

# Sample Size Planning for MLM

PSYC 575

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# Week Learning Objectives

- Describe the importance of having sufficient sample size for scientific research
- Describe conceptually the steps for sample size planning: precision analysis and power analysis
- Perform power analysis for MLM using the PowerUpR application and the `simr` package
- Understand the effect of uncertainty in parameter values and explore alternative approaches for sample size planning

# Why Sample Size?

# Small Sample Size is a Problem Because . . .

Low power

Misleading and noisy results<sup>1</sup>

- When coupled with publication bias (statistical significance filter)<sup>2 3</sup>

Nonreproducible findings

[1] See [Maxwell \(2004\)](#)

[2] See the graph on [this blog post](#)

[3] See also [Vasishth et al. \(2018\)](#)

# Review: Sampling distributions

Test yourself! -- Week 13 Quiz (ungraded)

What is the null distribution?

- Suppose we examine the effect of a therapy on eating disorder
- We test against the null hypothesis  $H_0 : \gamma_{01} = 0$ , where  $\gamma_{01}$  is the fixed effect of the therapy on eating disorder

What is the alternative distribution?

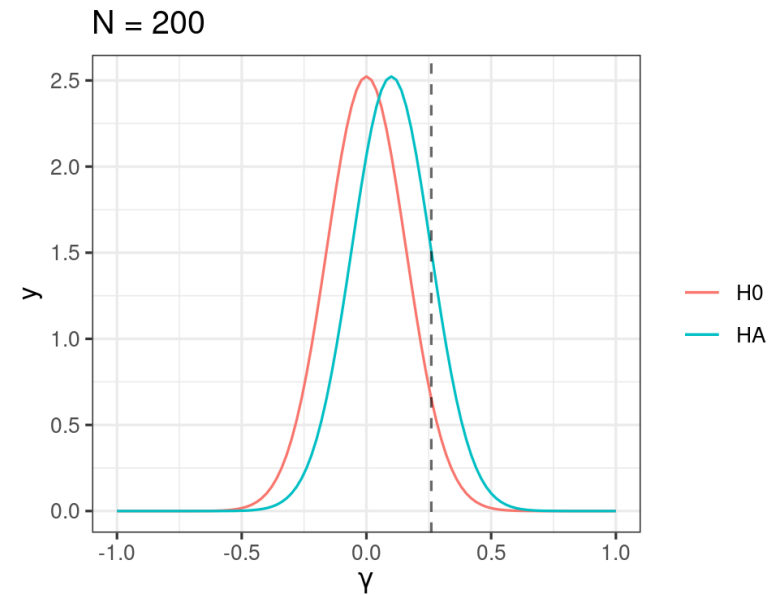
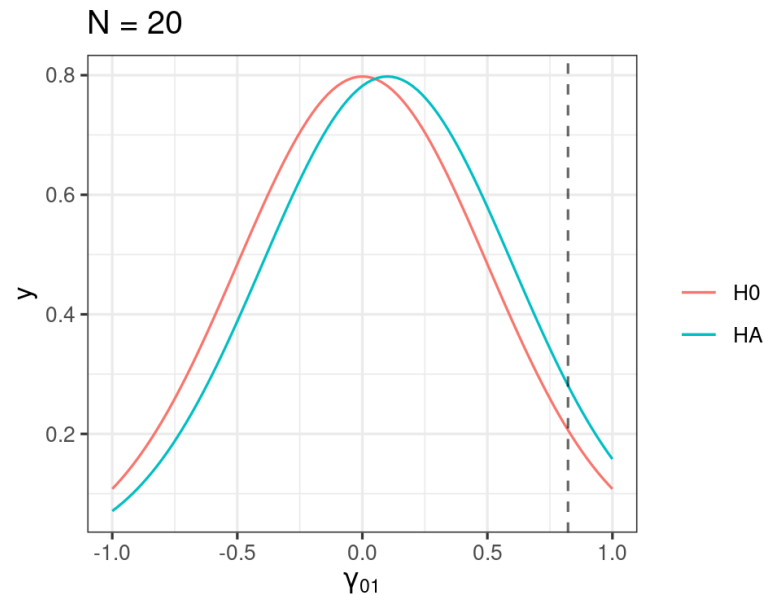
- Assume that the true effect of this therapy is  $\gamma_{01} = .1$

