Counterfactual Framework of Causal Inference

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Outline

Missing Data

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Random Experiments

Identification

- We have learned how to use samples to estimate and make statistical inference over some population quantity (e.g., P(X) or E(Y|X))
- What if we cannot observe some random variables?
- Statistical identification: use some observed random variables to infer properties about random variables that cannot be observed, or unobserved.
- To address identification problem, we need additional assumptions about our data

Missing Data

- Missing data is one common identification problem
- E.g., in a survey, people answer "Don't know"
- Let us work with the simplest case: we are interested in only one random variable Y.
- And we draw a sample of n points, Y_1, \dots, Y_n from population Y.
- Define R_i be an indicator for whether or not we observe Y_i
- General solutions:
 - 1. Bounds: possible ranges of Y
 - 2. Deletion: discard missing ones
 - 3. Imputation: predict missing Y

Missing data: bounds

- Assume we are interested in E(Y)
- How do we estimate E(Y) in the presence of missing data?
- Suppose we see a data that looks like the below

Unit	Y_{i}	R:
1	1	1
2	?	0
3	1	1
4	0	1
5	1	1
6	?	0

- And we know that Y can take values between [0,1] (Y can be continuous)
- What is the maximum possible value of E(Y)?

Missing data: bounds

• The largest value of Y is 1. We just fill in them, and calculate the largest possible value of E(Y)

Unit	Y_i	R_i
1	1	1
2	1	0
3	1	1
4	0	1
5	1	1
6	1	0

- The largest possible E(Y) is 5/6
- Likewise, we plug in the smallest value of Y
- The smallest possible value of E(Y) is 3/6
- We obtained bounds for E(Y|X): [3/6,5/6]; this is known as Manski bounds.
- Note that bounds are not confidence intervals. WHY?

Missing data: deletion

- Bounds can often be very wide, making them not that useful
- We can make stronger assumption to obtain more meaningful point estimation of E(Y)

Definition (MCAR: Missing Complete at Random, Rubin, 1976)

Y is missing completely at random if:

- 1. The missing $Y \perp \!\!\! \perp \!\!\! \perp R$ (Response is independent of the missing Y we are interested in).
- 2. P(R = 1) > 0 (non-zero response probability)

Missing data: deletion

MCAR assumption implies that

$$E(Y) = E(Y|R=1) \tag{3}$$

- The right hand side is something we can estimate: the sample mean for those we can observe (apply plug-in principle)
- Practical implication: if MCAR holds, we can safely delete missing Y, and E(Y|R=1) is an unbiased estimates of E(Y)

Missing data: imputation

• We can also impute missing values to estimate E(Y)

Unit	Y_i	R_i
1	1	1
2	?	0
2 3 4 5	1	1
4	0	1
5	1	1
6	?	0

Instead of deleting missing rows, we can fill in values

Imputation Method 1: Unconditional Mean Imputation

 Unconditional mean imputation fill in missing Y by the sample mean of observed Y

Unit	Y_i	R_i
1	1	1
2	$\hat{\mathrm{E}}\left[Y R=1\right] = \frac{3}{4}$	0
3	1	1
4	0	1
5	1	1
6	$\hat{\mathrm{E}}\left[Y R=1\right] = \frac{3}{4}$	0

- After unconditional mean imputation, the sample mean of imputed Y is an unbiased estimate of Y
 - Note: this is not the only way to make E(Y) = E(Y|R=1)
- Deletion and imputation all lead to unbiased estimate of E(Y)
- Their variance estimates are usually different!
 - $\hat{V}_{deletion}(Y) = 0.25$
 - $\hat{V}_{imputation}(Y) = 0.15$

MCAR in multivariate case

- When we have multiple variables, we can extend MCAR assumptions: each variable is independent of response.
- And with MCAR assumptions, we can perform listwise deletion by removing any row that has missing entries.

Unit	Y_i	R_i	X_i
1	1	1	0
2	?	0	0
3	1	1	0
4	0	1	0
5	1	1	?
6	?	0	1

 Or taking the imputation perspective, we can perform unconditional mean imputation for each variables

MAR

- MCAR is often too strong in multivariate case
 - If there are many variables, we can delete a lot of observations
 - Often these variable are correlated with each other;
- One weaker assumption is MAR, also known as ignorability

Definition (MAR: Missing at Random, Rubin, 1976)

Y is missing at random if:

- 2. P(R = 1) > 0 (non-zero response probability)
 - That is, Y is missing at random, once we condition on some control variables X.

