### **Section 2**

#### **Randomization Inference**

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**GOV 2003** 

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#### **Overview**

- Logistics:
  - Pset 2 released! Due at 11:59 pm (ET) on Sept 22
- Today's topics:
  - 1. Randomization inference (Design-based inference)
  - 2. Toy example: Donations encouragement (small/large sample)
  - 3. Inverting test to obtain CIs

#### Randomization inference

- Randomization inference (Design-based inference; permutation test)
  - Assignment mechanism:  $\rightarrow \Omega_0 = \{ \boldsymbol{d} : \mathbb{P}(\boldsymbol{D} = \boldsymbol{d}) > 0 \}.$ 
    - Bernoulli randomization → use rbinom(N, 1,.5)
    - Completely randomized experiment → use ri::genperms() or sample()
  - Sharp null hypothesis:  $H_0: \tau_i = Y_i(1) Y_i(0) = const. \forall i$ 
    - We can fill out the missing potential outcomes
  - We can compute/approximate the distribution of test statistics
     T(D, Y) under the null (randomization distribution)

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    - → We can fill out the missing potential outcomes
  - $\rightsquigarrow$  We can compute/approximate the distribution of test statistics  $T(\mathbf{D}, \mathbf{Y})$  under the null (randomization distribution)
- Model-based inference
  - Assumes a distribution for potential outcomes

### **Toy example: Donations encouragement**

- Setup:
  - N people
  - Encouragement by mail  $(0/1; D_i) \rightarrow \text{Donations to Harvard } (\$; Y_i)$
  - Let  $\Omega$  = set of  $2^N$  treatment vectors (any N-vector of 0s and 1s).  $\Omega_0 \subset \Omega$

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# Suppose **complete randomization** has been implemented N = 6 and $n_1 = \sum_{i=1}^{N} D_i = 3$ .

- $\sim \Omega_0 = \{\mathbf{d} \in \Omega : \sum_{i=1}^6 d_i = 3\} = \{(1,1,1,0,0,0), (1,1,0,1,0,0), \ldots\}$

$$\Omega_0 = id \cdot P(D=d) > 0$$

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  - $\rightarrow \Omega_0 = \{ \mathbf{d} \in \Omega : \sum_{i=1}^6 d_i = 3 \} = \{ (1, 1, 1, 0, 0, 0), (1, 1, 0, 1, 0, 0), \ldots \}$
- Test a sharp null of no effect:  $H_0: \tau_i = Y_i(1) Y_i(0) = 0 \quad \forall i$ .

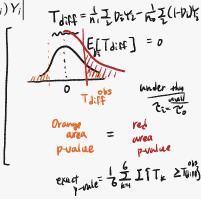
	Mailer	Contr.		
Unit	$D_i$	$Y_i$	$Y_i(0) =$	$Y_i(1)$
Jon	1	3	(3)	3
Sansa	1	5	(5)	5
Arya	1	0	(0)	0
Robb	0	4	4	(4)
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Rickon	0	1	1	(1)

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  - E.g.: absolute difference-in-means estimator → Lost

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- 2. Calculate observed test statistic:  $T^{obs} = T(\mathbf{D}, \mathbf{Y})$ .
- 3. List all the possible treatment vectors in  $\Omega_0$ :  $\{\widetilde{\mathbf{D}}_1,\ldots,\widetilde{\mathbf{D}}_K\}$  where  $K=|\Omega_0|$

$$= \binom{6}{3} = 20$$

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- 5. Observe the distribution of  $\widetilde{T} = {\widetilde{T}_1, \dots, \widetilde{T}_K}$ .
- 6. Calculate the exact p-value:  $p = \frac{1}{K} \sum_{k=1}^{K} \mathbb{I}(\widetilde{T}_k \geq T)$

#### 1. Calculate observed test statistic

	Mailer	Contr.		
Unit	$D_i$	$Y_i$	$Y_i(0)$	$Y_i(1)$
Jon	1	3	(3)	3
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$$T_{\rm diff}^{\rm obs} = |8/3 - 5/3| = 1$$

```
y \leftarrow c(3, 5, 0, 4, 0, 1)
D \leftarrow c(1, 1, 1, 0, 0, 0)
T_{obs} \leftarrow abs(mean(y[D == 1]) - mean(y[D == 0]))
T_{obs} \leftarrow cobs
(y,y) cobs y cobs y
## [1] 1
```

#### 2. Randomization distribution

 $\mathcal{V}^{o}$ 

- Possible treatment assignments  $\{\widetilde{\mathbf{D}}_1, \dots, \widetilde{\mathbf{D}}_{20}\}$
- Test statistics under the null  $\widetilde{T} = \{\widetilde{T}_1(\widetilde{\mathbf{D}}_1, \mathbf{Y}), \dots, \widetilde{T}_{20}(\widetilde{\mathbf{D}}_{20}, \mathbf{Y})\}$

	Mailer	Contr.		
Unit	$\widetilde{D}_1$	$Y_i$	$Y_i(0)$	$Y_i(1)$
Jon	1	3	(3)	3
Sansa	1	5	(5)	5
Arya	0	0	(0)	0
Robb	1	4	4	(4)
Bran	0	0	0	(0)
Rickon	0	1	1	(1)

$$\widetilde{T}_1 = |12/3 - 1/3| = 3.67$$

#### 2. Randomization distribution

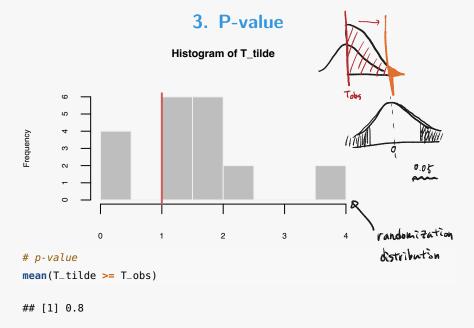
• Possible treatment assignments  $\{\widetilde{\mathbf{D}}_1, \dots, \widetilde{\mathbf{D}}_{20}\}$ • Test statistics under the null  $\widetilde{T} = \{3.67, \dots, \widetilde{T}_{20}(\widetilde{\mathbf{D}}_{20}, \mathbf{Y})\}$ 

