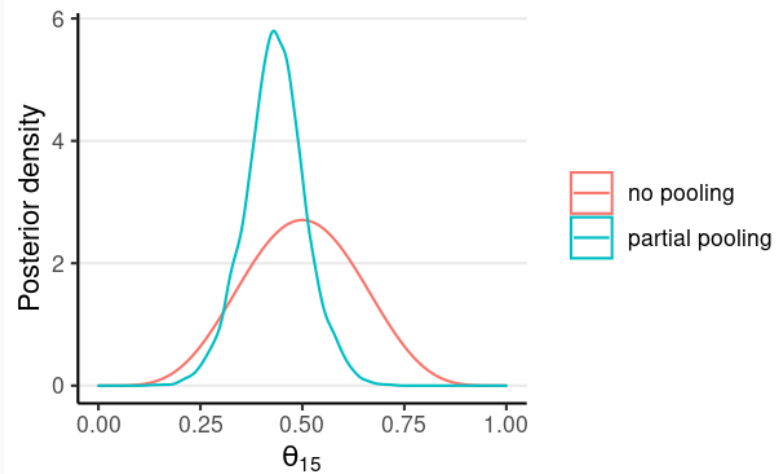
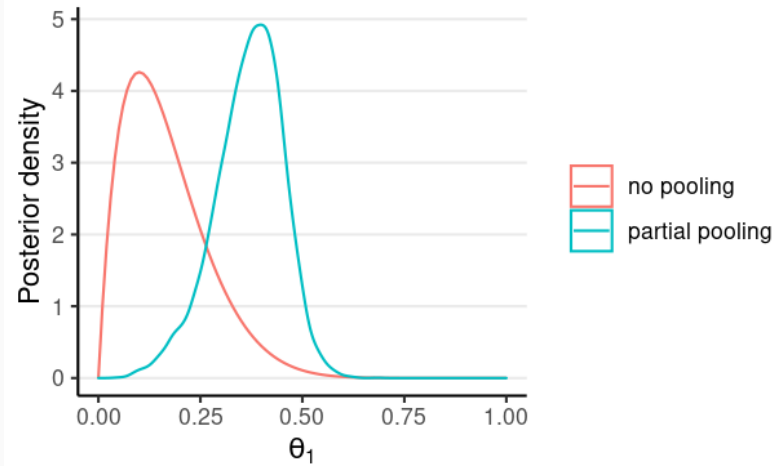
$$\kappa \sim \text{Gamma}(0.01, 0.01)$$

Posterior of Hyperparameters

```
library(bayesplot)
mcmc_dens(tt_fit, pars = c("mu", "kappa"))
```



Shrinkage



Multiple Comparisons?

Frequentist: family-wise error rate depends on the number of intended contrasts

Bayesian: only one posterior; hierarchical priors already express the possibility that groups are the same

Thus, Bayesian hierarchical model "completely solves the multiple comparisons problem."¹

[1]: see <https://statmodeling.stat.columbia.edu/2016/08/22/bayesian-inference-completely-solves-the-multiple-comparisons-problem/>

[2]: See more in ch 11.4 of Kruschke (2015)

Hierarchical Normal Model

Effect of coaching on SAT-V

School	Treatment Effect Estimate	Standard Error
A	28	15
B	8	10
C	-3	16
D	7	11
E	-1	9
F	1	11
G	18	10
H	12	18

Model

Stan code

Model:

