### Potential outcomes

• A feature vector  $X \in \mathbb{R}^k$ 

- A treatment assignment  $A \in \{0,1\}$
- Potential outcomes: Y(1), Y(0) Observe Y := Y(A), never Y(1 A)

### **Average Treatment Effect (ATE)**

$$\begin{split} ATE &= \mathbb{E}[Y(1) - Y(0)] \\ &= \mathbb{E}_{X \sim P_X} \left[ \mathbb{E}[Y(1) \,|\, X] - \mathbb{E}[Y(0) \,|\, X] \right] \\ &= \mathbb{E}_{X \sim P_X} \left[ \mu_1^{\star}(X) - \mu_0^{\star}(X) \right] \\ &= \mathbb{E}_{X \sim P_X} \left[ \mu_1^{\star}(X) - \mu_0^{\star}(X) \right] \\ &= : \mathbb{E}_{X \sim P_X} \left[ \mu_1^{\star}(X) - \mu_0^{\star}(X) \right] \\ \end{split}$$

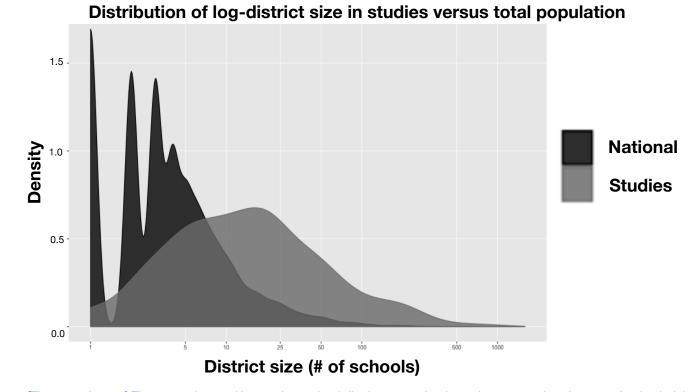
 $\bullet$   $P_X$  is the data generating distribution for X

# Problem I: population shifts

a.k.a. X-shift, covariate shit

### Problem I: what if $P_X$ changes?

 Even for carefully designed randomized trials, "statistics" starts only at treatment assignment, with big biases in selection into study



[Tipton et al. 2019] The convenience of large urban school districts: a study of recruitment practices in 37 randomized trials

# Problem I: what if $P_X$ changes?

- "Clinical trials for new drugs skew heavily white" [Oh et al. '15, Burchard et al. '15, SA Editors '18]
  - Out of 10,000+ cancer trials, less than 5% of participants were non-white
- Even large clinical trials suffer from these biases. Recently, two large trials with n = 5K-10K had opposite findings on a treatment to lower blood pressure on cardiovascular disease [Leigh et al. '16, Imai et al. '13, Gijsberts et al. '15, Basu et al. '17, Duan et al. '19]

## Problem II: unobserved confounders

a.k.a. Y | X shift

### Unobserved confounders

 There always exists unobserved confounders that simultaneously affect potential outcomes and treatment assignments

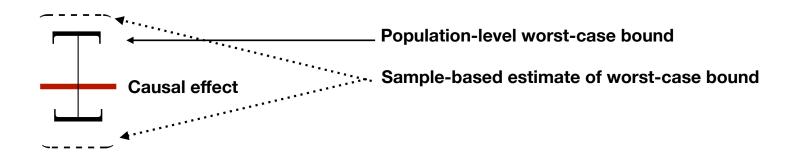
Judges are more lenient after taking a break, study finds theguardian [Danziger '11]

Overlooked factors in the analysis of parole decisions [Weinshall-Margel '11]

- Visual observations used in clinical decisions and drugs preferentially prescribed those who can tolerate them
  - Not properly recorded even at the resolution of large databases