Post-stratification estimator of sample mean

• Under MAR, we can estimate the mean of *Y* using post-stratification estimator

$$E(Y) = \sum_{x} E(Y|R = 1, X = x)p(X = x)$$
 (5)

- In other words, we estimate E(Y) as the weighted mean of the conditional expectation of Y given X in observed data, with weights P(X=x)
- Both terms on the right hand side can be estimated from samples (plug-in sample analog)
- Note: post-stratification estimator does not impute; directly estimate E(Y)

MAR vs MCAR

• Under MCAR: $\hat{E}[Y_i] = 3/4$

Unit	Y_i	R_i	X_i
1	1	1	0
2	?	0	0
3	1	1	0
4	0	1	0
5	1	1	1
6	?	0	1

• Under MAR, with stratification estimator, $\hat{E}[Y_i] = 7/9$

$$\hat{\mathbf{E}}[Y] = \hat{\mathbf{E}}[Y|R = 1, X = 0]\hat{\mathbf{P}}[X = 0] + \hat{\mathbf{E}}[Y|R = 1, X = 1]\hat{\mathbf{P}}[X = 1]$$
$$= \frac{2}{3} \cdot \frac{4}{6} + 1 \cdot \frac{2}{6} = \frac{7}{6}$$

- MCAR and MAR will yield different estimates of E(Y)
- Each estimate is unbiased estimate only if the corresponding assumption is true

Imputation method 2: Conditional Mean Imputation

- With MAR, we can also impute Y using conditional mean imputation: use the conditional mean of Y as our guesses of the missing Y
- $Y_i = \hat{E}(Y|R=1, X=X_i)$

Unit	Y_i	R_i	X_i
1	1	1	0
2	$\hat{\mathbf{E}}\left[Y_i X_i=0\right]=\tfrac{2}{3}$	0	0
3	1	1	0
4	0	1	0
5	1	1	1
6	$\hat{\mathrm{E}}\left[Y_{i} X_{i}=1\right]=1$	0	1

- Then we can calculate sample mean over imputed Y
- Under conditional mean imputation, $\hat{E}(Y)$ is again 7/9
- The below two gives the same estimate of E(Y):
 - conditional mean imputation of Y, and then take sample mean of imputed Y
 - post-stratification estimator

Conditional Mean Imputation using linear regression

- If we further assume all assumptions of linear regression are correct: E(Y|R=1,X=x) is linear in X
- Then conditional mean imputation just uses predicted values of linear regression as imputed values

Conditional Mean Imputation using regression

Unit	Y_i	R_i	$X_{[1]i}$	$X_{[2]i}$	
1	1	1	0	3	
2	$\hat{\beta}_0 + \hat{\beta}_1 \cdot 0 + \hat{\beta}_2 \cdot 7$	0	0	7	
3	1	1	0	9	(8)
4	0	1	0	5	
5	1	1	1	4	
6	$\hat{\beta}_0 + \hat{\beta}_1 \cdot 1 + \hat{\beta}_2 \cdot 3$	0	1	3	

Conditional Mean Imputation using other methods

- The interpretation advantage of linear regression is not relevant now; we do not care about interpreting β ; we want our predictions of E(Y|R=1,X=x) to be more precise
- So you can use GLM to predict E(Y|R=1,X=x)
 - GLM
- Or other more complex machine learning algorithms. It's a prediction problem!
- These options are all provided in R package mice

Imputation Method 3: hot-deck imputation

- Hot-deck imputation uses nearest-neighbor matching
- For unit i with missing Y_i , and non-missing X_i
 - Find the X_j that has the smallest distance to/is closet X_i
 - Use the Y_j associated with j as the imputed Y value for i

Unit	Y_i	R_i	X_i
1	1	1	4
2	?	0	8
3	1	1	1
4	0	1	12
5	1	1	20
6	?	0	3

• Example: unit 6's X is closest to unit 1's X. So we impute Y_6 as $Y_1=1$

Hot-deck imputation using propensity scores

- When we have multivariate X, it is not easy to calculate their distances
- Instead, it is popular to estimate propensity score of response

$$P(R=1|X) \tag{9}$$

- Propensity score of response provides an single-number summary of multivariate X
- Hot-deck imputation based on nearest propensity score, not based on original distances between X
 - In other words, you want to match units whose response propensity are similar
- Estimation of propensity scores
 - Logistic regression is the default choice
 - But apparently other machine learning methods are acceptable