	Mailer	Contr.		
Unit	$\widetilde{D}_{20}$	$Y_i$	$Y_i(0)$	$Y_i(1)$
Jon	0	3	(3)	3
Sansa	0	5	(5)	5
Arya	0	0	(0)	0
Robb	1	4	4	(4)
Bran	1	0	0	(0)
Rickon	1	1	1	(1)

$$\widetilde{T}_{20} = |5/3 - 8/3| = 1$$

#### 2. Randomization distribution

```
D_bold <- ri::genperms(D)
K <- ncol(D_bold)
T_tilde <- rep(NA, times = K)
for (i in 1:K) {
    D_tilde <- D_bold[, i]
    T_tilde[i] <- abs(mean(y[D_tilde == 1]) - mean(y[D_tilde == 0]))
}</pre>
```



In a large sample,

(omplete rand

- 1. Choose a sharp null hypothesis and a test statistic:
- 2. Calculate observed test statistic:  $T^{\text{obs}} = T(\mathbf{D}, \mathbf{Y})$ .
- 3. Too many possible treatment vectors in  $\Omega_0 \to \text{take } K = 1000$  samples!
- 4. Calculate  $\widetilde{T}_k = T(\widetilde{\mathbf{D}}_k, \mathbf{Y})$  for each k under the sharp null.
- 5. Observe the distribution of  $\widetilde{T} = {\widetilde{T}_1, \dots, \widetilde{T}_K}$ .
- 6. Calculate the p-value:  $p = \frac{1}{K} \sum_{k=1}^{K} \mathbb{I}(\widetilde{T}_k \geq T)$

### **Choosing test statistics**

- Difference in means
- Rank statistic
  - when we have many outliers
  - → wilcox.test() for rank-sum statstic
- $S = \sum_{i=1}^{N} D_i Y_i(1)$ 
  - when  $Y_i$  is binary, Fisher's exact test (recall Lady Tasting Tea)
  - → fisher.test()
- Using absolute values under the sharp null of no effect
  - > testing against a two-sided alternative hypothesis

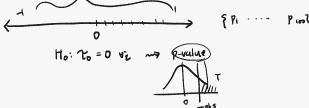
$$H_0: \tau_i = 0 \ \forall i$$
  $H_1: \tau_i \neq 0 \ \text{for some } i$ 

### **Confidence Intervals: Inverting the Test**

/const

- For a sharp null  $\tau_i = Y_i(1) Y_i(0) = \tau_0$   $\forall i$ , we can conduct the test and calculate the p-value.
- Repeat the above with different values of  $\tau_0$

• 95% CI: The range of  $\tau_0$  that we cannot reject the null at the 0.05 level.



# **Example Code for Inverting the Test**

```
# Data
 Yi <- large_sample<mark>$</mark>factor # Observed outcome
Di <- recode(large_sample$canvass, `Placebo` = 0, `Full Intervention` = 1)
N \leftarrow length(Yi); n1 \leftarrow sum(Di); n0 \leftarrow sum(1-Di)
 # Pick candiate taus on a grid
                                                 tau_{cand} = c(-0.5, -\cdots, 0.5)
 tau_cand <- seq(-0.5, 0.5, by = 0.01)
save_pval <- rep(NA, length(tau_cand)) # to save the p-value below</pre>
# 1. Calculate the observed statistics
T_obs <- sum(Di*Yi)/n1 - sum((1-Di)*Yi)/n0</pre>
```

```
Example Code for Inverting the Test pin - 2-4
T000 2: Create function for computing p-value given tau and observed star
TODO 2-1: Calculate Yi(1) using Yi, Di, and tau
   <- NULL
                                                 Yzu)- Yzlo) = 0,1
  TODO 2-2: Calculate Yi(0) using Yi, Ti, and tau
Y0)<- NULL
                                        null
Ttilde_ls <- rep(NA, n_sim)</pre>
# Simulation:
for (s) in 1:n_sim) {
                                                        (0.6)
                                    7: ~ No
  # TODO 2-3: Randomly sample treatment vectors
  Dtilde_s <- NUI
  # TODO 2-4: For each treatment vector,
                                     compute
  Ttilde_ls[s] <- NULL
                                             Y2(0) =0.2
                               h:M
# 2-5: Calculate and return the p-value
-pval <- 2 * min(mean(Ttilde_ls >= t_obs), mean(Ttilde_ls <= t_bbs
return(pval)
                                                               15
```

### **Example Code for Inverting the Test**

```
# TODO 3: Loop over each candidate tau
set.seed(123)
for (t in 1:length(tau_cand)) {
  save_pval[t] <- your_fun(tau_t, T_obs)</pre>
}
# 4. Obtain the upper / lower bound of 95% CI
lb <- tau_cand[min(which(save_pval >= 0.025))]
ub <- tau_cand[max(which(save_pval >= 0.025))]
# TODO 5: Print the 95% CI (lb. ub)
```

# Visualization of p-values from test inversion