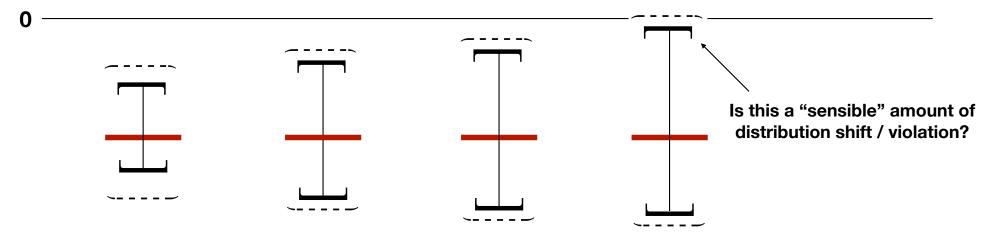
## Worst-case approach

- Posit a set of "plausible" distribution shifts, and take worst-case over them
- If effects are still valid under plausible shifts, we can certify robustness
- Sensitivity of a finding: magnitude of shift when endpoint crosses a threshold
- Today: worst-case bounds on the Doubly Robust estimator



## Worst-case approach

- Posit a set of "plausible" distribution shifts, and take worst-case over them
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# Part I: External validity

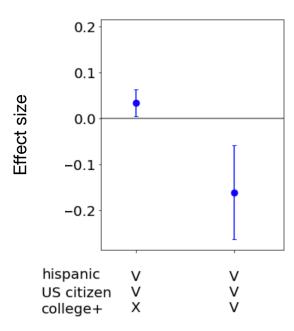
a.k.a. X-shift, covariate shit

## Challenges

- X-shift problematic when treatment effect is heterogeneous
  - Healthcare: across demographics, comorbidities, and concomitant drugs
- Option 1: Directly estimate conditional average treatment affect (CATE)?
  - ML models unstable on underrepresented groups; resulting inference underpowered
- Option 2: Subgroup analysis?
  - Difficult due to intersectionality



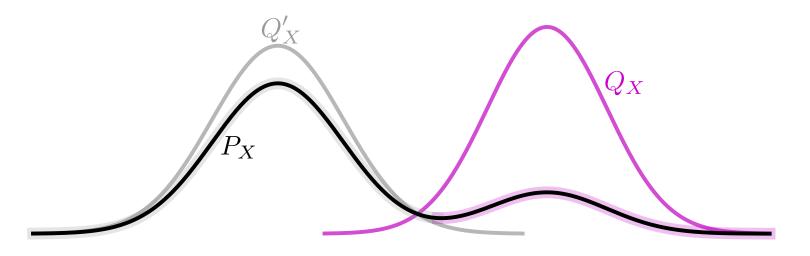
### Effect of Medicaid enrollment on doctor's office utilization



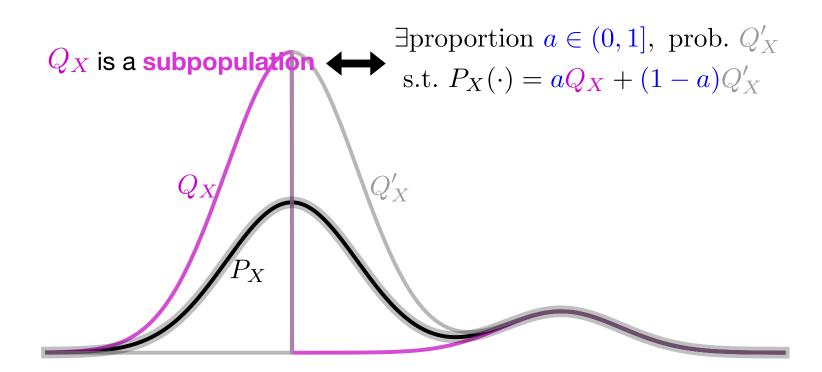
[Leigh et al. '16, Imai et al. '13, Gijsberts et al. '15, Basu et al. '17, Baum et al. '17, Duan et al. '19, Nie and Wager '20]

Automatically find worst-off subpopulations and measure treatment effect on them

$$Q_X$$
 is a subpopulation  $\longleftrightarrow$   $\exists \text{proportion } a \in (0,1], \text{ prob. } Q_X'$  s.t.  $P_X(\cdot) = aQ_X + (1-a)Q_X'$ 