Hot-deck example

Unit	Y_i	R_i	$X_{[1]i}$	$X_{[2]i}$	$P(R_i = 1 X_i)$	
1	2	1	0	3	0.33	
2	?	0	0	7	0.14	
3	3	1	0	9	0.73	(10)
4	10	1	0	5	0.35	
5	12	1	1	4	0.78	
6	?	0	1	3	0.70	

• Unit 6's propensity score of response is closest to unit 3's propensity score. Thus Y_6 is imputed as $Y_3 = 3$

Deletion vs Imputation

- In practice, assume we want to run a regression based on Y and 10 predictors X
- Solution 1: Listwise deletion
 - Both R and Stata uses this strategy by default
 - Pros: simple; unbiased if MAR is true
 - Cons: large standard errors (since you will drop many cases)
- Solution 2: mean imputation (unconditional or conditional)
 - Pros:
 - give you more cases to work with
 - also unbiased if MCAR/MAR is true
 - Cons: small standard errors. Why?
 - Artificially fix the missing Y to its mean.
- Solution 3: hot-deck imputation
 - Pros: preserve the support of original data
 - Bear similarity to propensity score matching
 - Cons: how to estimate propensity scores?

Stochastic Imputation

- Problem of mean imputation: small standard error issues when we use sample mean for imputation
- A workaround—stochastic imputation—add some random noise to the sample mean
 - Say, we still use regression to impute Y, but add some random noises to your predicted Y
 - If we are working with complex machine learning models, there
 may be some inherent stochastic component (results are not
 the same every time)
- Problem: stochastic imputation also have some uncertainty, based on what noises you use
 - These random noises are not added to your final analysis, thus still producing small standard error estimates

Multiple Imputation

- Rubin, 1977, Multiple Imputation
 - Stochastic imputation for m times; ending up with m imputed datasets.
 - Analysis: Run your model (regression Y on 10 X) m times
 - Pooling: parameter estimates for m different models can be used for estimation and inference:
 - The final parameter estimates of β is the mean of β across m models
 - The standard error of final β is more complex in math
 - basically it's the (within model standard error of β) + (between model standard error of β)
 - Or use bootstrap if *m* is large enough

Missing data by Chained Equations

- In practice, more than one X can have missing values
- Assume we have 5 X; we use 1, 2, 3, 4 to impute the 5th variable, and then use 1, 2, 3, 5 to impute the 4th variable, and so on and so forth
 - Imputed values are allowed to use in the next step

Prediction vs Causation

- Correlation \neq causation
- We can use X to predict Y, and use Y to predict X
- $Y = g(X) \iff X = g^{-1}(Y)$
- This does not capture the intuitive idea that X causes Y



https://www.tylervigen.com/spurious-correlations

Counterfactual

- Does college education lead to higher wages?
- Observed (Factorial): on average, college graduates indeed earn more than people with only high school education
- Critique:
 - people who can go to college have higher ability
 - even if they did not go to education, they could still earn more
 - Therefore, correlation does not mean causation
- Counterfactual thinking:
 - Guess the (counterfactual) earning of college graduates if they did not go to college
 - If the counterfactual earning equals to the factual earning, then college education does not matter; there is no causal effect
 - Alternatively, if the factual earning is higher than counterfactual earning, then college education indeed lead to increase in wages

Neyman-Rubin Causal Model: potential outcomes

- Neyman-Rubin Causal Model formally write down the counterfactual idea
- We have a binary treatment D; D = 1 if treated and 0 otherwise
- For a person i in the population, her outcome Y_i is assumed to be:

Definition (Neyman-Rubin model)

$$Y_{i} = \begin{cases} Y_{i}^{0} : D_{i} = 0 \\ Y_{i}^{1} : D_{i} = 1 \end{cases}$$
$$= Y_{i}^{0} + D_{i}(Y_{i}^{1} - Y_{i}^{0})$$

- Y_i is observed outcome
- Y_i^0 is the potential outcome if i is not treated
- Y_i^1 is the potential outcome if i is treated

Individual Level Treatment Effect

Definition (Unit-Level Treatment Effect)

For i in the population, the causal effect of treatment for unit i is : $\rho_i = Y_i^1 - Y_i^0$

- $Y_i = Y_i^0 + D_i(Y_i^1 Y_i^0)$
- So the Neyman Rubin model suggests that the observed Y for a treated unit i are the two sums:
 - counterfactual outcome if i were not treated
 - individual level treatment effect