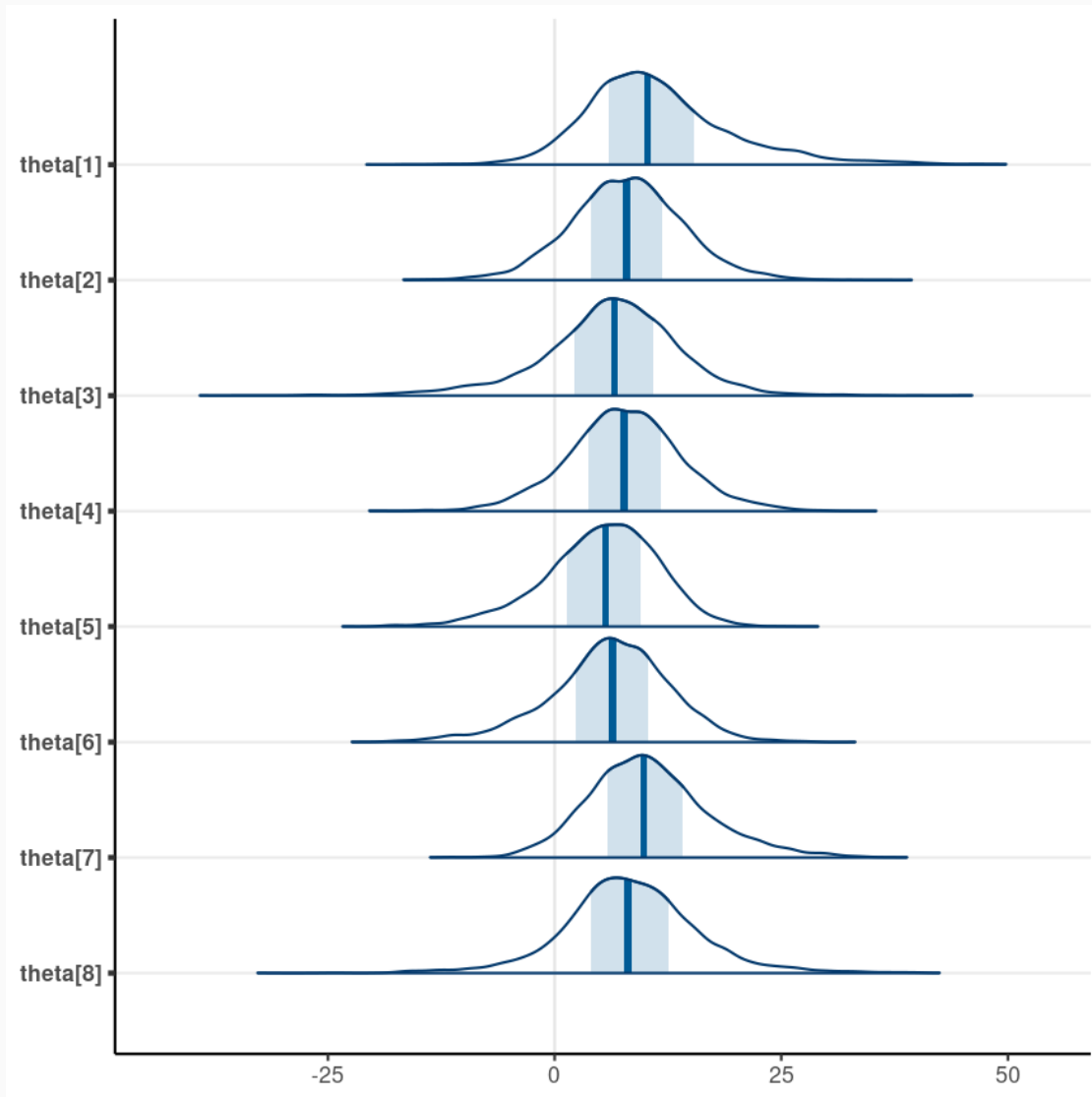
$$d_j \sim N(\theta_j, s_j)$$

$$\theta_j \sim N(\mu, \tau)$$

Prior:

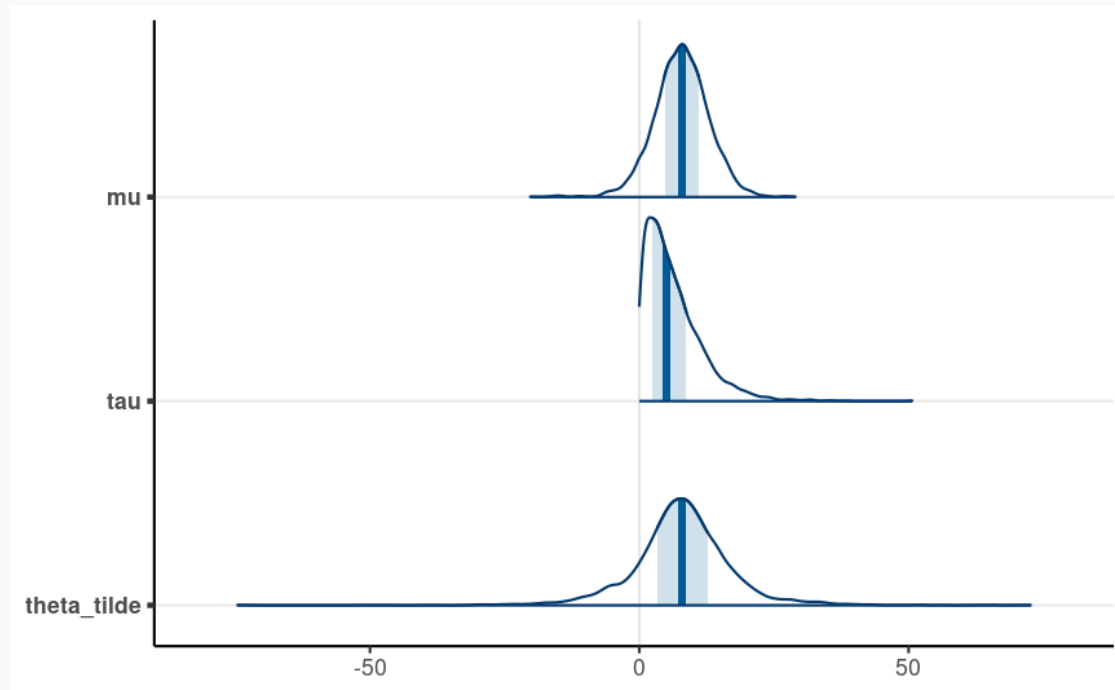
$$\mu \sim N(0, 100)$$

$$\tau \sim t_4^+(0, 100)$$



Prediction Interval

Posterior distribution of the true effect size of a new study, $\tilde{\theta}$



See <https://onlinelibrary.wiley.com/doi/abs/10.1002/jrsm.12> for an introductory paper on random-effect meta-analysis