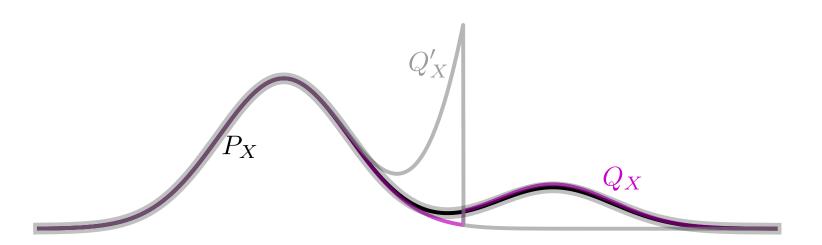


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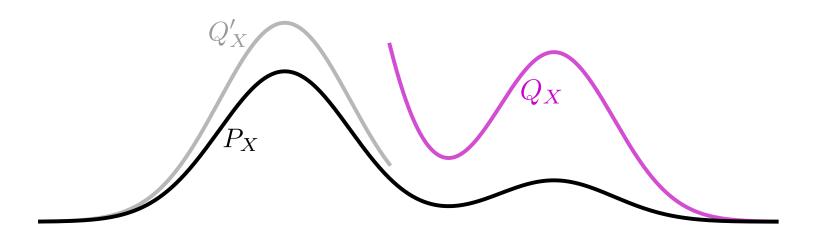
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## **Worst-case subpopulation**

#### Recap

- Covariates: X
- ► Treatment assignment: A
- ▶ Potential outcome: Y(0), Y(1)
- Response Y := Y(Z)

$$Q_X \succeq \alpha \iff$$

subpopulation with proportion larger than  $\alpha \in (0, 1]$ 

### **Worst-case Subpopulation Treatment Effect**

$$WTE_{\alpha} := \sup_{Q_X \succeq \alpha} \mathbb{E}_{Q_X} [\mu^{\star}(X)]$$

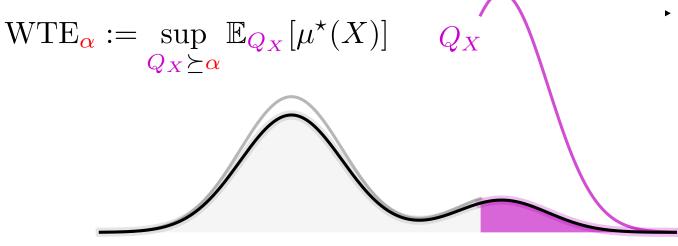
where 
$$\mu^{\star}(X) := \mathbb{E}[Y(1) - Y(0) \mid X]$$

is the conditional average treatment effect (CATE).

## WTE = Tail-average

#### Recap

- Covariates: X
- Treatment assignment: A
- Potential outcome: Y(0), Y(1)
- CATE  $\mu^{\star}(X) = \mathbb{E}[Y(1) Y(0) \mid X]$



Lemma (Shapiro et al. '09)

$$\sup_{Q_X \succeq \alpha} \mathbb{E}_{Q_X} [\mu^*(X)] = \mathbb{E}[\mu^*(X) \mid \mu^*(X) \ge q_\alpha^*]$$

 $(1-\alpha)$ -quantile of  $\mu^{\star}(X)$ 

### Main Result

$$\omega(\mu^*) := \sup_{Q_X \succeq \alpha} \mathbb{E}_{Q_X}[\mu^*(X)]$$

- Use any ML method to fit  $\mu^*(X) = \mathbb{E}[Y|X, A=1] \mathbb{E}[Y|X, A=0]$
- Debiasing: Correct plug-in estimator using the first-order error

$$\omega(\hat{\mu}) - \omega(\mu^*) = \nabla \omega(\hat{\mu})^{\mathsf{T}} (\hat{\mu} - \mu^*) + \mathsf{Rem}$$

Debiased estimator automatically only has second-order error

#### Theorem (Jeong & N. '20)

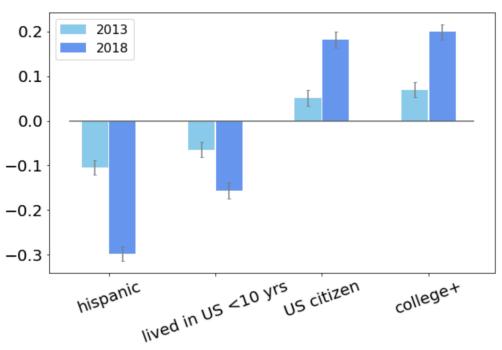
- 1. Even when nuisance parameters converge more slowly,  $\sqrt{n}(\hat{\omega} \omega) \Rightarrow N(0, \sigma^2)$
- 2.  $\sigma^2$  is the **optimal** asymptotic variance