# Sampling Distribution as a Function of Sample Size

Assume true effect is  $\gamma_{01} = 0.10$

Let's say

- when  $N = 20$ ,  $p < .05$  when  $\hat{\gamma} \geq 0.82$
- when  $N = 200$ ,  $p < .05$  when  $\hat{\gamma} \geq 0.26$



# Steps for Sample Size Planning

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1. Write down your model equations
2. List out all parameters in the model
3. Determine if you want to achieve a desired level of
  - a. Power, or
  - b. Precision



# Step 1: Write down model equations

Group-based therapy for eating disorder (cluster-randomized trial)

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Level-1

$$Y_{ij} = \beta_{0j} + \beta_{1j}X_{cmc_{ij}} + e_{ij}$$

$$e_{ij} \sim N(0, \sigma)$$

Level-2

$$\beta_{0j} = \gamma_{00} + \gamma_{01}W_j + u_{0j}$$

$$\beta_{1j} = \gamma_{10} + \gamma_{11}W_j + u_{1j}$$

$$\begin{bmatrix} u_{0j} \\ u_{1j} \end{bmatrix} \sim N \left( \begin{bmatrix} 0 \\ 0 \end{bmatrix}, \begin{bmatrix} \tau_0^2 & \\ \tau_{01} & \tau_1^2 \end{bmatrix} \right)$$

- $\gamma_{10}$ :  $X$  (purely level-1 with ICC = 0)
- $\gamma_{01}$ :  $W$  (level-2)
- $\gamma_{11}$ :  $W \times X$  (cross-level interaction)

# Step 2: List out all parameters

1. Fixed effects:  $\gamma_{00}, \gamma_{01}, \gamma_{10}, \gamma_{11}$

2. Random effects:  $\tau_0^2, \tau_1^2, \tau_{01}$

3. Number of clusters:  $J$

4. Cluster size:  $n$

Level-1

$$Y_{ij} = \beta_{0j} + \beta_{1j}X_{\text{cmc}}c_{ij} + e_{ij}$$

$$e_{ij} \sim N(0, \sigma)$$

Level-2

$$\beta_{0j} = \gamma_{00} + \gamma_{01}W_j + u_{0j}$$

$$\beta_{1j} = \gamma_{10} + \gamma_{11}W_j + u_{1j}$$

$$\begin{bmatrix} u_{0j} \\ u_{1j} \end{bmatrix} \sim N \left( \begin{bmatrix} 0 \\ 0 \end{bmatrix}, \begin{bmatrix} \tau_0^2 & \\ & \tau_1^2 \end{bmatrix} \right)$$

# Standard Error and Precision Analysis

# Sample Size and $SE/Post. SD$

In the previous graph, when  $N = 20$ , the sample estimate is likely to be anywhere between -0.4 and 0.6

$$SE \propto \frac{1}{\sqrt{N}}$$

One goal of sample size planning is to

- Have sufficient sample size to get precise (low  $SE$ ) sample estimates of an effect

# Analytic Formulas of $SE$

$J$  = Number of clusters;  $n$  = Cluster size

- E.g.,  $J = 100$  schools;  $n = 10$  students per school

Assuming  $\tau_{01} = 0$

$$SE(\gamma_{01}) = \sqrt{\frac{1}{S_W^2} \left( \frac{\tau_0^2}{J} + \frac{\sigma^2}{Jn} \right)}$$

$$SE(\gamma_{10}) = \sqrt{\frac{\tau_1^2}{J} + \frac{\sigma^2}{JnS_X^2}}$$

$$SE(\gamma_{11}) = \sqrt{\frac{1}{S_W^2} \left( \frac{\tau_1^2}{J} + \frac{\sigma^2}{JnS_X^2} \right)}$$

# Precision Analysis

Group-based therapy for eating disorder (cluster-randomized trial)

- Intervention at group level
- 10 participants per group
- Outcome standardized (i.e.,  $SD = \sqrt{\tau_0^2 + \sigma^2} = 1$ )
  - $\gamma =$  Cohen's  $d$
- ICC = .3 (i.e.,  $\tau_0^2 = .3$ )
- Goal: estimate  $J$  such that  $SE(\gamma_{10}) \leq .1$ 
  - E.g., if we estimated the sample effect size to be  $d = .25$ , the 95% CI would be approximately [.05, .45].

