STA 640 — Causal Inference

Chapter 2.1. Randomized Experiments: Fisher's and Neyman's Mode of Inference

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Role of Randomization

- ► In randomized experiments, assignment mechanism is known and controlled by investigators
- Randomization does:
 - ▶ balance observed covariates: $Z \perp \!\!\! \perp \!\!\! \perp X$
 - ▶ balance unobserved covariates: $Z \perp \!\!\! \perp \!\!\! U$
 - balance potential outcomes, i.e. guarantee ignorability or unconfoundedness (Rubin 1978)

$$Z \perp \!\!\! \perp (Y(1), Y(0))$$

▶ Note on terminology: the terms *covariates*, *pretreatment variables*, *baseline characteristics* are used exchangeably in the literature, referring to *X*

Role of Randomization

Under randomization, causal effects are (nonparametrically)
 identified, because we can show

$$Pr(Y(z)) = Pr(Y|Z = z), z = 0, 1$$

Under randomization, association does imply causation (of course within the potential outcome framework with assumptions), for example:

$$\tau^{\text{ATE}} \equiv \mathbb{E}[Y_i(1) - Y_i(0)] = \mathbb{E}[Y|Z=1] - \mathbb{E}[Y|Z=0]$$

 Randomization ensures model-free validity for model-based statistics (explain later)

Types of Randomized Experiments

- 1. Bernoulli trials
- 2. Completely randomized experiment
- 3. Stratified (or block) randomized experiment
- 4. Paired randomized experiment
- 5. Cluster randomized experiment

Bernoulli Trials

- 1. *Bernoulli trials*: for each unit we flip a fair coin to determine their assignment
 - ► The assignment mechanism is

$$p(\mathbf{Z}|\mathbf{X}, \mathbf{Y}(0), \mathbf{Y}(1)) = \left(\frac{1}{2}\right)^{N}$$

- ► Here $p(Z_i = 1 | \mathbf{X}, \mathbf{Y}(0), \mathbf{Y}(1)) = \frac{1}{2}$
- ▶ The number of possible assignments is 2^N

Completely Randomized Experiment

- 2. Completely randomized experiment: N_1 out of N units are randomly chosen to receive the treatment ($N_0 = N N_1$) is the no. of control)
 - ► The assignment mechanism is

$$p(\mathbf{Z} \mid \mathbf{X}, \mathbf{Y}(0), \mathbf{Y}(1)) = \begin{cases} \begin{pmatrix} N_1 \\ N \end{pmatrix}^{-1} \left(\frac{N_1}{N}\right)^{N_1} \left(\frac{N_0}{N}\right)^{N_0} & \text{if } \sum_{i=1}^{N} Z_i = N_1 \\ 0 & \text{otherwise.} \end{cases}$$

- ► Here $p(Z_i = 1 | \mathbf{X}, \mathbf{Y}(0), \mathbf{Y}(1)) = \frac{N_1}{N}$
- Most often $N_1 = N/2$, so that half the units receive the active treatment and half receive the control treatment
- ► The number of possible assignments is $\begin{pmatrix} N \\ N_1 \end{pmatrix}$

Stratified (block) Randomized Experiment

- 3. *Stratified or block randomized experiment*: First partition into blocks or strata, similar in some sense (e.g. covariates), and then within each block, perform a completely randomized trial
 - Assignments are independent across blocks
 - Example: block 1 women; block 2- men
 - "Block what you can; randomize what you cannot" George Box
- 4. Paired randomized experiments: a special case of stratified trials, with 2 units within each block. Total number of possible assignments $2^{N/2}$.
- Clustered randomized experiments: treatment assigned at cluster level, all units in one unit get the same treatment (requires additional notations, more later)

Fisher's Randomization Inference

- ► Fundamental idea: inference based *solely on the assignment*mechanism
- ► *Y*(0), *Y*(1) are all fixed, randomness only comes from the assignment of *Z*
- ► RA Fisher was the first to grasp the importance of randomization for credibly assessing causal effects (1925, 1935)
- Given data from such a randomized setting, Fisher was intent on testing sharp null hypotheses