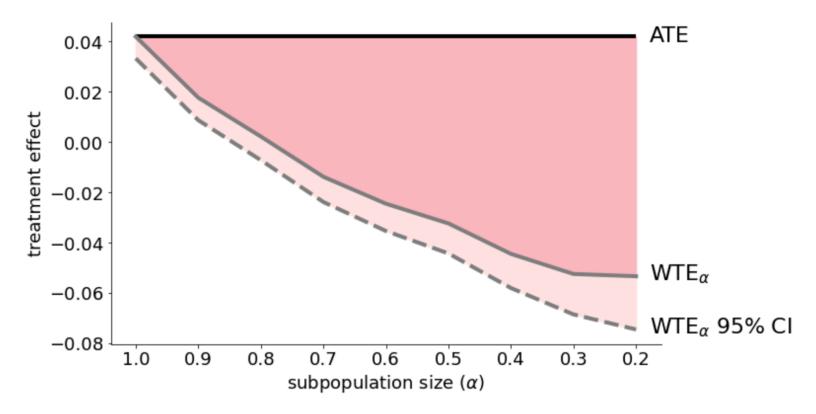
- Evaluate effect of Medicaid enrollment on doctors' office utilization
- Medicaid costs \$553 billion/yr; need to ensure valid effects through time
- Outcome: visit to doctors in the two-weeks prior to a random survey date
- Control for demographics, medical history, employment, earnings, insurance, government assistance etc (d = 396)
- Take the viewpoint of an analyst in 2009 (n = 82,993)

### Demographic compositions shift over time

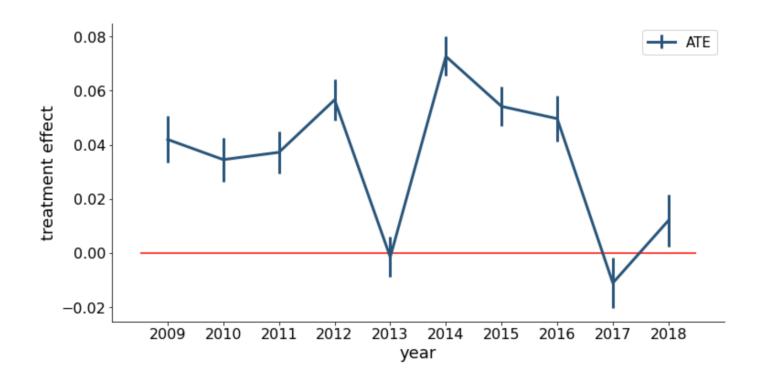
#### Change in share from 2009



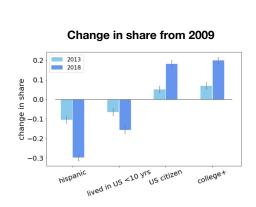
• Evaluate effect of Medicaid enrollment on doctors' office utilization in 2009

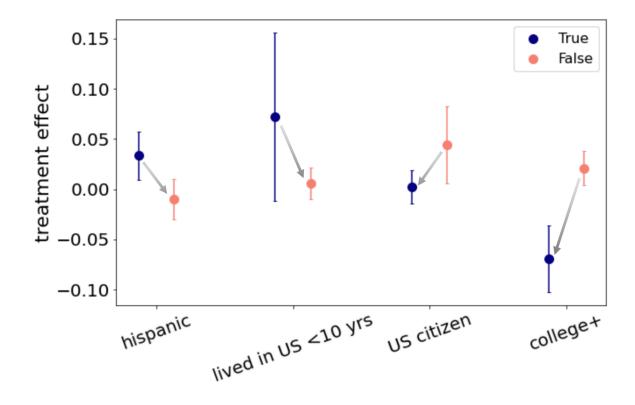


Evaluate effect of Medicaid enrollment on doctors' office utilization



Evaluate effect of Medicaid enrollment on doctors' office utilization



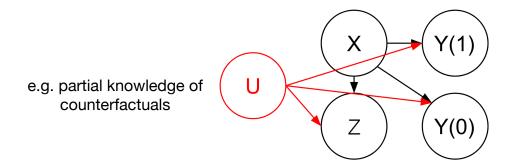


## Part II: unobserved confounders

a.k.a. Y | X shift

## Bounded unobserved confounding

What if there's a hidden variable U that wasn't observed?