# Calculating $J$

When the predictor is binary (e.g., treatment-control), if half of the groups is in one condition,  $S_W^2 = 0.25$

- Otherwise, if 30% in one condition,  $S_W^2 = 0.3 \times 0.7$
- $\tau_0^2 = 0.3, \sigma^2 = 0.7, n = 10$

E.g., if  $J = 30$

$$SE(\gamma_{01}) = \sqrt{\frac{1}{S_W^2} \left( \frac{\tau_0^2}{J} + \frac{\sigma^2}{Jn} \right)} = \sqrt{\frac{1}{0.25} \left( \frac{0.3}{30} + \frac{0.7}{(30)(10)} \right)} = 0.2221111$$

Keep trying, and you'll find ...

When  $J = 148$ ,  $SE(\gamma_{01}) = 0.1$

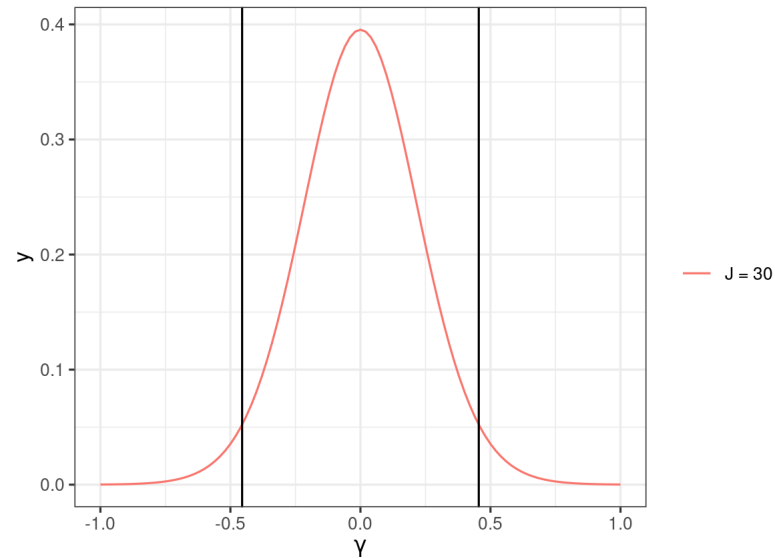
So you'll need 148 groups (74 treatment, 74 control)



# Power Analysis

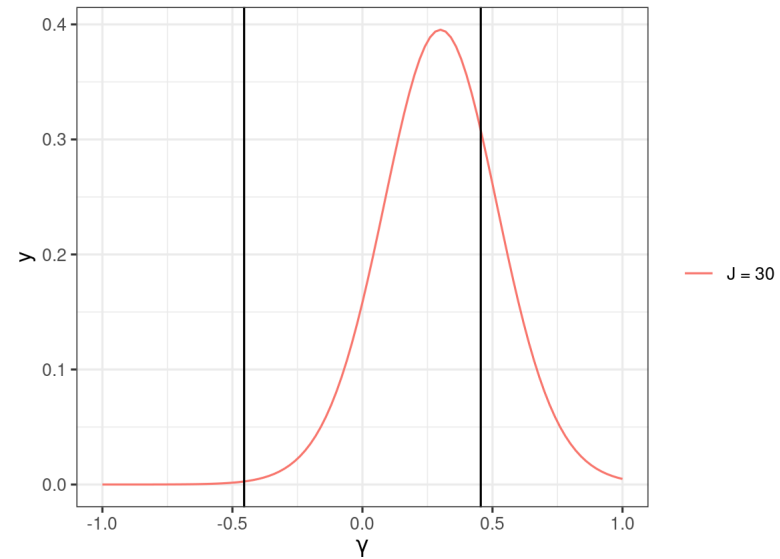
Two-tailed test,  $\alpha = .05$

$$H_0 : \gamma_{01} = 0$$



Critical region:  $\hat{\gamma}_{01} \leq -0.45$  or  $\hat{\gamma}_{01} \geq 0.45$

