MS&E 226: "Small" Data

Lecture 19: Additional topics in causal inference (v2)

Ramesh Johari ramesh.johari@stanford.edu

Regression analysis of experiments

Regression analysis of an experiment

Recall using OLS to fit the following model:

$$Y_i \approx \hat{\beta}_0 + \hat{\beta}_1 W_i,$$

where $W_i \in \{0,1\}$ is the assignment in a randomized experiment (0 is control, 1 is treatment), and Y_i is the corresponding observed outcome for individual i.

As we showed:

- $lackbox{}\hat{eta}_0$ is the average outcome in the control group.
- $\hat{eta}_0 + \hat{eta}_1$ is the average outcome in the treatment group.
- $\qquad \qquad \hat{\beta}_1 = \widehat{\mathsf{ATE}}.$

In this lecture we consider what happens when we have additional covariates we can exploit in our analysis.

Suppose in addition to Y(0), Y(1), and W, each individual also has a vector of observed covariates \vec{X} .

In this lecture we consider what happens when we have additional covariates we can exploit in our analysis.

Suppose in addition to Y(0), Y(1), and W, each individual also has a vector of observed covariates \vec{X} .

There are two ways in which the regression approach to experimental analysis is powerful:

In this lecture we consider what happens when we have additional covariates we can exploit in our analysis.

Suppose in addition to Y(0), Y(1), and W, each individual also has a vector of observed covariates \vec{X} .

There are two ways in which the regression approach to experimental analysis is powerful:

Controlling for observed covariates helps improve estimation of the ATE.

In this lecture we consider what happens when we have additional covariates we can exploit in our analysis.

Suppose in addition to Y(0), Y(1), and W, each individual also has a vector of observed covariates \vec{X} .

There are two ways in which the regression approach to experimental analysis is powerful:

- Controlling for observed covariates helps improve estimation of the ATE.
- Interactions with the treatment effect allow us to see how the treatment effect varies among individuals with different covariate vectors.

In this lecture we consider what happens when we have additional covariates we can exploit in our analysis.

Suppose in addition to Y(0), Y(1), and W, each individual also has a vector of observed covariates \vec{X} .

There are two ways in which the regression approach to experimental analysis is powerful:

- Controlling for observed covariates helps improve estimation of the ATE.
- Interactions with the treatment effect allow us to see how the treatment effect varies among individuals with different covariate vectors.

Warning: The covariates \vec{X} must be observed pre-treatment!

Controlling for observables: An example

I created a synthetic experiment where $n_0 = n_1 = 150$.

For each individual i, $X_i \sim \mathcal{N}(0,1)$ is a pre-existing covariate, and W_i is the treatment indicator.

I contructed Y_i as:

$$Y_i = 10 + 0.5 \times W_i + X_i + \varepsilon_i,$$

where $\varepsilon_i \sim \mathcal{N}(0,1)$.

In this example:

- ► The true ATE is 0.5—it does not vary depending on X.
- However, some of the variation in Yi's is explained the X's as well.

Controlling for observables: An example

Suppose we regress Y on the treatment indicator W alone:

Controlling for observables: An example

Now suppose we include the covariate X in the regression:

Notice that the standard error is smaller on the coefficient of W.

Controlling for observables: Interpretation

In the specification Y $\tilde{\ }$ 1 + W + X, we still interpret the coefficient on W as an estimate of the population-level ATE.

The point is that adding X to the regression gives us a better estimate of the baseline Y(0) for each individual.

Essentially, this regression says that for an individual with covariate ${\cal X}$:

- $Y(0) \approx \hat{\beta}_0 + \hat{\beta}_2 X.$
- $Y(1) \approx \hat{\beta}_0 + \hat{\beta}_1 + \hat{\beta}_2 X.$

Controlling for observables

Controlling for observed covariates has another effect as well:

If the randomization was less than perfect, controlling for observed covariates can reduce the sampling bias.

How this works:

▶ Suppose, e.g., individuals with higher *X* were more likely to receive the treatment.

What are the limitations to this process?

Controlling for observables

Controlling for observed covariates has another effect as well:

If the randomization was less than perfect, controlling for observed covariates can reduce the sampling bias.

