

# MACHINE LEARNING & PUBLIC POLICY

## LECTURE V: MACHINE LEARNING FOR CAUSAL INFERENCE

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# PREDICTION FOR POLICY

**Observation 1:** Sometimes correlation is valuable on its own

# ANOMALY DETECTION

**Observation II:** Sometimes goal is one of detection or discovery

# CAUSAL INFERENCE

## SETUP AND NOTATION

- We follow the Neyman-Rubin Causal Model
- We have an *iid* sample  $\mathcal{N}$  of units  $(1, \dots, N)$  from a population  $\mathcal{P}$
- Every unit  $i$  is described by the tuple  $(Y_i(0), Y_i(1), X_i, W_i)$ 
  - $(Y_i(0), Y_i(1))$  are the set of potential outcomes under (binary) treatment conditions (e.g., test scores)
  - $X_i$  is the vector of pre-treatment covariates (e.g., demographics)
  - $W_i$  is the (binary) treatment indicator (e.g., if received tutoring or not)
  - $Y_i^{obs} = Y_i(W_i)$  (e.g., the test score we observe)

## CHALLENGES TO CASUAL INFERENCE

- Estimands of interest are?

- $\tau_{ATE} := \mathbb{E}[Y_i(1) - Y_i(0)]$

- What would be a good (unbiased and consistent) estimator?

$$\tau_{ATE} := \frac{1}{N} \sum Y_i(1) - \frac{1}{N} \sum Y_i(0)$$

- Is this feasible? What estimator this feasible?

$$\hat{\tau}_{ATE} := \frac{1}{N_T} \sum_{W_i=1} Y_i^{obs} - \frac{1}{N_C} \sum_{W_i=0} Y_i^{obs}$$

- What is the problem with what is feasible?

- Selection:  $(Y_i(0), Y_i(1)) \perp\!\!\!\perp W_i$
  - Students who have educated parents, may be more likely to have higher test scores, and are very motivated so they select to get tutoring
  - Athletes may be more likely to have lower test scores, and are forced to get tutoring

## ARE WE AT AN IMPASSE?

- We need additional assumptions, e.g., selection on  $X_i$  (unconfoundedness)
  - $(Y_i(0), Y_i(1)) \perp\!\!\!\perp W_i \mid X_i$
- Students who have educated parents, may be more likely to have higher test scores, and are very motivated so they select to get tutoring
  - If  $X_i$  contains the educational level of parents we are ok
- Athletes may be more likely to have lower test scores, and are forced to get tutoring
  - If  $X_i$  contains an indicator of the student if an athlete we are ok
- Also,  $X_i$  can contain information highly predictive of  $P(W_i)$
- Randomized Experiments are the gold standard, if possible, because
  - $(Y_i(0), Y_i(1)) \perp\!\!\!\perp W_i$
  - $\hat{\tau}_{ATE} := \frac{1}{N_T} \sum_{W_i=1} Y_i^{obs} - \frac{1}{N_C} \sum_{W_i=0} Y_i^{obs}$  therefore, unbiased & consistent estimate of  $\tau_{ATE}$

## TREATMENT EFFECT HETEROGENEITY





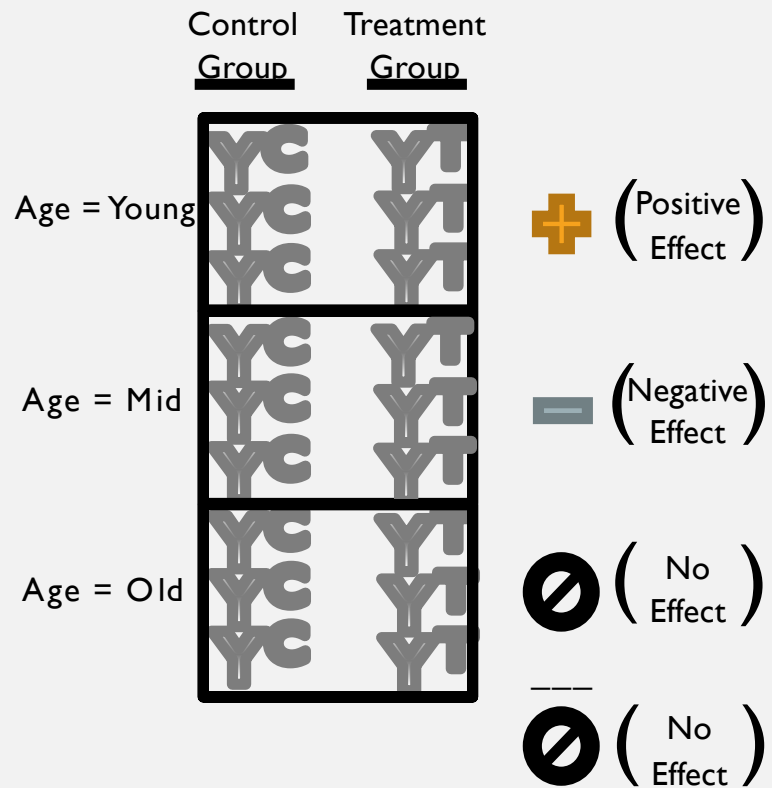
## TREATMENT EFFECT HETEROGENEITY



# TREATMENT EFFECT HETEROGENEITY

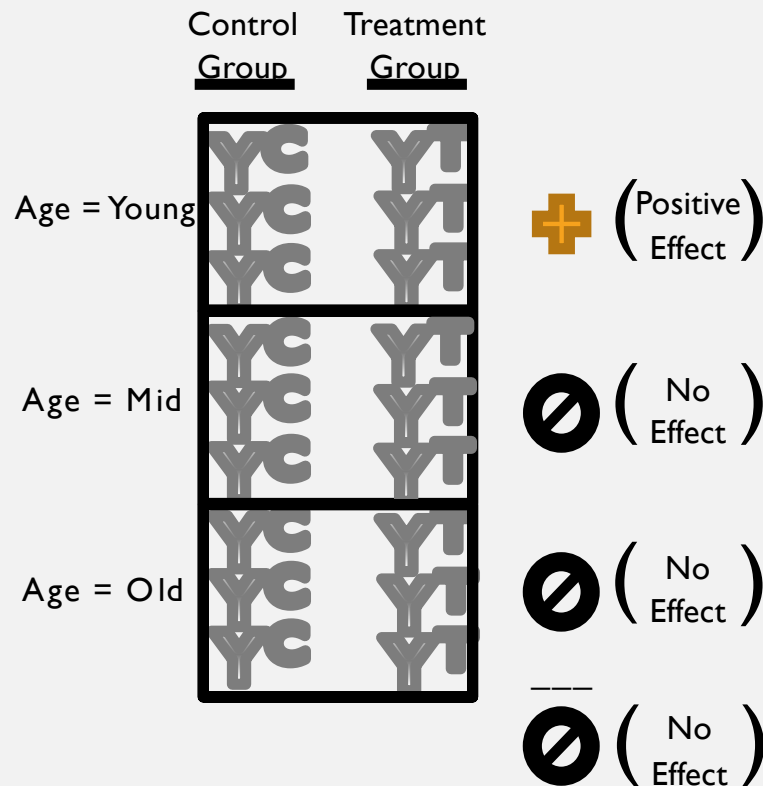
	<u>Control</u> <u>Group</u>	<u>Treatment</u> <u>Group</u>
Age = Young	YC YC YC	YT YT YT
Age = Mid	YC YC YC	YT YT YT
Age = Old	YC YC YC	YT YT YT