Fisher's Exact Test of Sharp Null Hypothesis

- Null Hypothesis (H_0): An initial supposition regarding the nature of the treatment effect, usually specified to test its consistency against observed data
- ► Sharp Null Hypothesis: A null hypothesis articulated with specificity sufficient to allow the researcher to fill in a hypothetical value for each unit's missing potential outcome
- ► Test Statistic ($T = T(\mathbf{Y}(1), \mathbf{Y}(0), \mathbf{Z})$): A function of the data that the researcher uses to determine the consistency of the null hypothesis with observed data

Fisher's Exact Test: Three Steps

- 1. Specify the null hypothesis
- 2. Choose the statistic
- 3. Measure the extremeness by the p-value (under the randomization distribution)

Fisher's Exact Test of Sharp Null: Example

Imbens and Rubin (2009): A small subset of data from a randomized job training experiment — the Greater Avenues to INdependence (GAIN) programs. Earnings in First Year Post-randomization:

Individual	Potentia	al Outcomes	Actual	Observed Outcome	
	$Y_i(0)$	$Y_i(1)$	Treatment	Y_i	
1	66	?	0	66	
2	0	?	0	0	
3	0	?	0	0	
4	?	0	1	0	
5	?	607	1	607	
6	?	436	1	436	

Fisher's Exact Test: Example

We first test the sharp null hypothesis that the program had absolutely no effect on reading scores:

$$H_0: Y_i(0) = Y_i(1)$$
 for all $i = 1, ..., 6$

▶ Under H_0 , the unobserved potential outcomes are equal to the observed outcomes for each unit

Individual	Potentia	l Outcomes	Actual	Observed Outcome	
	$Y_i(0)$	$Y_i(1)$	Treatment	Y_i	
1	66	(66)	0	66	
2	0	(0)	0	0	
3	0	(0)	0	0	
4	(0)	0	1	0	
5	(607)	607	1	607	
6	(436)	436	1	436	

Fisher's Exact Test: Example

- Consider testing the null hypothesis against the alternative hypothesis that $Y_i(0) \neq Y_i(1)$ for some units
- ➤ We use the difference in average outcomes by treatment status as test statistic:

$$T = T(\mathbf{Z}, Y^{obs}) = \frac{1}{3} \sum_{i=1}^{6} Z_i \cdot Y_i^{obs} - \frac{1}{3} \sum_{i=1}^{6} (1 - Z_i) \cdot Y_i^{obs}$$

- ► For the observed data the value of the test statistic is 325.6
- ▶ Under the null hypothesis, we can calculate the value of this statistic under each vector of treatment assignments **Z** (20 different assignment vectors)

Z_1	Z_2	Z_3	Z_4	Z_5	Z_6	level
0	0	0	1	1	1	325.
0	0	1	0	1	1	325.
0	0	1	1	0	1	-79.0
0	0	1	1	1	0	35.0
0	1	0	0	1	1	325.
0	1	0	1	0	1	-79.0
0	1	0	1	1	0	35.0
0	1	1	0	0	1	-79.0
0	1	1	0	1	0	35.0
0	1	1	1	0	0	369.
1	0	0	0	1	1	369.
1	0	0	1	0	1	-35.0
1	0	0	1	1	0	79.0
1	0	1	0	0	1	-35.0
1	0	1	0	1	0	79.0
1	0	1	1	0	0	325.
1	1	0	0	0	1	-35.0
1	1	0	0	1	0	79.0
1	1	0	1	0	0	325.
1	1	1	0	0	0	325.