$$H_1 : \gamma_{01} = 0.3$$



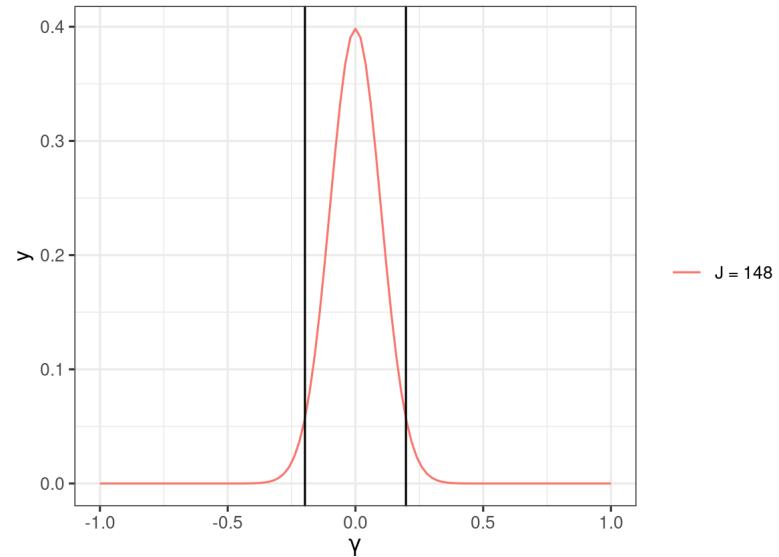
Power<sup>1</sup>

$$\approx P(\hat{\gamma}_{01} \leq -0.45) + P(\hat{\gamma}_{01} \geq 0.45) = 0.2465731$$

[1] In practice, we need to incorporate the sampling variability of the standard error as well, so this power calculation is only a rough approximation.

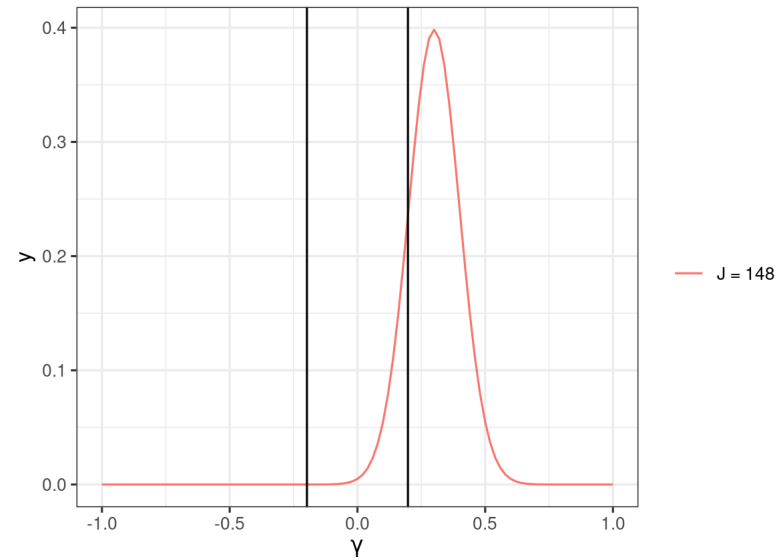
Two-tailed test,  $\alpha = .05$

$$H_0 : \gamma_{01} = 0$$



Critical region:  $\hat{\gamma}_{01} \leq -0.2$  or  $\hat{\gamma}_{01} \geq 0.2$

$$H_1 : \gamma_{01} = 0.3$$



Power

$$\approx P(\hat{\gamma}_{01} \leq -0.2) + P(\hat{\gamma}_{01} \geq 0.2) = 0.8461551$$

# Tools for Power Analysis

## 1. Stand-alone programs

- [Optimal Design](#)
- [PinT](#)

## 2. R packages

- `simr`

## 3. Spreadsheet/Webapp

- [PowerUp!](#)

See more discussion in [Arend & Schäfer \(2019\)](#)

# PowerUpR Shiny App

<https://powerupr.shinyapps.io/index/>

# Monte Carlo Simulation for Power Analysis

- Simulate a large number (e.g.,  $R = 1,000$ ) of data sets based on given effect size, ICC, etc
- Fit an MLM to each simulated data
- Power  $\approx$  Proportion of times  $p < \alpha$

See sample R code for using `simr`

# Uncertainty in Parameter Values

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In the PowerUpR demo, to calculate the number of clusters  $J$  need to achieve 80% power, we determined

1. Type I error rate = .05
2. Two tailed test = TRUE
3.  $g_2, r_{21}, r_{22} = 0$ , as we did not include any covariates
4.  $p = .5$ , for a balanced design (half treatment, half control)