# TREATMENT EFFECT HETEROGENEITY



- Positive and negative effects can cancel

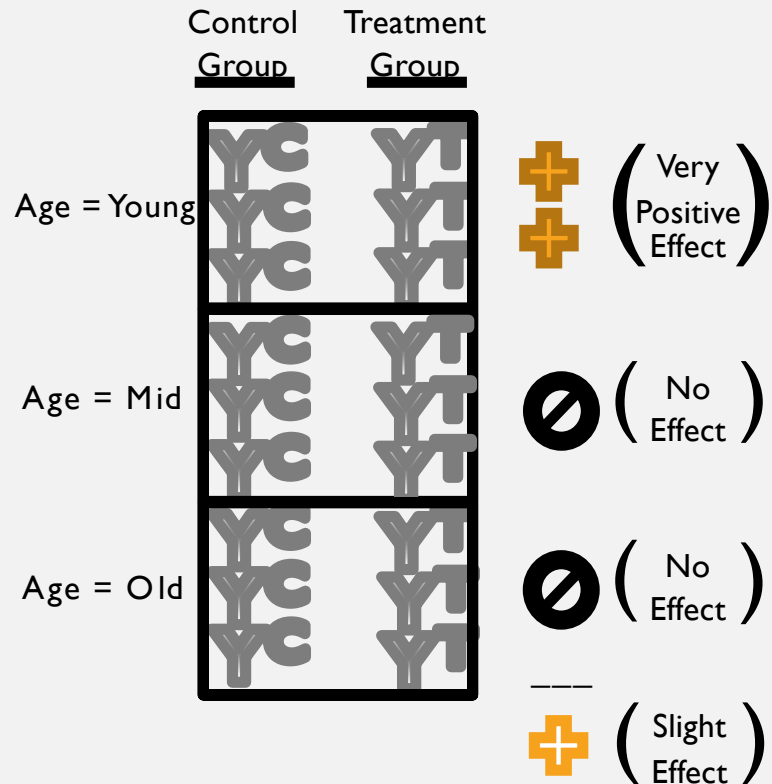
# TREATMENT EFFECT HETEROGENEITY



- Positive and negative effects can cancel

- True effect can be masked

# TREATMENT EFFECT HETEROGENEITY

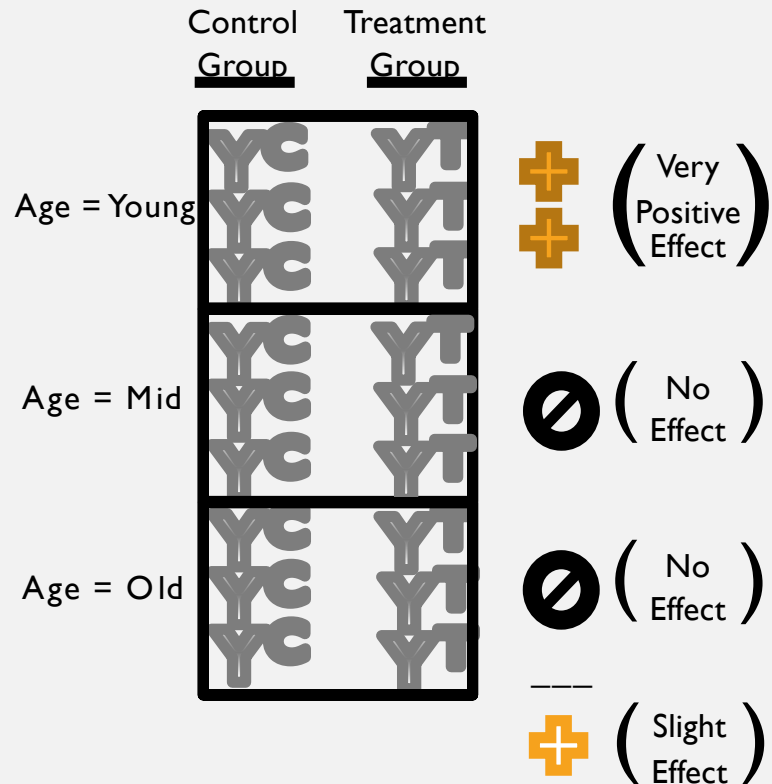


- Positive and negative effects can cancel

- True effect can be masked

- Effects could really be driven by a subpopulation

# TREATMENT EFFECT HETEROGENEITY



- Positive and negative effects can cancel
- True effect can be masked
  - Ex: FDA Approved BiDil Drug
- Effects could really be driven by a subpopulation
  - Ex: Perry Preschool

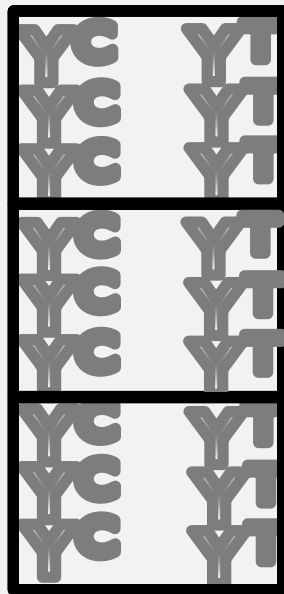
# TREATMENT EFFECT HETEROGENEITY

	<u>Control</u> <u>Group</u>	<u>Treatment</u> <u>Group</u>
Age = Young	YC YC YC	YT YT YT
Age = Mid	YC YC YC	YT YT YT
Age = Old	YC YC YC	YT YT YT

- We need richer estimands of interest
  - $\tau_{CATE}(x) := \mathbb{E}[Y_i(1) - Y_i(0) | X_i = x]$
- With a good (unbiased and consistent) estimator
  - $\hat{\tau}_{CATE}(x) := \frac{1}{N_{T,x}} \sum_{X_i=x} W_i=1 Y_i^{obs} - \frac{1}{N_{C,x}} \sum_{X_i=x} W_i=0 Y_i^{obs}$

# TREATMENT EFFECT HETEROGENEITY

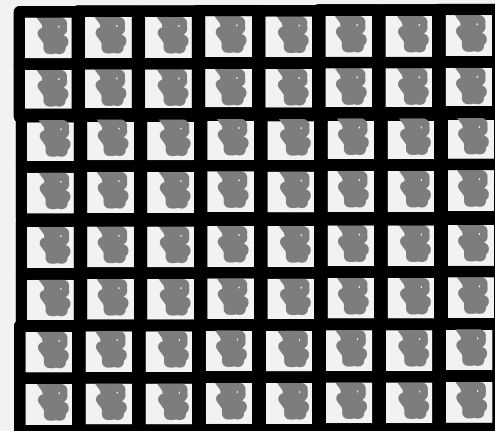
Control  
Group



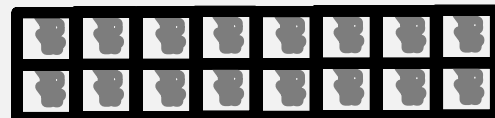
Treatment  
Group



Google  
facebook



...





# ML FOR HTE

**Observation III:** Sometimes correlation can be (forced into) causation

# MACHINE LEARNING'S CONTRIBUTIONS

- Regression Methods
  - OLS and Regularized Regression (e.g., LASSO)
  - Imai and Ratkovic (2013)
- Single Tree Methods
  - Athey and Imbens (2017)
- Ensemble Methods
  - Wager and Athey (2017)
- Anomalous Pattern Detection
  - McFowland et al. (2018)

## (SPARSE) REGRESSION METHOD

- OLS

$$y = \gamma_0 + \gamma_1 X_1 + \gamma_2 X_2 + \dots + \beta_1 W + \beta_2 X_1 W + \beta_3 X_1^2 W + \dots + \varepsilon$$

- Difficult to estimate, so formulate as a problem of variable selection

- LASSO:

$$\min_{\alpha} \|y - X\alpha\|_2^2 + \lambda \|\alpha\|_1 \text{ for } \alpha = (\beta, \gamma)$$

- The goals of  $\beta$  and  $\gamma$  are fundamentally different

- $\beta$  meant to capture the (heterogeneity in) treatment effect that exist
- $\gamma$  meant to explain how the outcome in the control condition behaves

- Imai and Ratkovic (2013) (R-package: FindIt) I

$$\min_{\alpha=(\beta, \gamma)} \|y - X\alpha\|_2^2 + \lambda_{\beta} \|\beta\|_1 + \lambda_{\gamma} \|\gamma\|_1$$

- See Belloni et al, 2014 for how to perform inference after sparse selection
- Issues with Regression?
  - Have to pre-specify the model (and the dimensions of heterogeneity)

## TREE METHODS

- Trees offer a flexible means to estimate  $Y_i = f(X_i) + \epsilon_i$ 
  - Where  $Y_i$  is an outcome of interest and  $X_i$  are features along which  $Y_i$  can vary
- What is the outcome of interest in HTE?