#### Relaxed assumption: Bounded unobserved confounding

There exists  $\Gamma > 1$ , and U such that  $Y(1), Y(0) \perp A \mid X, U$ ,

$$u\mapsto \frac{\mathbb{P}(A=1\mid X,U=u)}{\mathbb{P}(A=0\mid X,U=u)}$$
 can vary by at most a factor of  $\Gamma$  [Rosenbaum '02]

## Bounded unobserved confounding

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• Equivalent to a logit model: for some function  $\kappa(\cdot) \in [0,1], g(\cdot),$ 

$$\log \frac{\mathbb{P}(A=1\mid X,U)}{\mathbb{P}(A=0\mid X,U)} = g(X) + \log \Gamma \cdot \kappa(X,U)$$

### **FAQs**

#### Relaxed assumption: Bounded unobserved confounding

There exists  $\Gamma > 1$ , and U such that  $Y(1), Y(0) \perp A \mid X, U$ ,

$$u\mapsto \frac{\mathbb{P}(A=1\mid X,U=u)}{\mathbb{P}(A=0\mid X,U=u)}$$
 can vary by at most a factor of  $\Gamma$  [Rosenbaum '02]

- How do I choose  $\Gamma$ ?
- **→** Domain expertise (e.g. clinical intuition)
- $\implies$  Sensitivity: what would be a clinically significant result? what value of  $\Gamma$  would change its significance?
- Is this the only natural confounding model?
- → No. Today: modern semiparametric framework.

### Lower bound for $\mathbb{E}[Y(1) \mid X]$

#### Recap

- Treatment assignment: A
- ▶ Potential outcome: Y(0), Y(1)
- Response Y := Y(Z)

#### unobservable

ullet Lower bound unobservables under  $\Gamma$ -bounded unobserved confounding

$$\mathbb{E}[Y(1) \mid X, A = 0] = \mathbb{E}[YL(Y \mid X) \mid X, A = 1]$$
 **observable**  $L(\cdot \mid X) := \frac{dP(Y(1) \in \cdot \mid X, A = 1)}{dP(Y(1) \in \cdot \mid X, A = 0)}$ 

**Lemma** Under  $\Gamma$ -confounding,  $y \mapsto L(y \mid x)$  can vary by at most a factor of  $\Gamma$ 

Minimizing over the above set of likelihood ratios,

$$\mathbb{E}[Y(1) \mid X, A = 0] \ge \inf_{L \in \mathcal{L}_1} \mathbb{E}[YL(Y|X) \mid X, A = 1] =: \theta_1^{\star}(X)$$
 Bound is tight

### **Convex Duality**

#### Recap

► Treatment assignment: Z

▶ Potential outcome: Y(0), Y(1)

• Response Y := Y(Z)

**Lemma** Under  $\Gamma$ -confounding,  $y \mapsto L(y \mid x)$  can vary by at most a factor of  $\Gamma$ 

• Minimizing over the above set of likelihood ratios,

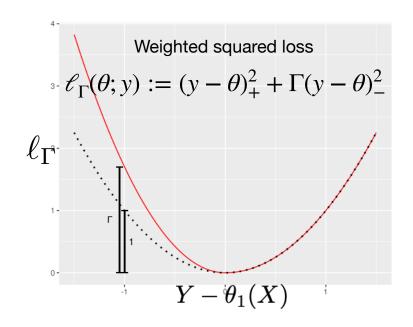
$$\mathbb{E}[Y(1) \mid X, A = 0] \ge \inf_{L \in \mathcal{L}_1} \mathbb{E}[YL(Y|X) \mid X, A = 1] =: \theta_1^{\star}(X)$$
 Bound is tight

One-dimensional dual for each X

Lemma  $\theta_1^*(X) = \sup \{ \mu : \mathbb{E}[(Y(1) - \mu)_+ - \Gamma(Y(1) - \mu)_- \mid X, A = 1] \ge 0 \}$ 