How this works:

- ▶ Suppose, e.g., individuals with higher *X* were more likely to receive the treatment.
- ▶ Ignoring this fact will lead to a biased estimate of the ATE: part of the variation in the observed Y's is explained by variation in the X's, not by the variation in the treatment. (This is an omitted variable bias.)

What are the limitations to this process?

Controlling for observables

Controlling for observed covariates has another effect as well:

If the randomization was less than perfect, controlling for observed covariates can reduce the sampling bias.

How this works:

- ▶ Suppose, e.g., individuals with higher *X* were more likely to receive the treatment.
- ▶ Ignoring this fact will lead to a biased estimate of the ATE: part of the variation in the observed Y's is explained by variation in the X's, not by the variation in the treatment. (This is an omitted variable bias.)
- Controlling for X removes the omitted variable bias.

What are the limitations to this process?

The preceding slides suggest one limitation of merely controlling for observed covariates:

What if the treatment effect itself varies depending on the covariates observed?

To address this issue we employ interactions with the treatment indicator.

Suppose given a covariate X, we add the interaction term $W\times X$ to the model:

$$Y_i \approx \hat{\beta}_0 + \hat{\beta}_W W_i + \hat{\beta}_X X_i + \hat{\beta}_{WX} W_i X_i.$$

With the addition of this term we can interpret the model as follows:

Suppose given a covariate X, we add the interaction term $W\times X$ to the model:

$$Y_i \approx \hat{\beta}_0 + \hat{\beta}_W W_i + \hat{\beta}_X X_i + \hat{\beta}_{WX} W_i X_i.$$

With the addition of this term we can interpret the model as follows:

Suppose given a covariate X, we add the interaction term $W \times X$ to the model:

$$Y_i \approx \hat{\beta}_0 + \hat{\beta}_W W_i + \hat{\beta}_X X_i + \hat{\beta}_{WX} W_i X_i.$$

With the addition of this term we can interpret the model as follows:

$$Y(0) \approx \hat{\beta}_0 + \hat{\beta}_X X.$$

Suppose given a covariate X, we add the interaction term $W \times X$ to the model:

$$Y_i \approx \hat{\beta}_0 + \hat{\beta}_W W_i + \hat{\beta}_X X_i + \hat{\beta}_{WX} W_i X_i.$$

With the addition of this term we can interpret the model as follows:

- $Y(0) \approx \hat{\beta}_0 + \hat{\beta}_X X.$
- $Y(1) \approx \hat{\beta}_0 + \hat{\beta}_W + (\hat{\beta}_X + \hat{\beta}_{WX})X.$

Suppose given a covariate X, we add the interaction term $W \times X$ to the model:

$$Y_i \approx \hat{\beta}_0 + \hat{\beta}_W W_i + \hat{\beta}_X X_i + \hat{\beta}_{WX} W_i X_i.$$

With the addition of this term we can interpret the model as follows:

- $Y(0) \approx \hat{\beta}_0 + \hat{\beta}_X X.$
- $Y(1) \approx \hat{\beta}_0 + \hat{\beta}_W + (\hat{\beta}_X + \hat{\beta}_{WX})X.$
- ▶ The estimated causal effect is $\approx \hat{\beta}_W + \hat{\beta}_{WX} X$.

Suppose given a covariate X, we add the interaction term $W \times X$ to the model:

$$Y_i \approx \hat{\beta}_0 + \hat{\beta}_W W_i + \hat{\beta}_X X_i + \hat{\beta}_{WX} W_i X_i.$$

With the addition of this term we can interpret the model as follows:

For an individual with covariate X,

- $Y(0) \approx \hat{\beta}_0 + \hat{\beta}_X X.$
- $Y(1) \approx \hat{\beta}_0 + \hat{\beta}_W + (\hat{\beta}_X + \hat{\beta}_{WX})X.$
- ▶ The estimated causal effect is $\approx \hat{\beta}_W + \hat{\beta}_{WX} X$.

This allows us to measure *heterogeneous treatment effects* across the population.