$$\begin{aligned}\tau_i &= Y_i(1) - Y_i(0) \\ &= f(X_i)\end{aligned}$$

- Cannot learn a tree directly because  $\tau_i$  is not observed for any unit in the data
- Recall

$$\begin{aligned}\tau_{CATE}(x) &= \mathbb{E}[Y_i(1) - Y_i(0) | X_i = x] \\ &= \mu(1, x) - \mu(0, x)\end{aligned}$$

$$\mu(w, x) = \mathbb{E}[Y_i^{obs} | W_i = w, X_i = x]$$

## TREE METHODS

$$\tau_{CATE}(x) = \mu(1, x) - \mu(0, x)$$

$$\mu(w, x) = \mathbb{E}[Y_i^{obs} | W_i = w, X_i = x]$$

- Approach I: Use 1 tree to estimate  $\hat{\mu}(w, x)$ , including  $W$  as feature
- Approach II: Use 2 trees to estimate  $\hat{\mu}(1, x)$  and  $\hat{\mu}(0, x)$  separately

$$\hat{\tau}_{CATE}(x) = \hat{\mu}(1, x) - \hat{\mu}(0, x)$$

- $Y_i^{obs} \perp\!\!\!\perp W_i$ , therefore  $\hat{\mu}(w, x)$  is unbiased & consistent (for A.I and A.II)
- Künzel et al. (2018) explains when each Approach is better
  - Also proposes a way to combine them to combine their strengths and avoid their weaknesses.
- Issues?
  - Neither approach is actually optimizing for fitting  $\hat{\tau}_{CATE}(x)$
  - Tree may never actually split on  $W$  (A.I) and lose interpretable subpopulations (A.II)

## TREE METHODS

- Athey and Imbens (2017) proposes Causal Tree
  - Directly optimize tree splitting criteria to minimize  $MSE(\hat{\tau}, \tau)$ 
    - Proposes using separate data when splitting for “Honest Estimation”
  - Key assumption:  $\mathbb{E}_{Test}[\tau_i | X_i] = \mathbb{E}_{Test}[\hat{\tau}(x_i) | X_i]$
  - Provides valid asymptotic confidence intervals
- Benefits of Trees?
  - Does not require pre-specification
- Issues with Trees?
  - Greedy, Unstable, Piece-wise constant function estimation

## ENSEMBLE METHODS

- Many methods propose to simply use ensemble methods, instead of trees, for Approaches I and II
  - Ex: Bayesian Additive Regression Trees (Hill, 2012), Combining super learners (Grimmer, 2018)
  - Same issues still remain: not actually optimizing for fitting  $\hat{\tau}_{CATE}(x)$
- Wager and Athey (2017) propose CausalForest
  - Uses random forest for honest treatment effect estimation for individual units
  - Directly optimize tree splitting criteria to minimize  $MSE(\hat{\tau}, \tau)$
  - Proves asymptotic normality of random forest, providing valid confidence intervals
- Benefits?
  - Stable and smooth function estimation
- Issues?
  - No natural subpopulations or groups.

## LIMITATIONS

- Regression Methods
  - Pre-specification of the model
- Single Tree Methods
  - Greedy and unstable
- Ensemble Methods
  - Fairly uninterpretable/no natural subpopulations
- General Limitations
  - The mean and only the mean
    - Other moments can be effected
    - Simpsons Paradox
  - Risk minimization not effect maximization
    - Small number of subpopulations considered
    - No guarantee on their “interestingness”
  - No “discovery”, only model inspection

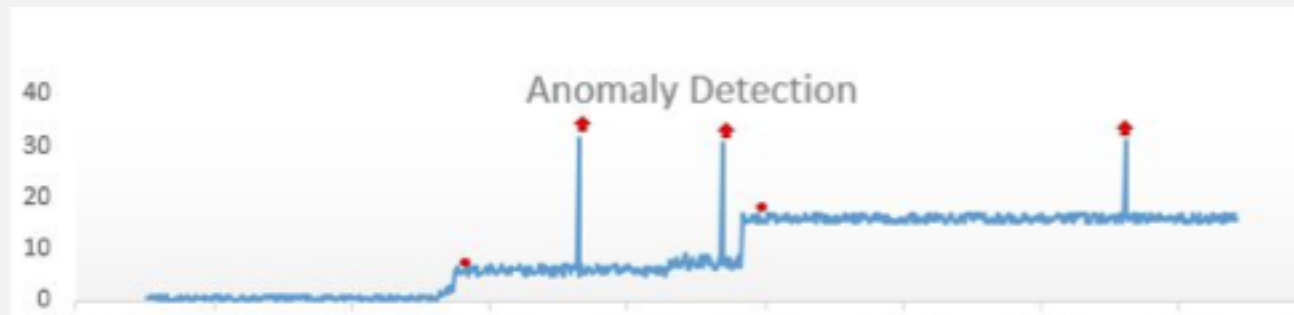


# ANOMALY DETECTION FOR HTE

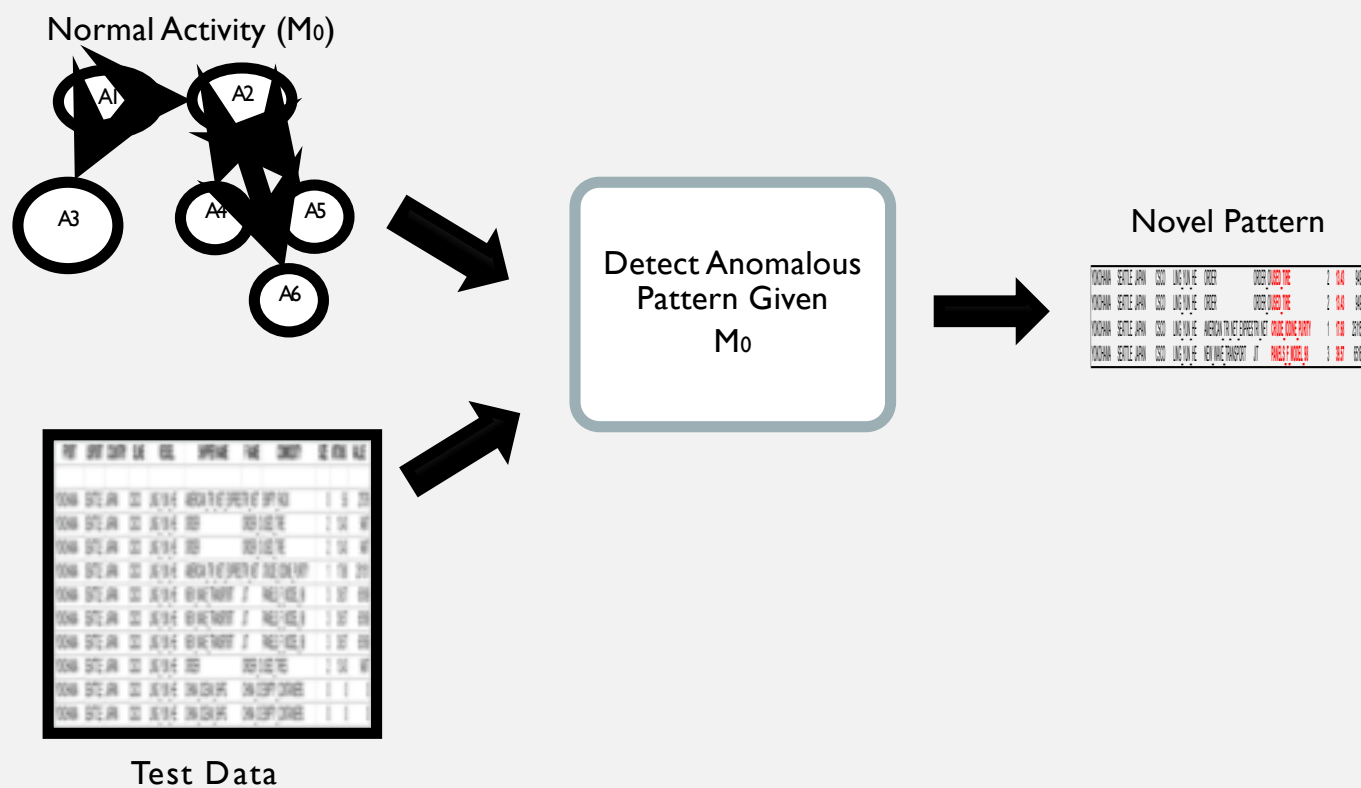
**Observation III:** Sometimes correlation can be (forced into) causation

# ANOMALY DETECTION PARADIGM

- Identifying when a “system” deviates away from its expected behavior.