Fisher's Exact Test: Example

- ► How likely is (under the randomization distribution) to observe a value of the test statistic that is as large in absolute value as the one actually observed?
- **Exact p-value:** $Pr(T \ge T^{obs}) = 8/20 = 0.40$ for our sharp null hypothesis, our test statistic, and our definition of unusual
- ► In this example, no difference between one-way and two-way tests. But usually matters
- Under the null hypothesis of no effect of the program the observed difference could therefore well be due to chance

Fisher's Exact Test: Definition

- ► Randomization Distribution: All possible values of the test statistic and the probabilities associated with each such value under the (randomized) assignment mechanism, assuming the null hypothesis to be true
- p-value: Under the assumption that the null hypothesis is correct, the probability of observing a value of the test statistic as extreme or more extreme than the one actually observed
- Fisher Interval: The set of possible values of the causal quantity of interest corresponding to test statistics with p-values that fall within some range set by the researcher; usually interpreted to be a set of plausible values of the average causal effect

Fisher's Exact Test: Choice of Statistic

t statistic

$$T = \frac{1}{n_1} \sum_{i=1}^{n} Z_i Y_i - \frac{1}{n_0} \sum_{i=1}^{n} (1 - Z_i) Y_i$$

ightharpoonup Wilcoxon statistic based on ranks R_i 's of Y_i 's

$$W = \frac{1}{n_1} \sum_{i=1}^{n} Z_i R_i - \frac{1}{n_0} \sum_{i=1}^{n} (1 - Z_i) R_i$$

 Special case with binary outcome: equivalent to Fisher's exact test

Covariate adjustment in randomization test

- ► What if there is chance imbalance in baseline (pretreatment) covariates?
- ► Regression adjustment:
 - \triangleright OLS (or other model) fitting of the Y_i 's on pretreatment covariates
 - obtain residuals ϵ_i 's
- Key facts under the sharp null $Y_i(1) = Y_i(0)$ for all i
 - ightharpoonup values of Y_i 's and X_i 's are fixed
 - ▶ values of ϵ_i 's are fixed
- ▶ Replace Y_i by ϵ_i in the test statistic
- Or replace R_i by the rank of ϵ_i
- Validity of randomization test does not rely on the linear model assumption (Rosenbaum 2002 Stat Sci)

Pros and cons of randomization test

► Pros

- exact, no need for asymptotics
- works for any test statistic
- test statistics = any functions of data (fitted parameters of any models)
- can deal with heavy-tailed data

Cons

- exact inference is attractive, but sharp null seems dull (and restrictive). Extended to less restrictive null, i.e.
 - $H_0: Y(1) = Y(0) + \tau$ by Ding et al. (2016), but still dull and restrictive
- computationally intensive, especially in large experiments
- often less powerful than methods mentioned later
- no guidance for choice of test statistic (power?)
- confidence interval is hard unless constant causal effects

Neyman's Repeated Sampling Approach

- ► Focus on methods for estimating *average treatment effects*
- ► Neyman's two questions:
 - ▶ What would the average outcome be if all units were exposed to the treatment?
 - ► How did that compare to the average outcome if all units were exposed to the control?
- Neyman's procedure (classic frequentist's view):
 - 1. Find an *unbiased estimator* of the ATE
 - 2. Obtain variance and interval estimates of the unbiased estimator (derive the estimator's distribution under *repeated sampling* based on the randomization distribution of the assignment vector *Z*)

Neyman's Repeated Sampling: Basic concepts

- **Estimand**: average treatment effects (ATE).
- ► Note: in randomized experiments, ATE is the same as ATT or ATC (average treatment effect for the control). Why?
- Unbiased estimator of the estimand: A statistic whose average value over all possible randomizations equals the true treatment effect
- Confidence (Neyman) Interval: An interval which, over all possible randomizations, includes the true value of the quantity of interest at least as often as advertised. For example, a 95% confidence interval includes the true value 95% of the time or more
- ► Variance: find the *sampling* variance of the unbiased estimator of the effect and an unbiased estimator of this variance

Estimand: Population Average Treatment Effect (PATE)