(12)

Counterfactual Outcome as Missing Data Problem

Y_i	D_i
1	1
1	0
1	1
0	1
1	1
0	0
	1 1 0

Unit	Y_i^0	Y_i^1	D_i
1	?	1	1
2	1	?	0
3	?	1	1
4	?	0	1
5	?	1	1
6	0	?	0

Fundamental Problem of Causal Inference

Definition (Fundamental Problem of Causal Inference, Holland, 1986)

At any given time, we only observe one of the potential outcomes for unit i — either Y_i^1 or Y_i^0 —but not both. Thus unit-level treatment effect ρ_i is not identified.

- Similar to missing data problems, we have to make assumptions (here, assumptions about potential outcomes) to allow identification.
 - In particular, identification of the average of treatment effect ρ_i because identifying the effect for every unit can be extremely hard

ATE and ATT

Definition (Average Treatment Effect (ATE))

$$ATE = E(\rho) = E(Y^1 - Y^0) = E(Y^1) - E(Y^0)$$

- ATE is the mean of unit-level treatment effect
- ATE is the difference between the mean of two potential outcomes

Definition (Average Treatment Effect on the Teated (ATT))
$$ATT = E(\rho|D=1) = E(Y^1 - Y^0|D=1) = E(Y^1|D=1) - E(Y^0|D=1)$$

ATT is the mean of unit-level treatment effect for treated units

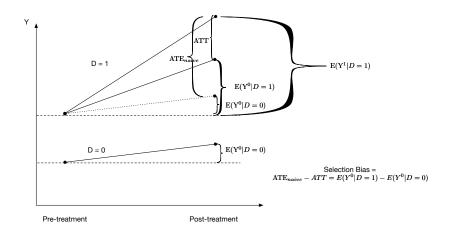
Naive estimate of ATE

 Naive estimate of ATE is just the difference in means of observed data

$$ATE_{naive} = E[Y|D=1] - E[Y|D=0]$$
 (13)

- ullet For instance, D=1 for college education and D=0 for less than college education
- ATE_{naive} is the mean earning of college education mean earning of non-college educated
- In general, $ATE \neq ATT \neq ATE_{naive}$; what is their connection?

Selection Bias



Selection bias

- ATE_{naive} neither estimates ATE nor ATT
- ATE_{naive} differs from ATT by selection bias

selection bias =
$$ATE_{naive} - ATT = E(Y^0|D=1) - E(Y^0|D=0)$$

- Intuitively, this is the counterfactual earning of college educated if they did not go to college, minus the factual earning of non-college educated
 - This selection bias could be caused by ability, for example
- If selection bias is 0, $ATE_{naive} = ATT$

Random Assignment

 In randomized controlled experiments, we randomly assign subjects into treatment and control groups; we have random assignment

Definition ((Completely) Random Assignment)

- $Y_i^0, Y_i^1 \perp \!\!\! \perp D_i$ (Potential outcome is independent of treatment assignment)
- P(D=1) > 0 (non-zero treatment probability)

- Cautions:
 - Observed outcome Y is not independent of treatment assignment.

Random Assignment Solves the Identification Problem

• Under random assignment of *D*, we have:

$$ATE_{naive} = E[Y|D=1] - E[Y|D=0] = ATE$$
 (14)

• Proof (the first line to second line is due to independence between D and Y^0, Y^1)

$$E[Y|D = 1] - E[Y|D = 0] = E[Y^{1}|D = 1] - E[Y^{0}|D = 0]$$

$$= E[Y^{1}|D = 1] - E[Y^{0}|D = 1]$$

$$= E[Y^{1} - Y^{0}|D = 1]$$

$$= E[Y^{1} - Y^{0}]$$

$$= E[Y^{1}] - E[Y^{0}]$$
(15)

Non-parametric estimator: difference-in-means

- With random assignment, estimating ATE is very simple: ATE_{naive}, which is just the difference in mean outcome of the treatment and the control group
- This is a non-parametric estimator
- Another important observation: ATT = ATE for randomized experiments

Experiment as Imputation

Unit	Y_i^0	Y_i^1	D_i
1	?	1	1
2	1	?	0
3	?	1	1
4	?	0	1
5	?	1	1
6	0	?	0

- Random assignment implies that we can impute the missing values using observed sample mean; similar to the MCAR assumption in missing data
 - But here, random assignment is a fact, not an assumption