# Anomalous Pattern Detection Procedure



# HTE Pattern Detection

## Control Group

Item	Item No	Qty	Unit	Desc	Price	Quantity
1001	1	1000	EA	WASHING MACHINE	1000	1000
1002	1	1000	EA	WASHING MACHINE	1000	1000
1003	1	1000	EA	WASHING MACHINE	1000	1000
1004	1	1000	EA	WASHING MACHINE	1000	1000
1005	1	1000	EA	WASHING MACHINE	1000	1000
1006	1	1000	EA	WASHING MACHINE	1000	1000
1007	1	1000	EA	WASHING MACHINE	1000	1000
1008	1	1000	EA	WASHING MACHINE	1000	1000
1009	1	1000	EA	WASHING MACHINE	1000	1000
1010	1	1000	EA	WASHING MACHINE	1000	1000
1011	1	1000	EA	WASHING MACHINE	1000	1000
1012	1	1000	EA	WASHING MACHINE	1000	1000
1013	1	1000	EA	WASHING MACHINE	1000	1000
1014	1	1000	EA	WASHING MACHINE	1000	1000
1015	1	1000	EA	WASHING MACHINE	1000	1000
1016	1	1000	EA	WASHING MACHINE	1000	1000
1017	1	1000	EA	WASHING MACHINE	1000	1000
1018	1	1000	EA	WASHING MACHINE	1000	1000
1019	1	1000	EA	WASHING MACHINE	1000	1000
1020	1	1000	EA	WASHING MACHINE	1000	1000

## Treatment Group

Item	Item No	Qty	Unit	Desc	Price	Quantity
1001	1	1000	EA	WASHING MACHINE	1000	1000
1002	1	1000	EA	WASHING MACHINE	1000	1000
1003	1	1000	EA	WASHING MACHINE	1000	1000
1004	1	1000	EA	WASHING MACHINE	1000	1000
1005	1	1000	EA	WASHING MACHINE	1000	1000
1006	1	1000	EA	WASHING MACHINE	1000	1000
1007	1	1000	EA	WASHING MACHINE	1000	1000
1008	1	1000	EA	WASHING MACHINE	1000	1000
1009	1	1000	EA	WASHING MACHINE	1000	1000
1010	1	1000	EA	WASHING MACHINE	1000	1000
1011	1	1000	EA	WASHING MACHINE	1000	1000
1012	1	1000	EA	WASHING MACHINE	1000	1000
1013	1	1000	EA	WASHING MACHINE	1000	1000
1014	1	1000	EA	WASHING MACHINE	1000	1000
1015	1	1000	EA	WASHING MACHINE	1000	1000
1016	1	1000	EA	WASHING MACHINE	1000	1000
1017	1	1000	EA	WASHING MACHINE	1000	1000
1018	1	1000	EA	WASHING MACHINE	1000	1000
1019	1	1000	EA	WASHING MACHINE	1000	1000
1020	1	1000	EA	WASHING MACHINE	1000	1000

Detect Anomalous  
Pattern Given  
 $M_0$

## Novel Pattern

1001	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1002	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1003	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1004	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1005	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1006	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1007	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1008	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1009	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1010	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1011	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1012	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1013	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1014	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1015	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1016	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1017	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1018	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1019	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4
1020	1	1000	EA	WASHING MACHINE	1000	1000	1	0.4	0.4

# HTE Pattern Detection

## Control Group

Item	Item No	Qty	Unit	Desc	Part No	Part Desc	Part Unit	Part Qty
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100

## Treatment Group

Item	Item No	Qty	Unit	Desc	Part No	Part Desc	Part Unit	Part Qty
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100

Detect Anomalous  
Subpopulation  
Given  
 $M_0$

## Novel Pattern

100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100
100	1	1	1	100	100	100	100	100

# HTE Pattern Detection

Control  
Group

Access Date	Time	Latitude	Long	DEPTH	TIME	COND
2017-08-01	12:00	41.250	-89.250	10	0.00	0.00
2017-08-01	12:01	41.250	-89.250	10	0.00	0.00
2017-08-01	12:02	41.250	-89.250	10	0.00	0.00
2017-08-01	12:03	41.250	-89.250	10	0.00	0.00
2017-08-01	12:04	41.250	-89.250	10	0.00	0.00
2017-08-01	12:05	41.250	-89.250	10	0.00	0.00
2017-08-01	12:06	41.250	-89.250	10	0.00	0.00
2017-08-01	12:07	41.250	-89.250	10	0.00	0.00
2017-08-01	12:08	41.250	-89.250	10	0.00	0.00
2017-08-01	12:09	41.250	-89.250	10	0.00	0.00
2017-08-01	12:10	41.250	-89.250	10	0.00	0.00
2017-08-01	12:11	41.250	-89.250	10	0.00	0.00
2017-08-01	12:12	41.250	-89.250	10	0.00	0.00
2017-08-01	12:13	41.250	-89.250	10	0.00	0.00
2017-08-01	12:14	41.250	-89.250	10	0.00	0.00
2017-08-01	12:15	41.250	-89.250	10	0.00	0.00
2017-08-01	12:16	41.250	-89.250	10	0.00	0.00
2017-08-01	12:17	41.250	-89.250	10	0.00	0.00
2017-08-01	12:18	41.250	-89.250	10	0.00	0.00
2017-08-01	12:19	41.250	-89.250	10	0.00	0.00
2017-08-01	12:20	41.250	-89.250	10	0.00	0.00
2017-08-01	12:21	41.250	-89.250	10	0.00	0.00
2017-08-01	12:22	41.250	-89.250	10	0.00	0.00
2017-08-01	12:23	41.250	-89.250	10	0.00	0.00
2017-08-01	12:24	41.250	-89.250	10	0.00	0.00
2017-08-01	12:25	41.250	-89.250	10	0.00	0.00
2017-08-01	12:26	41.250	-89.250	10	0.00	0.00
2017-08-01	12:27	41.250	-89.250	10	0.00	0.00
2017-08-01	12:28	41.250	-89.250	10	0.00	0.00
2017-08-01	12:29	41.250	-89.250	10	0.00	0.00
2017-08-01	12:30	41.250	-89.250	10	0.00	0.00
2017-08-01	12:31	41.250	-89.250	10	0.00	0.00
2017-08-01	12:32	41.250	-89.250	10	0.00	0.00
2017-08-01	12:33	41.250	-89.250	10	0.00	0.00
2017-08-01	12:34	41.250	-89.250	10	0.00	0.00
2017-08-01	12:35	41.250	-89.250	10	0.00	0.00
2017-08-01	12:36	41.250	-89.250	10	0.00	0.00
2017-08-01	12:37	41.250	-89.250	10	0.00	0.00
2017-08-01	12:38	41.250	-89.250	10	0.00	0.00
2017-08-01	12:39	41.250	-89.250	10	0.00	0.00
2017-08-01	12:40	41.250	-89.250	10	0.00	0.00
2017-08-01	12:41	41.250	-89.250	10	0.00	0.00
2017-08-01	12:42	41.250	-89.250	10	0.00	0.00
2017-08-01	12:43	41.250	-89.250	10	0.00	0.00
2017-08-01	12:44	41.250	-89.250	10	0.00	0.00
2017-08-01	12:45	41.250	-89.250	10	0.00	0.00
2017-08-01	12:46	41.250	-89.250	10	0.00	0.00
2017-08-01	12:47	41.250	-89.250	10	0.00	0.00
2017-08-01	12:48	41.250	-89.250	10	0.00	0.00
2017-08-01	12:49	41.250	-89.250	10	0.00	0.00
2017-08-01	12:50	41.250	-89.250	10	0.00	0.00
2017-08-01	12:51	41.250	-89.250	10	0.00	0.00
2017-08-01	12:52	41.250	-89.250	10	0.00	0.00
2017-08-01	12:53	41.250	-89.250	10	0.00	0.00
2017-08-01	12:54	41.250	-89.250	10	0.00	0.00
2017-08-01	12:55	41.250	-89.250	10	0.00	0.00
2017-08-01	12:56	41.250	-89.250	10	0.00	0.00
2017-08-01	12:57	41.250	-89.250	10	0.00	0.00
2017-08-01	12:58	41.250	-89.250	10	0.00	0.00
2017-08-01	12:59	41.250	-89.250	10	0.00	0.00
2017-08-01	13:00	41.250	-89.250	10	0.00	0.00

