

hw3

1

```
dgp_p1 <- function() {  
  ### Define number of schools, and number of students in the school  
  G <- 50 # number of schools  
  ng <- 10 # number of students in each school  
  n <- G*ng # total n-size  
  
  ### Assign students to schools  
  school <- rep(seq(1,G), ng)  
  school <- sort(school)  
  
  ### Gamma (school-varying intercept for PO's)  
  gamma <- rnorm(G, sd=sqrt(2))[school]  
  
  ### Generate potential outcomes  
  y0 <- gamma + rnorm(n)  
  y1 <- 2 + gamma + rnorm(n)  
  
  ### Put everything into a data-frame  
  data <- data.frame("school" = school, "y0" = y0, "y1" = y1)  
  
  return(data)  
}
```

a

There are 10 students per school and 50 schools total, meaning that there are 500 total students. The ATE here is 2, and the factors that influence the potential outcomes are some random noise from the $\mathcal{N}(0, 1)$ instances added to each one, as well as the school specific random intercept called ‘gamma’.

b

- 1) Complete: Randomly draw 250 students to be treated
- 2) Bernoulli: Probability of treatment for each student is 0.5
- 3) Cluster: Randomly draw 25 schools to be treated
- 4) Stratified: Within each school, randomly draw 5 students to be treated

```
library(tidyverse)

# empty results df to fill
res <- tibble(
  complete = rep(0, 1000),
  bernoulli = rep(0, 1000),
  cluster = rep(0, 1000),
  stratified = rep(0, 1000)
)

for (i in 1:1000) {
  data <- dgp_p1()

  # complete
  row_ids <- sample(1:500, 250)
  complete <- data %>%
    mutate(
      id = row_number(),
      D = if_else(id %in% row_ids, 1, 0)
    ) %>%
    select(-id)

  # bernoulli
  bernoulli <- data %>%
    mutate(D = rbinom(500, 1, 0.5))

  # cluster
  school_ids <- sample(1:50, 25)
  cluster <- data %>%
    group_by(school) %>%
    mutate(
      group_id = cur_group_id(),
      D = if_else(group_id %in% school_ids, 1, 0)
    ) %>%
    ungroup() %>%
```

```

    select(-group_id)

# stratified
within_school_ids <- data %>%
  group_by(school) %>%
  mutate(chosen_id = row_number()) %>%
  slice_sample(n = 5) %>%
  ungroup() %>%
  select(school, chosen_id) %>%
  mutate(D = 1)

stratified <- data %>%
  group_by(school) %>%
  mutate(
    id = row_number()
  ) %>%
  ungroup() %>%
  left_join(
    within_school_ids, by = c("id" = "chosen_id", "school")
  ) %>%
  mutate(D = replace_na(D, 0)) %>%
  select(-id)

data_list <- list(complete, bernoulli, cluster, stratified)

dim <- function(data){

  return(mean(data[data$D == 1, ]$y1) - mean(data[data$D == 0, ]$y0))

}

res[i, ] <- data_list %>%
  map(.f = dim)

}

res %>%
  pivot_longer(cols = everything(), names_to = "sampling_mechanism", values_to = "dim") %>%
  ggplot(aes(x = dim, fill = sampling_mechanism)) +
  geom_density(alpha = 0.4, show.legend = F) +
  facet_wrap(~ sampling_mechanism, ncol = 1) +

```

```
theme_minimal()
```

