Overview

Submit your writeup, including any code, as a PDF via gradescope.¹ We recommend reading through the entire homework beforehand and carefully using functions for testing procedures, plotting, and running experiments. Taking the time to reuse code will help in the long run!

Data science is a collaborative activity. While you may talk with others about the homework, please write up your solutions individually. If you discuss the homework with your peers, please include their names on your submission. Please make sure any handwritten answers are legible, as we may deduct points otherwise.

1. Observational Data on Infant Health

The Infant Health and Development Program (IHDP) was an experiment treating low-birth-weight, premature infants with intensive high-quality childcare from a trained provider. The goal is to estimate the causal effect of this treatment on the child's cognitive test scores. The data *does not* represent a randomized trial with randomly allocated treatment, so there may be confounders between treatment and outcome. In this problem, we devise a propensity score model to control for observed confounders.

- (a) (2 points) The CSV file ihdp.csv has 27 columns:
 - Column 1 is the treatment $z_i \in \{0,1\}$, which indicates whether or not the treatment was given to the infant.
 - Column 2 is the outcome $y_i \in \mathbb{R}$, the child's cognitive test score.
 - Columns 3-27 contain 25 features of the mother and child (e.g. the child's birth weight, whether or not the mother smoked during pregnancy, her age and race). Since this dataset was not collected by a randomized trial, these features could all confound z_i and y_i , and are denoted by $x_i \in \mathbb{R}^{25}$.

In this part, you'll estimate $\hat{e}(x)$ (the predicted probability that $z_i = 1$) by fitting a logistic regression model that predicts z_i from x_i . Specifically:

- 1. Read the data in ihdp.csv (e.g. using the csv package in Python) into three arrays: $Z \in \{0,1\}^n$ containing the treatments, $Y \in \mathbb{R}^n$ containing the outcomes, and $X \in \mathbb{R}^{n \times 25}$ containing the features.
- 2. To fit a logistic regression model, use the scikit-learn package in Python, which is imported as sklearn. Start with the following two lines: from sklearn.linear_model import LogisticRegression as LR lr = LR(penalty='none', max_iter=200, random_state=0)
- 3. Use the lr.fit() method to fit the logistic regression model $\hat{e}(x)$

See the documentation here.

(b) (2 points) Write a function estimate_treatment_effect to estimate treatment accounting for the propensity. It should take as arguments a fitted regression model

 $^{^1\}mathrm{In}$ Jupyter, you can download as PDF or print to save as PDF

(the LogisticRegression object lr from the previous part), X, Y, and Z, and output a single value, which is the estimate of the average treatment effect.

Hint: Use the inverse propensity weighted estimator:

$$\hat{\tau} = \frac{1}{n} \sum_{i=1}^{n} \left(\frac{z_i y_i}{\hat{e}(x_i)} - \frac{(1-z_i) y_i}{1-\hat{e}(x_i)} \right). \tag{1}$$

See the LogisticRegression object's predict_proba method.

- (c) (3 points) Use the function estimate_treatment_effect from the previous part to estimate the treatment effect on the IHDP dataset. Report this estimate. According to the estimate, did the treatment have a beneficial causal effect on the outcome (i.e. cause cognitive test scores to increase)?
- (d) (3 points) The naïve estimator is the difference between the sample means:

$$\tilde{\tau} = \frac{1}{n_1} \sum_{i=1}^{n} y_i z_i - \frac{1}{n_0} \sum_{i=1}^{n} y_i (1 - z_i), \tag{2}$$

where $n_1 = \sum_{i=1}^n z_i$ and $n_0 = n - n_1$. Report this estimate on the IHDP dataset. Why is it different from the estimate you computed in the previous part? Are there any circumstances under which these two estimators should produce the same estimates?

2. Causal Inference Potpourri

A research team wants to estimate the effectiveness of a new veterinary drug for sick seals. They ask aquariums across the country to volunteer their sick seals for the experiment. Since the team offers monetary compensation for volunteering, zoos with less income decide to volunteer their sick seals, whereas zoos with more income are less compelled to volunteer their seals.

It turns out that zoos with less income feed their seals less nutritious diets (regardless of whether they are sick or healthy), due to budgetary constraints. Less nutritious diets prevent seals from recovering as effectively.

- (a) (2 points) **Draw** a causal graph between variables Z, Y, I and N which denote receiving the drug, recovering, the income level of the zoo, and how nutritious a seal's diet is, respectively. Justify each edge in your graph.
- (b) (3 points) We saw in lecture that if we can identify and condition on (adjust for) all confounding variables, then we can use the unconfoundedness assumption to compute the average treatment effect (ATE).

The backdoor criterion provides a way to determine which variables are confounders. In particular, we simply need to "block" all the confounding pathways in the graphical model between Z and Y.

In a causal graph, we define a *path* between two nodes X and Y as a sequence of nodes beginning with X and ending with Y, where each node is connected to the next by an edge (pointed in either direction).

Given an ordered pair of variables (Z, Y), a set of variables S satisfies the backdoor criterion relative to (Z, Y) if no node in S is a descendant of Z (to prevent us from conditioning on colliders), and S blocks every path between Z and Y that contains an arrow into Z.

Using the causal graph in the previous part, determine all possible sets of variables that satisfy the backdoor criterion relative to (Z, Y).