Treatment  
Group

[illegible]

Detect Anomalous  
Subpopulation  
Given  
 $M_0$

### Subpopulation

1000	6	20320	LINE 100-HE	AMERICAN TR	NET EXPRESS	TR	NET	EMPTY RACK
2000	6	20320	LINE 100-HE	AMERICAN TR	NET EXPRESS	TR	NET	EMPTY RACK
2000	6	20320	LINE 100-HE	AMERICAN TR	NET EXPRESS	TR	NET	EMPTY RACK
1000	6	20320	LINE 100-HE	AMERICAN TR	NET EXPRESS	TR	NET	EMPTY RACK
5000	6	20320	LINE 100-HE	AMERICAN TR	NET EXPRESS	TR	NET	EMPTY RACK

## THE GOAL

	Male	Female
Black		
White		
Hispanic		
Asian		
Native American		
Other		

Detect a subpopulation (subsets of attribute values), which correspond to anomalous outcomes for subjects in the treatment group

## THE GOAL

	Male	Female
Black		
White		
Hispanic		
Asian		
Native American		
Other		

Detect a subpopulation (subsets of attribute values), which correspond to anomalous outcomes for subjects in the treatment group

### The Optimization

$$S_I \subseteq \{a_1 \dots a_t\}, \dots, S_M \subseteq \{a_1 \dots a_p\}$$



## THE GOAL

	Male	Female
Black		
White		
Hispanic		
Asian		
Native American		
Other		

Detect a subpopulation (subsets of attribute values), which correspond to anomalous outcomes for subjects in the treatment group

### The Optimization

$$sI \subseteq \{a1...at\}, \dots, sM \subseteq \{a1...ap\}$$

$$S = sI \times \dots \times sM$$

## THE GOAL

	Male	Female
Black		
White		
Hispanic		
Asian		
Native American		
Other		

Detect a subpopulation (subsets of attribute values), which correspond to anomalous outcomes for subjects in the treatment group

### The Optimization

$$s_1 \subseteq \{a_1 \dots a_t\}, \dots, s_M \subseteq \{a_1 \dots a_p\}$$

$$S = s_1 \times \dots \times s_M$$

$$S^* = \operatorname{argmax}_S F(S)$$

# Treatment Effects Subset Scan (TESS)

	Male	Female
Black	$\Upsilon^{BM}$	$\Upsilon^{BF}$
White	$\Upsilon^{WM}$	$\Upsilon^{WF}$
Hispanic	$\Upsilon^{HM}$	$\Upsilon^{HF}$
Asian	$\Upsilon^{AM}$	$\Upsilon^{AF}$
Native American	$\Upsilon^{NM}$	$\Upsilon^{NF}$
Other	$\Upsilon^{OM}$	$\Upsilon^{OF}$

I. Compute the statistical anomalousness of each treatment group subject

II. Detect subpopulation that is collectively the most anomalous

# Treatment Effects Subset Scan (TESS)

	Male	Female
Black	$p^{BM}$	$p^{BF}$
White	$p^{WM}$	$p^{WF}$
Hispanic	$p^{HM}$	$p^{HF}$
Asian	$p^{AM}$	$p^{AF}$
Native American	$p^{NM}$	$p^{NF}$
Other	$p^{OM}$	$p^{OF}$

I. Compute the statistical anomalousness of each treatment group subject  
-- **This measurement will be a p-value**

II. Detect subpopulation that is collectively the most anomalous  
-- **Many subjects with significant p-values**

# Treatment Effects Subset Scan (TESS)

	Male	Female
Black	$\gamma^{BM}$	$\gamma^{BF}$
White	$\gamma^{WM}$	$\gamma^{WF}$
Hispanic	$\gamma^{HM}$	$\gamma^{HF}$
Asian	$\gamma^{AM}$	$\gamma^{AF}$
Native American	$\gamma^{NM}$	$\gamma^{NF}$
Other	$\gamma^{OM}$	$\gamma^{OF}$
	Control Group	

I. Compute the statistical anomalousness of each treatment group subject

I. Estimate Conditional Distribution Under  $H_0$

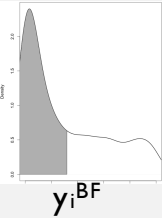
# Treatment Effects Subset Scan (TESS)

	Male	Female
Black	$\gamma^{BM}$	$\gamma^{BF}$
White	$\gamma^{WM}$	$\gamma^{WF}$
Hispanic	$\gamma^{HM}$	$\gamma^{HF}$
Asian	$\gamma^{AM}$	$\gamma^{AF}$
Native American	$\gamma^{NM}$	$\gamma^{NF}$
Other	$\gamma^{OM}$	$\gamma^{OF}$
Treatment Group		

- I. Compute the statistical anomalousness of each treatment group subject
  1. Estimate Conditional Distribution Under  $H_0$
  2. Compute empirical p-values

# Treatment Effects Subset Scan (TESS)

	Male	Female
Black	$\gamma^{BM}$	$\gamma^{BF}$
White	$\gamma^{WM}$	$\gamma^{WF}$
Hispanic	$\gamma^{HM}$	$\gamma^{HF}$
Asian	$\gamma^{AM}$	$\gamma^{AF}$
Native American	$\gamma^{NM}$	$\gamma^{NF}$
Other	$\gamma^{OM}$	$\gamma^{OF}$
Treatment Group		



I. Compute the statistical anomalousness of each treatment group subject

1. Estimate Conditional Distribution Under  $H_0$
2. Compute empirical p-values

# Treatment Effects Subset Scan (TESS)

	Male	Female
Black	$p^{BM}$	$p^{BF}$
White	$p^{WM}$	$p^{WF}$
Hispanic	$p^{HM}$	$p^{HF}$
Asian	$p^{AM}$	$p^{AF}$
Native American	$p^{NM}$	$p^{NF}$
Other	$p^{OM}$	$p^{OF}$
Treatment Group		

- I. Compute the statistical anomalousness of each treatment group subject
  1. Estimate Conditional Distribution Under  $H_0$
  2. Compute empirical p-values



# Treatment Effects Subset Scan (TESS)

	Male	Female
Black	$p^{BM}$	$p^{BF}$
White	$p^{WM}$	$p^{WF}$
Hispanic	$p^{HM}$	$p^{HF}$
Asian	$p^{AM}$	$p^{AF}$
Native American	$p^{NM}$	$p^{NF}$
Other	$p^{OM}$	$p^{OF}$
Treatment Group		

- I. Compute the statistical anomalousness of each treatment group subject
  1. Estimate Conditional Distribution Under  $H_0$
  2. Compute empirical p-values
    - i. Maps each bin's distribution to the same interval

# Treatment Effects Subset Scan (TESS)

	Male	Female
Black	$p^{BM}$	$p^{BF}$
White	$p^{WM}$	$p^{WF}$
Hispanic	$p^{HM}$	$p^{HF}$
Asian	$p^{AM}$	$p^{AF}$
Native American	$p^{NM}$	$p^{NF}$
Other	$p^{OM}$	$p^{OF}$
	Treatment Group	

- I. Compute the statistical anomalousness of each treatment group subject
  1. Estimate Conditional Distribution Under  $H_0$
  2. Compute empirical p-values
    - i. Maps each bin's distribution to the same interval
    - ii.  $P^{ij} \sim \text{Uniform}[0,1]$  under  $H_0$

# Treatment Effects Subset Scan (TESS)

	Male	Female
Black	$p^{BM}$	$p^{BF}$
White	$p^{WM}$	$p^{WF}$
Hispanic	$p^{HM}$	$p^{HF}$
Asian	$p^{AM}$	$p^{AF}$
Native American	$p^{NM}$	$p^{NF}$
Other	$p^{OM}$	$p^{OF}$
Treatment Group		