Unit	Y^0	Y^1	D	
1	$\hat{\mathbf{E}}\left[Y D=0\right]=\tfrac{1}{2}$	1	1	
2	1	$\hat{\mathrm{E}}\left[Y D=1\right]=\tfrac{3}{4}$	0	
3	$\hat{\mathbf{E}}\left[Y D=0\right]=\tfrac{1}{2}$	1	1	(17)
4	$\hat{\mathbf{E}}\left[Y D=0\right]=\frac{1}{2}$	0	1	
5	$\hat{\mathbf{E}}\left[Y D=0\right]=\frac{1}{2}$	1	1	
6	0	$\hat{\mathrm{E}}\left[Y D=1\right]=\tfrac{3}{4}$	0	

Regression estimator of ATE

• We can rewrite Y_i in the following way (MHE, 2.3.1)

$$Y_{i} = E\left(Y_{i}^{0}\right) + \left(Y_{i}^{1} - Y_{i}^{0}\right) D_{i} + Y_{i}^{0} - E\left(Y_{i}^{0}\right)$$

$$= \alpha + \rho_{i} D_{i} + \eta_{i}$$
(18)

- This equation looks like linear regression! But each individual has its own regression coefficient ρ_i , which is the individual-level treatment effect
- Constant treatment assumption: assume that ρ_i is the same for every one, ρ , $ATE = E(\rho_i) = \rho$

$$E[Y_i|D_i = 1] = \alpha + \rho + E[\eta_i|D_i = 1]$$

$$E[Y_i|D_i = 0] = \alpha + E[\eta_i|D_i = 0]$$
(19)

$$ATE_{naive} = E[Y_i|D_i = 1] - E[Y_i|D_i = 0]$$

$$= \underbrace{\rho}_{ATE} + \underbrace{E[\eta_i|D_i = 1] - E[\eta_i|D_i = 0]}_{\text{selection bias}} (20)$$

Regression estimator of ATE

- Selection bias is 0, since $Y^0 \perp \!\!\! \perp D$ under random assignment
- Therefore,

$$ATE_{naive} = E[Y_i|D_i = 1] - E[Y_i|D_i = 0] = \underbrace{\rho}_{ATE}$$

- Therefore, if you are running a random experiment,
 - Non-parametric estimator: the difference in mean outcome of treated and control units
 - Parametric estimator:
 - assume constant treatment effect
 - run a regression of observed outcome on treatment D, and use coefficient of D as the estimate of ATE

Regression as Imputation

 The regression estimator of ATE is implicitly making counterfactual imputation using linear regression:

Unit	Y_i^0	Y_i^1	D_i	$X_{[1]i}$	$X_{[2]i}$
1	?	2	1	1	7
2	5	?	0	8	2
3	?	3	1	9	3
4	?	10	1	3	1
5	?	2	1	5	2
6	0	?	0	7	0

Regression as Imputation

• Fit a regression $Y = \beta_0 + \beta_1 D_i + \beta_2 X_{[1]i} + \beta_3 X_{[2]i}$, and impute counterfactual outcome using the linear regression:

Unit	Y_i^0	Y_i^1	D_i	$X_{[1]i}$	$X_{[2]i}$
1	$\hat{\beta}_0 + \hat{\beta}_1 \cdot 0 + \hat{\beta}_2 \cdot 1 + \hat{\beta}_3 \cdot 7$	2	1	1	7
2	5	$\hat{\beta}_0 + \hat{\beta}_1 \cdot 1 + \hat{\beta}_2 \cdot 8 + \hat{\beta}_3 \cdot 2$	0	8	2
3	$\hat{\beta}_0 + \hat{\beta}_1 \cdot 0 + \hat{\beta}_2 \cdot 9 + \hat{\beta}_3 \cdot 3$	3	1	9	3
4	$\hat{\beta}_0 + \hat{\beta}_1 \cdot 0 + \hat{\beta}_2 \cdot 3 + \hat{\beta}_3 \cdot 1$	10	1	3	1
5	$\hat{\beta}_0 + \hat{\beta}_1 \cdot 0 + \hat{\beta}_2 \cdot 5 + \hat{\beta}_3 \cdot 2$	2	1	5	2
6	0	$\hat{\beta}_0 + \hat{\beta}_1 \cdot 1 + \hat{\beta}_2 \cdot 7 + \hat{\beta}_3 \cdot 0$	0	7	0
					(21)