Interactions: Example

In the earlier example, there should be no meaningful change in the treatment effect across individuals with different X's.

```
Call:
lm(formula = Y ~ 1 + W + X + X * W, data = df)
...
```

Coefficients:

```
Estimate Std. Error t value Pr(>|t|)
(Intercept) 9.91161 0.08705 113.860 < 2e-16 ***
W1 0.61730 0.12323 5.009 9.4e-07 ***
X 1.06204 0.09365 11.340 < 2e-16 ***
W1:X -0.09945 0.12986 -0.766 0.444
```

Interactions: Example

Now suppose we change the model so that in the population, changing \boldsymbol{X} also changes the treatment effect.

In particular, suppose:

$$Y_i = 10 + (0.5 + X_i)W_i + X_i + \varepsilon_i,$$

where $\varepsilon_i \sim \mathcal{N}(0,1)$.

What happens when we estimate a model with interactions on the resulting experimental data?

Interactions: Example

```
The result:
Call:
lm(formula = Y ~ 1 + W + X + X * W, data = df)
Residuals:
   Min
         1Q Median
                     3Q
                          Max
-2.32757 -0.73146 0.05078 0.62216 2.85012
Coefficients:
       Estimate Std. Error t value Pr(>|t|)
W1
        X
        1.06476 0.09155 11.630 < 2e-16 ***
W1:X
```

SUTVA and interference

Interference

Implicitly throughout our discussion of causal inference, we have assumed there is no *interference* between treatment and control:

Whether or not individual i receives treatment or control has no impact on the causal effect of treatment on another individual j.

When might this fail?

Interference

Suppose Airbnb decides to A/B test a new feature that dramatically simplifies the booking process for a guest.

In the test, guests are randomized at when they start the booking process; control is the old experience, treatment is the new experience.

It is found that customers with the new experience book much more frequently than customers with the old experience, but the estimated \widehat{ATE} is an *overestimate*. Why?

Interference

Both treatment and control see the same inventory of host listings!

So if treatment individuals book more often, that *reduces* the inventory available to control individuals, and implies their booking rates will be lower.

SUTVA

If interference is present, the "potential outcomes" for an individual are much more complicated: they depend on not just the treatment a single individual received, but also on the treatment *other* individuals received.

With n individuals, this is 2^n potential outcomes for each individual!

SUTVA

If interference is present, the "potential outcomes" for an individual are much more complicated: they depend on not just the treatment a single individual received, but also on the treatment *other* individuals received.

With n individuals, this is 2^n potential outcomes for each individual!

The assumption that there is no interference between treatment and control is part of the *stable unit treatment value assumpton* (SUTVA) in econometrics and causal inference.

The other part of SUTVA is that there is only one form of treatment or control: e.g., if treatment is "taking a drug", there should be no variation in the treatment group as to *how much* of the drug is taken.

Paradoxes

A new treatment for a disease is introduced, and compared against the existing standard of care (control).

Let W=0,1 denote control or treatment, respectively.

Let Y=0,1 denote the outcome disease or no disease, respectively.

Let Z be the gender of the indivdual (M or F).

You run an experiment with a large sample size, and equal numbers of men and women.

	Men		Women	
		Disease	No disease	Disease
	(Y=1)	(Y=0)	(Y=1)	(Y=0)
Treatment ($W=1$)	0.1500	0.2250	0.1000	0.0250
${\sf Control}\;(W=0)$	0.0375	0.0875	0.2625	0.1125

(Here the numbers are the fractions of individuals in each category.)

Analyzing the results:

▶ On average, $\mathbb{P}(Y=1|W=1)=0.5$ while $\mathbb{P}(Y=1|W=0)=0.6$, so the treatment appears detrimental.

Analyzing the results:

- ▶ On average, $\mathbb{P}(Y=1|W=1)=0.5$ while $\mathbb{P}(Y=1|W=0)=0.6$, so the treatment appears detrimental.
- ▶ On the other hand, $\mathbb{P}(Y=1|W=1,Z=M)=0.4$, while $\mathbb{P}(Y=1|W=0,Z=M)=0.3$, so the treatment appears to be beneficial to men.