I. Compute the statistical anomalousness of each treatment group subject

1. Estimate Conditional Distribution Under  $H_0$

2. Compute empirical p-values

i. Maps each bin's distribution to the same interval

ii.  $P_{ij} \sim \text{Uniform}[0,1]$  under  $H_0$

iii. For any  $N$  p-values, we expect  $N*\alpha$  to be significant at level  $\alpha$

# Treatment Effects Subset Scan (TESS)

	Male	Female
Black	$p^{BM}$	$p^{BF}$
White	$p^{WM}$	$p^{WF}$
Hispanic	$p^{HM}$	$p^{HF}$
Asian	$p^{AM}$	$p^{AF}$
Native American	$p^{NM}$	$p^{NF}$
Other	$p^{OM}$	$p^{OF}$
	Treatment Group	

I. Compute the statistical anomalousness of each treatment group subject

1. Estimate Conditional Distribution Under  $H_0$

2. Compute empirical p-values

i. Maps each bin's distribution to the same interval

ii.  $P_{ij} \sim \text{Uniform}[0,1]$  under  $H_0$

iii. For any  $N$  p-values, we expect  $N*\alpha$  to be significant at level  $\alpha$

Higher Criticism:

$$F(S) = \max_{\alpha} \frac{N_{\alpha} - N\alpha}{\sqrt{N\alpha(1-\alpha)}}$$

# Treatment Effects Subset Scan (TESS)

	Male	Female
Black	$p^{BM}$	$p^{BF}$
White	$p^{WM}$	$p^{WF}$
Hispanic	$p^{HM}$	$p^{HF}$
Asian	$p^{AM}$	$p^{AF}$
Native American	$p^{NM}$	$p^{NF}$
Other	$p^{OM}$	$p^{OF}$
	Treatment Group	

- I. Compute the statistical anomalousness of each treatment group subject
- II. Discover subsets of attribute values that define the most anomalous outcomes
  - I. Maximize  $F(S)$  over all subsets of  $s_1 \times \dots \times s_M$ 
    - Naïve search is infeasible  $O(2^{\sum |A_i|})$

# Treatment Effects Subset Scan (TESS)

## Nonparametric Scan Statistic (NPSS)

Have:  $S \subseteq \{A_1 \times \dots \times A_M\}$   
 $= \{s_1 \times \dots \times s_M\}$

- I. Compute the statistical anomalousness of each treatment group subject
- II. Discover subsets of attribute values that define the most anomalous outcomes
  - I. Maximize  $F(S)$  over all subsets of  $s_1 \times \dots \times s_M$ 
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# Treatment Effects Subset Scan (TESS)

## Nonparametric Scan Statistic (NPSS)

Have:  $S \subseteq \{A_1 \times \dots \times A_M\}$   
 $= \{s_1 \times \dots \times s_M\}$

Select:  $F(S) = \max_{\alpha} \phi(\alpha, N_{\alpha}(S), N(S))$

Want:  $\max_S F(S)$

Assume:  $\phi \uparrow$  w.r.t  $N_{\alpha}$   
 $\phi \downarrow$  w.r.t  $N$  and  $\alpha$   
 $\phi$  is convex

- I. Compute the statistical anomalousness of each treatment group subject
- II. Discover subsets of attribute values that define the most anomalous outcomes
  - I. Maximize  $F(S)$  over all subsets of  $s_1 \times \dots \times s_M$ 
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Want:  $\max_S F(S)$

Assume:  $\phi \uparrow$  w.r.t  $N_{\alpha}$   
 $\phi \downarrow$  w.r.t  $N$  and  $\alpha$   
 $\phi$  is convex

There Exist:  $G(a_i)$

Such That:  $\max_{s_j \subseteq \{a_1, \dots, a_t\}} F(s_j | A_{-j}) = \max_{i=1 \dots t} F(\{a_{(1)} \dots a_{(t)}\} | A_{-j})$

Only Consider:  $\{a_{(1)}\}$

$\{a_{(1)}, a_{(2)}\}$

$\vdots$

$\{a_{(1)}, \dots, a_{(M)}\}$

I. Compute the statistical anomalousness  
of each treatment group subject

II. Discover subsets of attribute values  
that define the most anomalous outcomes

I. Maximize  $F(S)$  over all subsets of

$s_1 \times \dots \times s_M$

• Naïve search is infeasible  $O(2^{\sum |A_i|})$



# Treatment Effects Subset Scan (TESS)

## Nonparametric Scan Statistic (NPSS)

Have:  $S \subseteq \{A_1 \times \dots \times A_M\}$   
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Only Consider:  $\{\text{Black}\}$

$\{\text{Black, Hispanic}\}$

$\vdots$

$\{\text{Black, Hispanic, Asian, ..., White}\}$

- I. Compute the statistical anomalousness of each treatment group subject
- II. Discover subsets of attribute values that define the most anomalous outcomes

- I. Maximize  $F(S)$  over all subsets of

$s_1 \times \dots \times s_M$

•Naïve search is infeasible  $O(2^{\sum |A_i|})$

# Treatment Effects Subset Scan (TESS)

## Nonparametric Scan Statistic (NPSS)

Have:  $S \subseteq \{A_1 \times \dots \times A_M\}$   
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Select:  $F(S) = \max_{\alpha} \phi(\alpha, N_{\alpha}(S), N(S))$

Want:  $\max_S F(S)$

Assume:  $\phi \uparrow$  w.r.t  $N_{\alpha}$   
 $\phi \downarrow$  w.r.t  $N$  and  $\alpha$   
 $\phi$  is convex

There Exist:  $G(a_i) = \frac{1}{n(a_i)} \sum_{A_{-j}} I(p_{ij} \leq \alpha)$

Such That:  $\max_{s_j \subseteq \{a_1, \dots, a_t\}} F(s_j | A_{-j}) = \max_{i=1 \dots t} F(\{a_{(1)} \dots a_{(t)}\} | A_{-j})$

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$\{\text{Black, Hispanic}\}$

$\vdots$

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Intuitively:  $F(\{a_{(1)}, a_{(3)}\}) \leq F(\{a_{(1)}, a_{(2)}\})$   
 $G(a_{(3)}) \leq G(a_{(2)})$

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- II. Discover subsets of attribute values that define the most anomalous outcomes

- I. Maximize  $F(S)$  over all subsets of

$s_1 \times \dots \times s_M$

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I. Maximize  $F(S)$  over all subsets of

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Higher Criticism:

$$F(S) = \max_{\alpha} \frac{N_{\alpha} - N\alpha}{\sqrt{N\alpha(1-\alpha)}}$$

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## Nonparametric Scan Statistic (NPSS)

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Assume:  $\phi \uparrow$  w.r.t  $N_{\alpha}$   
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Intuitively:  $F(\{a_{(1)}, a_{(3)}\}) \leq F(\{a_{(1)}, a_{(2)}\})$   
 $G(a_{(3)}) \leq G(a_{(2)})$

I. Compute the statistical anomalousness  
of each treatment group subject

II. Discover subsets of attribute values  
that define the most anomalous outcomes

I. Maximize  $F(S)$  over all subsets of

$s_1 \times \dots \times s_M$

•NPSS over an attribute in  $O(t \log t)$

Higher Criticism:

$$F(S) = \max_{\alpha} \frac{N_{\alpha} - N\alpha}{\sqrt{N\alpha(1-\alpha)}}$$

# TESS Search Procedure

	Male	Female
Black	$p^{BM}$	$p^{BF}$
White	$p^{WM}$	$p^{WF}$
Hispanic	$p^{HM}$	$p^{HF}$
Asian	$p^{AM}$	$p^{AF}$
Native American	$p^{NM}$	$p^{NF}$
Other	$p^{OM}$	$p^{OF}$
	Treatment Group	

I. Compute the statistical anomalousness of each treatment group subject

1. Estimate Conditional Distribution Under  $H_0$
2. Compute empirical p-values

II. Discover subsets of attribute values that define the most anomalous outcomes

I. Maximize  $F(S)$  over all subsets of

$$s_1 \times \dots \times s_M$$

•NPSS over an attribute in  $O(t \log t)$

# TESS Search Procedure

	Male	Female
Black	$p^{BM}$	$p^{BF}$
White	$p^{WM}$	$p^{WF}$
Hispanic	$p^{HM}$	$p^{HF}$
Asian	$p^{AM}$	$p^{AF}$
Native American	$p^{NM}$	$p^{NF}$
Other	$p^{OM}$	$p^{OF}$