• Then ATE and ATT can be easily calculated as the difference in means of Y^1 and Y^0

Inference: Neyman Variance Estimator

- If we are running experiments, ATE can be estimated easily by taking the differences between E(Y|D=1) and E(Y|D=0)
- What about statistical inference?
- Neyman Variance Estimator
 - 1. assume constant treatment effect
 - 2. Then

$$V(ATE) = \frac{V_t}{N_t} + \frac{V_c}{N_c}$$

- 3. V_t is variance of Y for treated users, and N_t is number of treated users
- If treatment effect is not constant, true variance is usually smaller than $\frac{V_t}{N_t} + \frac{V_c}{N_c}$

Inference: Linear regression

• If we regress Y on D ($Y = \alpha + \rho D$), it can be shown that the regression estimates of the standard error of regression coefficient is exactly same as the Neyman Variance estimator

$$V(\rho) = V(ATE) = \frac{V_t}{N_t} + \frac{V_c}{N_c}$$

See Imbens and Rubin (chapter 7) for proof

Inference: Randomization Test

- Null distribution: D has no causal effect on Y
 - then if we shuffle the outcome, E(Y|D=1) E(Y|D=0) = 0
- Randomization test
 - Calculate ATE based on experimental data
 - Shuffle your observed Y, and recalculate ATE_{shuffle} based on the shuffled data
 - Say you shuffled 1000 times, and have 1000 ATE_{shuffle}.
 - Then you can easily calculate 95% confidence interval/standard errors of ATE estimates
 - The p value for observing ATE is just the probability that your shuffled $ATE_{shuffle}$ is larger than ATE: p-value = $P(ATE_{shuffle} > ATE)$
- Pros: do not need to assume constant treatment effect
- Cons: time consuming

- Researchers often collect some additional covariates (i.e., pre-treatment variables)
- With additional variables, it is easier to work with regression estimator

$$Y_i = \alpha + \rho D_i + \beta X_i + \epsilon_i \tag{22}$$

- $\hat{\rho}^{adj}$:covariate-adjusted estimate of treatment effect
- $\hat{\rho}$: difference-in-means of outcome variables across treatment and control (or regression coefficient by regressing Y on D without covariates)
- $\hat{\rho}_X$: difference-in-means of X across treatment and control

• It can be shown that (Li and Ding, 2019, J. R. Stat. Soc, or Imbens and Rubins, Chapter 7):

$$\hat{\rho}^{adj} = \hat{\rho} - \hat{\beta}^T \hat{\rho}_X$$

- $\hat{\rho}_X$: difference-in-means of X across treatment and control
 - · With completely randomized experiments
 - $\hat{\rho}^{adj}$ is biased; $\hat{\rho}$ is unbiased
 - $\hat{\rho}_X$ are usually not exactly 0, especially when the data size is not that large
 - Both are consistent
 - because with more and more data, $\hat{\rho}_X$ approaches 0; this is called covariate balance
- Be careful if your treatment and control groups are not balanced; in that case, the treatment effect estimates without and with covariates can differ a lot in finite sample

- Another classical justification to add covariates in regression is that $\hat{
 ho}^{adj}$ has smaller standard error than $\hat{
 ho}$
- For instance, MHE (p. 23): "Inclusion of the variable X... generate more precise estimates of the causal effect)"
- David A. Freedman, On regression adjustments to experimental data, Advances in Applied Mathematics 40 (2008), no. 2, 180–193
- It is not necessarily true!

Winston Lin, Agnostic notes on regression adjustments to experimental data: Reexamining Freedman's critique, The Annals of Applied Statistics **7** (2013), no. 1, 295–318. MR3086420

- adding covariate is guaranteed to lead to smaller standard error estimates, if
- 1. full interaction is added; and
- 2. robust standard errors are used

$$Y_i = \alpha + \rho D_i + \beta X_i + \frac{\gamma D_i X_i}{\epsilon_i} + \epsilon_i$$

 Note that condition 1 is not easy to follow in practice; if you have 10 covariates, you have to add 10 interaction terms

Recommended practice

- David A. Freedman, On regression adjustments to experimental data, Advances in Applied Mathematics 40 (2008), no. 2, 180–193
- Always present two treatment effects: without and with covariates
- "Regression estimates...should be deferred until rates and averages have been presented"
- Always check pre-treatment covariate balance
- Add interactions if covariance-balance is passed
- not only guarantees smaller standard error, but also detects treatment effect heterogeneity (next week)