However, we need to guess the values of

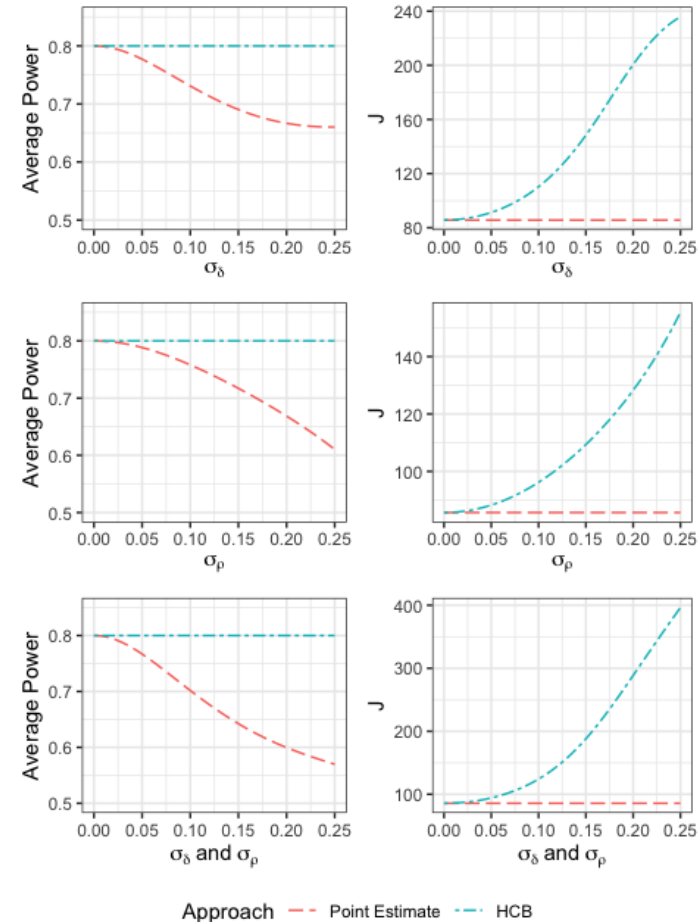
1. Effect size = .3?
2. ICC = .3?



# The Effect of Uncertainty in Power

## Ignoring uncertainty

- The more uncertainty we have but ignore about a parameter value, the more power loss we will have in our study (red curve)
- Uncertainty in both effect size and ICC can further reduce our power
- The more uncertainty we have, the more samples we need to achieve 80% power

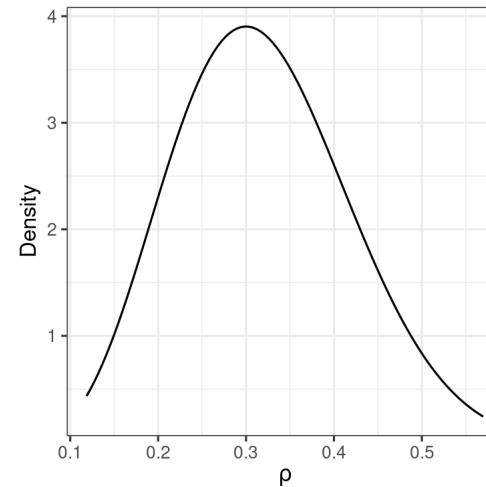
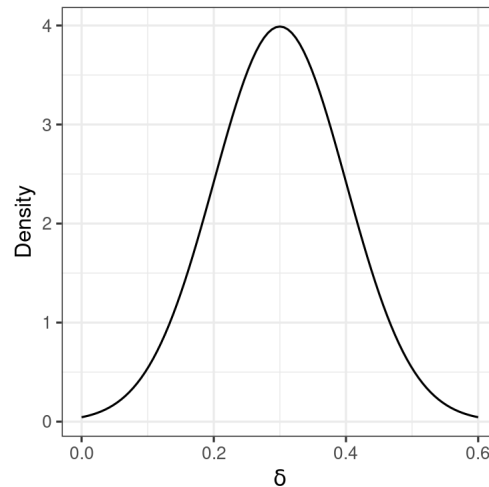


# Hybrid Classical-Bayesian approach

- Incorporates uncertainty for sample size planning
- Instead of plugging in a point value of a guess, we can specify how much uncertainty we have (e.g., standard error of  $\gamma_{01}$  from a previous study)

$$\delta \sim N(.3, .1) \quad \rho \sim \text{Beta}(a, b)$$

- where  $a, b$  can be calculated by  $\hat{\rho} = .3$  and  $\sigma_{\rho} = .1$  (estimate and uncertainty about  $\rho$ )



# hcbR Shiny App

[http://winnie-wy-tse.shinyapps.io/hcb\\_shiny](http://winnie-wy-tse.shinyapps.io/hcb_shiny)

# Additional Notes on Power

- Increasing  $J$  usually leads to higher power than increasing  $n$
- Balanced designs generally have higher power than unbalanced designs
- Larger sample size required for testing level-2 predictors
- Testing an interaction requires a much larger sample size
  - E.g., 16 times larger than for a main effect