(Score = 7.5)

I. Compute the statistical anomalousness of each treatment group subject

1. Estimate Conditional Distribution Under  $H_0$
2. Compute empirical p-values

II. Discover subsets of attribute values that define the most anomalous outcomes

1. Maximize  $F(S)$  over all subsets of

$$s_1 \times \dots \times s_M$$

•NPSS over an attribute in  $O(t \log t)$

# TESS Search Procedure

	Male	Female	
Black	$p^{BM}$	$p^{BF}$	←
White	$p^{WM}$	$p^{WF}$	
Hispanic	$p^{HM}$	$p^{HF}$	←
Asian	$p^{AM}$	$p^{AF}$	
Native American	$p^{NM}$	$p^{NF}$	
Other	$p^{OM}$	$p^{OF}$	

(Score = 8.1)

I. Compute the statistical anomalousness of each treatment group subject

1. Estimate Conditional Distribution Under  $H_0$
2. Compute empirical p-values

II. Discover subsets of attribute values that define the most anomalous outcomes

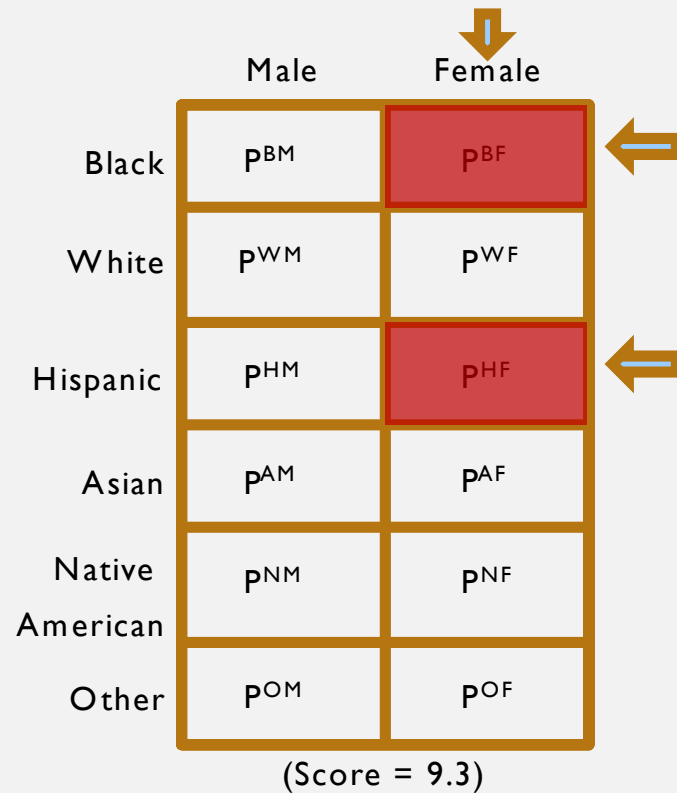
1. Maximize  $F(S)$  over all subsets of

$$S_1 \times \dots \times S_M$$

•NPSS over an attribute in  $O(t \log t)$



# TESS Search Procedure



	Male	Female
Black	$p^{BM}$	$p^{BF}$
White	$p^{WM}$	$p^{WF}$
Hispanic	$p^{HM}$	$p^{HF}$
Asian	$p^{AM}$	$p^{AF}$
Native American	$p^{NM}$	$p^{NF}$
Other	$p^{OM}$	$p^{OF}$

(Score = 9.3)

I. Compute the statistical anomalousness of each treatment group subject

1. Estimate Conditional Distribution Under  $H_0$
2. Compute empirical p-values

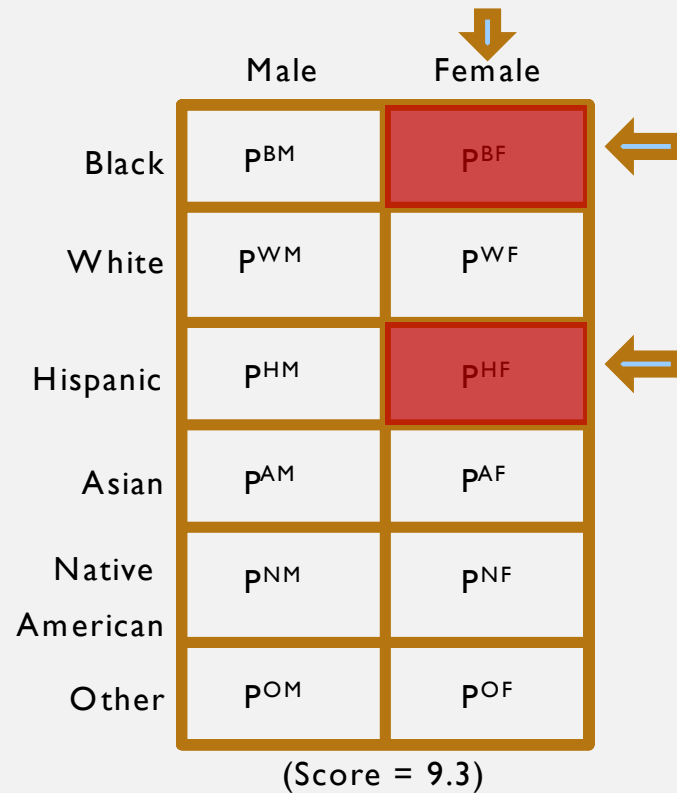
II. Discover subsets of attribute values that define the most anomalous outcomes

I. Maximize  $F(S)$  over all subsets of

$$S_1 \times \dots \times S_M$$

•NPSS over an attribute in  $O(t \log t)$

# TESS Search Procedure



	Male	Female
Black	$p^{BM}$	$p^{BF}$
White	$p^{WM}$	$p^{WF}$
Hispanic	$p^{HM}$	$p^{HF}$
Asian	$p^{AM}$	$p^{AF}$
Native American	$p^{NM}$	$p^{NF}$
Other	$p^{OM}$	$p^{OF}$

(Score = 9.3)

I. Compute the statistical anomalousness of each treatment group subject

1. Estimate Conditional Distribution Under  $H_0$
2. Compute empirical p-values

II. Discover subsets of attribute values that define the most anomalous outcomes

1. Maximize  $F(S)$  over all subsets of

$$S_1 \times \dots \times S_M$$

•NPSS over an attribute in  $O(t \log t)$

## Significance of our subpopulation

Compare subpopulation score to maximum scores of simulated datasets under  $H_0$

## CASE STUDY: TENNESSEE STAR

## TENNESSEE STAR ANALYSIS (1985)

- Effect of classrooms size on achievement (test scores)
- 4 year panel (kindergarten to 3rd grade)
- 6,500 students, 330 classrooms, 80 schools
  - Total of over 11,000 records
- Treatment Conditions (randomized within school)
  - Regular Size Class (20-25 students)
  - Regular Size + Aide Class (20-25 students)
  - Small Size Class (13-17 students)

## TENNESSEE STAR ANALYSIS

read	math	gender	ethnicity	lunch	grade	school	experience	degree	tethnicity	schoolid
439	463	male	afam	free	kindergarten	inner-city	0	bachelor	cauc	19
448	559	male	cauc	non-free	kindergarten	rural	16	bachelor	cauc	69
431	454	male	cauc	free	kindergarten	rural	8	bachelor	cauc	5
395	423	female	afam	free	kindergarten	inner-city	17	master	cauc	16
451	500	female	cauc	non-free	kindergarten	rural	3	bachelor	afam	56
430	473	male	cauc	non-free	kindergarten	rural	13	master	cauc	38
437	468	male	cauc	non-free	kindergarten	rural	6	master	cauc	69
490	528	male	cauc	non-free	kindergarten	suburban	18	bachelor	cauc	52
439	484	male	cauc	non-free	kindergarten	suburban	13	master	cauc	54
424	459	female	cauc	free	kindergarten	rural	12	bachelor	cauc	12
437	528	female	afam	free	kindergarten	suburban	1	bachelor	afam	21
424	559	male	cauc	free	kindergarten	rural	13	bachelor	cauc	79
431	454	male	cauc	non-free	kindergarten	rural	13	master	cauc	8
451	473	male	cauc	non-free	kindergarten	rural	3	bachelor	cauc	66
421	459	female	afam	free	kindergarten	inner-city	11	bachelor	cauc	31

## TENNESSEE STAR ANALYSIS

	(1)	(2)
Treatment	3.4791	-0.2909
	(2.547)	(2.277)
Sample	All 2 <sup>nd</sup> Grade	All 3 <sup>rd</sup> Grade
R-squared	0.000	0.000
Observations	4263	4063

Notes: All estimates are from OLS models.