### What can ML do?

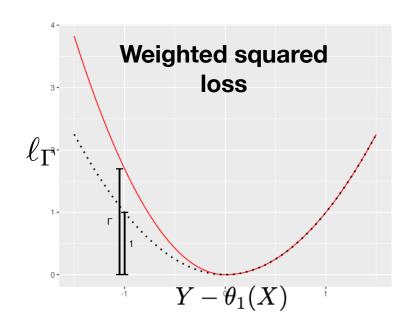
- Tremendous empirical success is curve-fitting tools in highdimensions, under noisy data
- Key ingredients: stochastic optimization & model selection



### Sensitivity of CATE via loss minimization

$$\bullet \, \mathbb{E}[Y(1) \mid X, A = 0] \ge \theta_1^{\star}(X) = \sup \left\{ \mu : \mathbb{E}[(Y(1) - \mu)_+ - \Gamma(Y(1) - \mu)_- \mid X, A = 1] \ge 0 \right\}$$

**Main result I:**  $\theta_1^{\star}$  is the unique solution to  $\min_{\theta(\cdot)} \mathbb{E}[\ell_{\Gamma}(\theta(X); Y(1)) \mid A = 1]$ 



- Estimate lower bound using flexible ML models
- Solve weighted regression problem using any black-box ML approach
- e.g., random forests, boosted trees, NNs

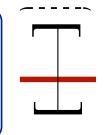
## Lower bound for $\mathbb{E}[Y(1)]$

#### Recap

- ► Treatment assignment: Z
- ► Potential outcome: *Y*(0), *Y*(1)
- Response Y := Y(Z)
- Similarly as before, we derive a **debiased estimator** for  $\mu_1^- = \mathbb{E}[AY(1) + (1-A)\theta_1^*(X)] \leq \mathbb{E}[Y(1)]$
- Bounds the doubly robust estimator for the ATE; equal when  $\Gamma=1$
- Value of prediction: DR estimator close to worst-case bound  $\mu_1^-$  (a.k.a. robust to confounding) when residuals  $Y \hat{\theta}_1(X)$  are small

**Theorem** Even when ML-based nuisance estimators converge at slower rates,

$$\sqrt{n}(\widehat{\mu}_1^- - \mu_1^-) \Rightarrow N(0,\sigma^2)$$

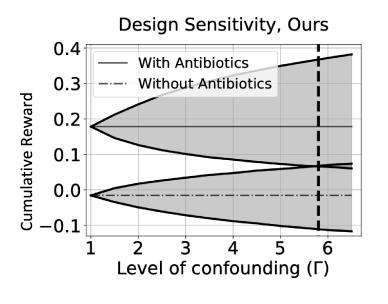


### Sepsis management in the ICU

- Sepsis in ICU patients accounts for 1/3 of deaths in hospitals [Howell and Davis '17]
- Automated approaches can manage important medication for sepsis [Futoma '18; Komorowski 18; Raghu 17]
- ICU data suffers from unobserved confounders
- ED physician: "initial treatment of antibiotics at admission to the hospital are often confounded by unrecorded factors that affect the eventual outcome (death or discharge from the ICU)."

### Proof of concept

- Whether to quickly begin antibiotic treatment is a topic of much discussion: balance early treatment vs. risks of over-prescription [Seymour '17; Sterling '15]
- Two policies: with or without antibiotics in the first step
- We use simulator developed by Obserst and Sontag (2019)



Our approach allows certifying robustness under realistic values of confounding

## Summary

- Worst-case bounds on the causal effect estimated through ML models
- Debiasing: CLT even when nuisance estimates converge slower; optimal
- Guard against brittle findings that do not hold under distribution shift

Assessing External Validity Over Worst-case Subpopulations.

Jeong & N. Under review. Short version appeared in COLT 2020.

Bounds on the conditional and average treatment effect with unobserved confounding factors.

Yadlowsky, N., Basu, Duchi, and Tian. Annals of Statistics, 2022.

Off-policy policy evaluation for sequential decisions under unobserved confounding.

N., Keramati, Yadlowsky, and Brunskill. NeurlPS 2020.