

# PROBLEM SET 1

## MGMT 737

Spring 2024

You should have gotten this homework assignment from the Github classroom environment (<https://classroom.github.com/classrooms/192971645-yale-mgmt-737-spring-2025-classroom>). In submitting your problem set, you should have two files in your Github repository:

1. `homework1-code.R`, which contains your code,
2. `homework1-writeup.pdf`, which contains your writeup

You may have other files / folders if necessary when there are images or auxiliary files. My very strong preference is for you to write this up in R. If this is not possible, you can use Python. Please let me know if you're planning on this. (This is not a coding preference, but mainly a grading issue.)

1. Randomization. This analysis will use the Dehijia and Wahba sample from the Lalonde dataset of the NSW experiment. The dataset is `lalonde_nsw.csv`. The outcome variable is `re78` (real earnings in 1978). The treatment indicator is `treat`. The remaining variables are potential covariates. Assume for the purposes of this problem set that `treat` is completely randomly assigned.
  - (a) Calculate the average treatment effect of the policy  $E(\tau_i)$  using a simple difference in means.  
→ Label the calculated value as `tau_ate` in your code.
  - (b) Calculate the average treatment effect on the treated of the policy  $E(\tau_i | \text{treat} = 1)$ . How does it compare to part a?  
→ Label the calculated value as `tau_att` in your code.
  - (c) Test the null of  $\tau_i = 0$  for all  $i$  using a randomization test.  
*N.B.* Hold fixed the number of treated and control (e.g. assume the treatment count would be held fixed) and permute the labels randomly 1000 times – you do not need to fully do every permutation (there would be too many). Report the quantile that your estimate from the previous question falls.  
→ Label the quantile value that `tau_att` falls into as `tau_att_quantile` in your code.
  - (d) Run a regression using robust standard errors (you may use canned software) of the outcome on the treatment dummy, and compare the p-values from this test to the previous answer.  
→ Report the difference in the quantile values as `tau_att_quantile_diff` in your code.