- (c) (3 points) Read the following paper: Safety of the BNT162b2 mRNA Covid-19 Vaccine in a Nationwide Setting and answer the following questions. As when reading any academic paper, you don't have to understand every single thing in the paper as you read: focus on connecting what you're reading to what we learned in class.
 - (i) Does the paper use one of the causal inference techniques we learned in class? If so, which one? Do you have any concerns with how it was applied?
 - (ii) The paper answers two distinct causal questions. For each of the two questions, what is the causal question? What is the treatment? What is the outcome(s)? What are the confounding variables? If any instrumental variables are used, what are they?
 - (iii) Identify at least one potential concern or limitation with this study.

 Hint: when reading any paper, the discussion and/or conclusion sections are often the most insightful.

3. Optimal Mean Estimation via Concentration Inequalities

Suppose we observe a sequence of i.i.d. random variables X_1, \ldots, X_n . Their distribution is unknown, and has unknown mean μ and known variance σ^2 . In this question, we will investigate two different estimators for the mean μ : the sample mean, and the so-called "median of means" estimator. In particular, we will analyze them in terms of how many samples n are required to estimate μ to a given precision ϵ and for a confidence threshold δ

We'll start with the sample mean for parts (a) - (c): in other words, we'll use X_1, \ldots, X_n to compute an estimate $S_n = \frac{1}{n} \sum_i X_i$ for the mean μ . We want to see what sample size n guarantees that $\mathbb{P}(|\hat{\mu} - \mu| \ge \epsilon) \le \delta$.

- (a) (2 points) Let $S_n = \frac{1}{n} \sum_{i=1}^n X_i$. Use Chebyshev's inequality to show that $n = \lceil \frac{\sigma^2}{\delta \epsilon^2} \rceil$ samples are sufficient for $|S_n \mu| \le \epsilon$ with probability at least 1δ .

 Note: $\lceil x \rceil$ is the ceiling function, which rounds x to the nearest larger integer.
- (b) (3 points) Now assume that each X_i is bounded between a and b. Use Hoeffding's inequality to compute the number of samples n sufficient for $|S_n \mu| \leq \epsilon$ with probability at least 1δ . In particular, show that the dependence of n on δ is $O(\log(1/\delta))$.
- (c) (2 points) Suppose we can't assume that each X_i is bounded. In that case, we can no longer apply Hoeffding's inequality, and can only use Chebyshev's inequality. This is a problem if we require a very high confidence level: in this part, you'll show why.

For this part only, assume that $\sigma^2=1$, a=-1,b=1, and $\epsilon=0.1$. Make a plot that shows, for particular values of δ , the number of samples n required based on your answers from parts (a) and (b). Your plot should show a range of ten δ values between 1/2 and 1/1000, using np.geomspace(1/2,1/1000,10), and should be shown on a log-log scale. What do you observe?

To overcome this problem, we'll replace the sample mean with another estimator, and construct bounded random variables that will help us reason about the new estimator. To construct this estimator, we'll start by considering m groups of X_i , each with fixed size n_0 . We'll compute the sample mean for each group, and call these sample means $S^{(1)}, \ldots, S^{(m)}$. Then, we'll use the median of all these group means as our estimate for the mean. The diagram below summarizes our approach.

$$X_1, X_2, \dots X_{n_0}$$
Sample mean $S^{(1)}$
Sample mean $S^{(2)}$
Sample mean $S^{(2)}$
Sample mean $S^{(m)}$

We do this because even though one such sample mean $S^{(i)}$ might be far from the true mean μ , we hope (and will show) that the median of all of them is more likely to be close to the true mean μ .

(d) (2 points) Fix a sample size $n_0 = \lceil \frac{4\sigma^2}{\epsilon^2} \rceil$. For each of the group means i, we define a binary random variable Z_i :

$$Z_i = \mathbb{1}(|S^{(i)} - \mu| \ge \epsilon).$$

In other words, Z_i is 0 if the corresponding group mean is close to the true mean μ (within ϵ), and 1 otherwise.

Show that $\mathbb{E}[Z_i] \leq 1/4$.

Hint: Z_i is a Bernoulli random variable.

(e) (2 points) We set $S_{\text{Med}} := \text{Median}(\{S^{(1)}, \dots, S^{(m)}\})$. This is called the *median-of-means estimator*. Explain in words why having $|S_{Med} - \mu| \ge \epsilon$ implies that $\sum_{i=1}^{m} Z_i \ge \frac{m}{2}$.

Hint: If y is the median of m numbers, it means that $\lceil m/2 \rceil$ of the numbers are greater than or equal to y, and similarly $\lceil m/2 \rceil$ of the numbers are less than or equal to y.

(f) (2 points) By taking probabilities, part (e) implies

$$\mathbb{P}(|S_{\text{Med}} - \mu| \ge \epsilon) \le \mathbb{P}\left(\frac{1}{m} \sum_{i=1}^{m} Z_i \ge \frac{1}{2}\right).$$

If we combine this fact with the result of (d), we can show that

$$\mathbb{P}(|S_{\text{Med}} - \mu| \ge \epsilon) \le \mathbb{P}\left(\frac{1}{m} \sum_{i=1}^{m} (Z_i - \mathbb{E}[Z_i]) \ge \frac{1}{4}\right).$$

Now use Hoeffding's inequality to compute what number m is sufficient to ensure that $|S_{\text{Med}} - \mu| \leq \epsilon$ with probability at least $1 - \delta$. What is the final number of samples of X required?