Analyzing the results:

- ▶ On average, $\mathbb{P}(Y=1|W=1)=0.5$ while $\mathbb{P}(Y=1|W=0)=0.6$, so the treatment appears detrimental.
- ▶ On the other hand, $\mathbb{P}(Y=1|W=1,Z=M)=0.4$, while $\mathbb{P}(Y=1|W=0,Z=M)=0.3$, so the treatment appears to be beneficial to men.
- ▶ In addition, $\mathbb{P}(Y=1|W=1,Z=F)=0.8$, while $\mathbb{P}(Y=1|W=0,Z=F)=0.7$, so the treatment appears to also be beneficial to women as well!

What happened? (This is called Simpson's paradox.)

Each man and woman has two potential outcomes Y(0) and Y(1), associated to control and treatment, respectively.

If we presume there was no sampling bias among men, (so W is uncorrelated with Y given Z=M) then the average causal effect among men is:

$$\begin{split} \mathbb{E}[Y(1) - Y(0)|Z &= M] \\ &= \mathbb{E}[Y(1)|Z = M, W = 1] - \mathbb{E}[Y(0)|Z = M, W = 0] \\ &= \mathbb{P}(Y = 1|Z = M, W = 1) - \mathbb{P}(Y = 1|Z = M, W = 0) \\ &= 0.1 \end{split}$$

Each man and woman has two potential outcomes Y(0) and Y(1), associated to control and treatment, respectively.

If we presume there was no sampling bias among men, (so W is uncorrelated with Y given Z=M) then the average causal effect among men is:

$$\begin{split} \mathbb{E}[Y(1) - Y(0)|Z &= M] \\ &= \mathbb{E}[Y(1)|Z = M, W = 1] - \mathbb{E}[Y(0)|Z = M, W = 0] \\ &= \mathbb{P}(Y = 1|Z = M, W = 1) - \mathbb{P}(Y = 1|Z = M, W = 0) \\ &= 0.1 \end{split}$$

Similarly the average causal effect among women is $\mathbb{E}[Y(1)-Y(0)|Z=F]=0.1.$

So what is the average causal effect overall?

$$\mathbb{E}[Y(1) - Y(0)]$$
= $\mathbb{E}[Y(1) - Y(0)|Z = M]\mathbb{P}(Z = M)$
+ $\mathbb{E}[Y(1) - Y(0)|Z = F]\mathbb{P}(Z = F)$
= 0.1

So there is no paradox: if the causal effect for men and women is separately positive, it must be positive overall.

The issue is that in this example:

$$\mathbb{E}[Y(1) - Y(0)] \neq \mathbb{E}[Y(1)|W = 1] - \mathbb{E}[Y(0)|W = 0].$$

The reason is that if we *ignore* gender, there *is* a sampling bias:

- Women are more likely to be in control than treatment; men are more likely to be in treatment than control.
- ▶ And women have *higher* potential outcomes on average than men: the average outcome of a woman in treatment (resp. control) is 0.8 (resp., 0.7), while the same for a man in treatment is 0.4 (resp., 0.3).
- ► This combination of effects lowers the average outcome in the treatment group relative to the overall population (since the treatment group is primarily men), and raises the average outcome in the control group relative to the overall population (since the control group is primarily women).

The preceding analysis shows that ignoring gender creates an *omitted variable bias* in our estimate of the average treatment effect.

Note that we assumed no further sampling bias beyond gender; the example makes clear that any such bias would only further cloud the true causal effect.

Another example: Berkeley admissions

Berkeley was sued for gender bias in admissions based on 1973 statistics: 44% of men were admitted, while only 35% of women were admitted.

But based on individual departments' admissions statistics, there did not appear to be statistically significant gender-based discrimination (in fact if anything, some departments tended to *favor* women).

What happened is that there was a sampling bias: women were systematically applying to majors that were much more competitive.

The moral

This example is meant to illustrate how to use potential outcomes to carefully describe the causal effect of interest.

Perfect randomization makes up for a lot of deficiencies, but sometimes things are less than perfect.

Taking care to think through potential outcomes and sampling bias carefully can help avoid incorrect inference!

Observational data

Natural experiments

How can we make causal inferences *without* randomized experiments?

As the preceding lecture shows, we need to find other ways to eliminate sampling bias.

The phrase "natural experiment" refers to the fact that we look for structure in the data we are given that "mimics" an experiment we would have wanted to conduct.

Examples

Some examples include:

- Regression discontinuity analysis
- Propensity score matching
- Instrumental variables