► Population Average Treatment Effect (PATE):

$$\tau^{\text{PATE}} \equiv \mathbb{E}[Y_i(1) - Y_i(0)]$$

► The sample under study is a simple random sample from a super population with sample size $\tilde{N} \gg N_0 + N_1$

Estimand: Sample Average Treatment Effect (SATE)

► Another estimand is Sample Average Treatment Effect (SATE)

$$\tau_{\text{SATE}} \equiv \frac{1}{N} \sum_{i=1}^{N} \{Y_i(1) - Y_i(0)\} = \bar{Y}(1) - \bar{Y}(0)$$

- ► SATE focuses on the sample (size $N = N_0 + N_1$) under study
- ► SATE is often of interest in randomized trials, but rarely in observational studies
- ► Subtle theoretical difference between PATE and SATE, see Imbens and Rubin (2015, Chapter 6), but little difference in practice
- ► We will mostly focus on PATE (referred to simply as ATE)

PATE: Unbiased Estimator

► An unbiased estimator of PATE is the difference-in-means between groups, also known as the unadjusted estimator:

$$\hat{\tau}^{\text{unadj}} = \sum_{i=1}^{N} \left(\frac{Z_i \cdot Y_i}{N_1} \right) - \sum_{i=1}^{N} \left(\frac{(1 - Z_i) \cdot Y_i}{N_0} \right) = \bar{Y}_1 - \bar{Y}_0$$

 Unbiasedness is with respect to all possible randomization assignments Z and sampling

$$\mathbb{E}[\hat{\tau}^{\text{unadj}}|Y(0),Y(1)] = \tau^{\text{PATE}}$$

PATE: Variance of Unadjusted Estimator

ightharpoonup Variance of the unadjusted estimator $\hat{\tau}^{\text{unadj}}$

$$\mathbb{V}[\hat{\tau}^{\text{unadj}}] = \frac{S_0^2}{N_0} + \frac{S_1^2}{N_1} - \frac{S_{01}^2}{\tilde{N}}$$

where \tilde{N} is the size of the super population, and

$$S_z^2 = \frac{1}{N-1} \sum_{i=1}^N (Y_i(z) - \bar{Y}(z))^2 \quad z = 0, 1$$

$$S_{01}^2 = \frac{1}{N-1} \sum_{i=1}^{N} (Y_i(1) - Y_i(0) - (\bar{Y}(1) - \bar{Y}(0)))^2$$

This estimator of the variance is unbiased if the super population size \tilde{N} is infinitely large

(Sample) Estimator of Variance

- ▶ Because we never observe both $Y_i(1)$ and $Y_i(0)$, S_{01} is not estimable from data
- ► So, as Neyman suggested, usually one just ignores S_{01} in estimating variance of ATE:

$$\mathbb{V}^{\text{Neyman}} = \frac{S_0^2}{N_0} + \frac{S_1^2}{N_1}$$

▶ A unbiased estimator of the Neyman variance $\mathbb{V}^{\text{Neyman}}$ is to replace S_z by its sample version

$$\hat{\mathbb{V}}^{\text{Neyman}} = \frac{s_0^2}{N_0} + \frac{s_1^2}{N_1}$$

where

$$s_z^2 = \frac{1}{N_z - 1} \sum_{i: Z = 1} (Y_i - \bar{Y}_z)^2, \quad z = 0, 1$$

Confidence Interval

- ▶ Usually based on a normal approximation to the randomization (and sampling) distribution of the estimate $\hat{\tau}^{\text{unadj}}/\mathbb{V}_{\text{Neyman}}$
- ▶ A standard confidence interval of level $(1 \alpha) \times 100\%$ is

$$CI^{1-\alpha}(\hat{\tau}) = \left(\hat{\tau}^{\text{unadj}} - z_{1-\alpha/2} \cdot \sqrt{\hat{\mathbb{V}}}, \hat{\tau}^{\text{unadj}} + z_{1-\alpha/2} \cdot \sqrt{\hat{\mathbb{V}}}\right)$$

► Can be improved by a t-distribution

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