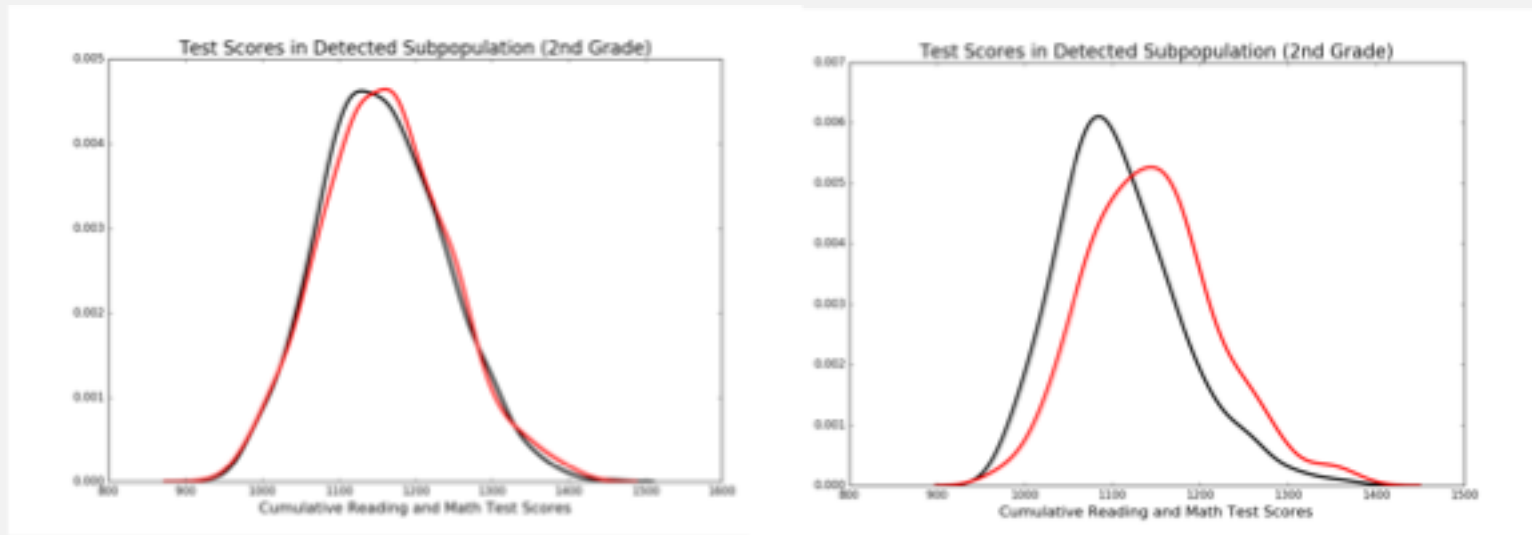
Standard errors are in parentheses.

\*\*\* p<0.01, \*\* p<0.05, \* p<0.1

## TENNESSEE STAR ANALYSIS

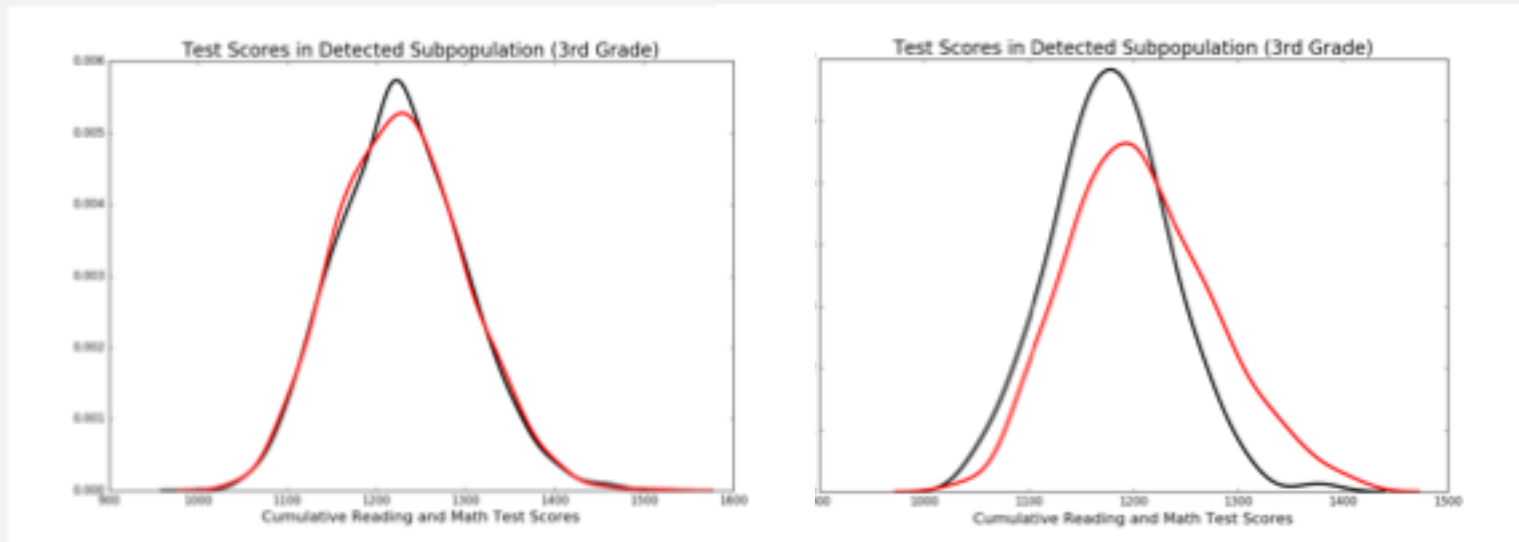
- Detected Subpopulation
  - grade:
    - 2nd or 3rd
  - school:
    - inner-city or urban
  - experience:
    - [10, infinity)
  - other features do not have differential effects

# TENNESSEE STAR ANALYSIS





# TENNESSEE STAR ANALYSIS



## TENNESSEE STAR ANALYSIS

	(1)	(2)	(3)
Treatment	3.4791	36.066***	1.309
	(2.547)	(6.055)	(2.772)
Sample	All 2 <sup>nd</sup> Grade	Detected Group (2 <sup>nd</sup> Grade)	Undetected Group (2 <sup>nd</sup> Grade)
P-value	0.172	<0.001	0.0637
Observations	4263	620	3643

Notes: All estimates are from OLS models.

Standard errors are in parentheses.

\*\*\*  $p < 0.001$ , \*\*  $p < 0.05$ , \*  $p < 0.1$

## TENNESSEE STAR ANALYSIS

	(1)	(2)	(3)
Treatment	-0.291	18.703***	0.1
	(2.277)	(5.18)	(2.478)
Sample	All 3 <sup>rd</sup> Grade	Detected Group (3 <sup>rd</sup> Grade)	Undetected Group (3 <sup>rd</sup> Grade)
P-value	0.898	<0.001	0.968
Observations	4063	706	3357

Notes: All estimates are from OLS models.

Standard errors are in parentheses.

\*\*\*  $p < 0.001$ , \*\*  $p < 0.05$ , \*  $p < 0.1$

THE END!



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## OTHER PAPERS/IDEAS

- Generalized Adversarial Method of Moments
- High-dimensional regression adjustments in randomized experiments
- Lasso adjustments of treatment effect estimates in randomized experiments
- A Simple Method for Estimating Interactions Between a Treatment and a Large Number of Covariates
- A Nonparametric Bayesian Analysis of Heterogenous Treatment Effects in Digital Experimentation
- Bayesian Nonparametric Modeling for Causal Inference
- Deep IV: A Flexible Approach for Counterfactual Prediction