

# Sensitivity Analysis for Unobserved Effect Modification when Generalizing Findings from Randomized Trials to Target Populations

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# Outline

- 1 Background on generalizability
- 2 Motivation for sensitivity analyses
- 3 Notation
- 4 Sensitivity Analysis for a Partially Unobserved Modifier ( $V$  Case)
- 5 Sensitivity Analysis for a Fully Unobserved Modifier ( $U$  Case)
- 6 Extension to Non-Continuous Outcomes
- 7 Conclusion



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# Making research results relevant: A range of policy or practice questions

- Medicare may be deciding whether or not to approve payment for a new treatment for back pain
- Interest in predicting overall population impacts of a broad public health media campaign around not switching car seats to forward facing until a child is 12 months old
- A health care system may be trying to predict the effects of an evidence-based intervention in their overall patient population
- A physician practice may be deciding whether training providers in a new intervention would be cost effective across their population



# Estimating Effects: Randomized Trials

- Randomized trials are gold standard for estimating causal effect of a treatment in a study sample
- Strong internal validity allows for unbiased estimation of the sample average treatment effect (SATE)
  - However, some policy questions require estimating the target population average treatment effect (TATE), not SATE



# Issue: RCTs May Not Generalize

- Randomized trials often have poor external validity (generalizability) (Weisberg et al., 2009)
- Potential reasons for this:
  - Strict exclusion criteria (Stuart et al., 2011)
  - Lack of attention to target population when implementing trial
- SATE may not equal TATE (Cole and Stuart, 2010)
  - If distribution of treatment effect modifier (TEM) differs in trial and target population
- Note: Similar issues arising in debates about non-probability samples in survey world



# Methods to Improve Generalizability

## Study Design

- Random sampling (great, but rare)
- Purposive sampling (not formally representative)
- Practical clinical trials (potentially useful, but expensive and still lacks formal representativeness)
  
- Main idea: select subjects for trial in a particular way



# Methods to Improve Generalizability

## Post-Hoc Analysis

- Weighting trial sample by odds of trial participation to resemble target population
  - Similar to propensity score ATT weighting in non-experimental studies
- Model outcome as (flexible) function of observed covariates in trial and predict outcome in target population
  - BART: Bayesian Additive Regression Trees (Hill, 2010; Kern et al., 2016), TMLE (Rudolph et al., 2014)
- Doubly robust methods, fitting models for both the outcome and the probability of trial participation (Dahabreh et al., 2018)





# Methods to Improve Generalizability

## Implementation

- Methods require full data sets of TEMs for individuals in both trial *and* target population (not just summary statistics!)
- Software in the works for easy implementation (Ackerman et al., 2019)

	S (trial membership)	Y (outcome)	A (treatment)	X <sub>1</sub> (covariate 1)	X <sub>2</sub> (covariate 2)	...	X <sub>p</sub> (covariate p)
Trial data set	1	✓	✓	✓	✓		✓
Population data set	0	×	×	✓	✓		✓



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# Assumptions\*

- 1 All members of target population have nonzero probability of trial selection
- 2 Range of effect modifiers in target population are covered by respective ranges in the trial
- 3 Treatment assignment is independent of sample selection, as well as potential outcomes, given pre-treatment covariates
- 4 No unmeasured variables associated with sample selection and treatment effect

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\*These are required for the standard weighting and outcome model based



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## What if

- there is an treatment effect modifier (TEM) observed in the trial but we don't have data on it from the target population?
- we are concerned there might be TEMs that are not even observed in the trial (fully unobserved)?



# Issue of Unobserved TEMs

- Unobserved TEMs likely very common in real data scenarios
- In real data, often few covariates observed consistently between trial and population data (Stuart and Rhodes, 2017)
- Today will focus on parametric sensitivity analysis strategy; bounding another possibility (Chan, 2018)



## Sensitivity Analyses for Generalizing RCT Findings to Population: Two Cases

- TEM observed in the trial but unobserved in the target population (*V case*)
- Any potential TEMs unobserved in both the trial *and* the target population (*U case*)

*analogous to sensitivity analyses for unobserved confounding in non-experimental settings (Rosenbaum and Rubin, 1983; VanderWeele, 2011)*



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# Notation

$A$ : treatment (0,1) randomized in trial

$Y$ : outcome (observed only in the trial)

$Y(a)$ : potential outcome under treatment  $a$

$S$ : sample membership

$$S_i = \begin{cases} 1 & \text{if individual } i \text{ is in trial} \\ 0 & \text{if individual } i \text{ is in target population} \end{cases}$$

$$SATE = E[Y(1) - Y(0) | S = 1]$$

$$TATE = E[Y(1) - Y(0) | S = 0]$$



# Notation

$X$ : non-effect modifying covariates

$Z$ : effect modifier, fully observed in trial and population

$V$ : partially unobserved effect modifier (observed in trial, not population)

$U$ : fully unobserved effect modifier



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## For a partially unobserved effect modifier ( $V$ case)

Assume the following outcome model:

$$E[Y_i] = \beta_0 + \beta_a A_i + \beta_{za} Z_i A_i + \beta_{va} V_i A_i + f_{xzv}(X_i, Z_i, V_i)$$

Therefore,

$$\text{TATE} = \beta_a + \beta_{za} E[Z | S = 0] + \beta_{va} E[V | S = 0]$$



# For a partially unobserved effect modifier ( $V$ case)

$$\text{TATE} = \beta_a + \beta_{za}E[Z | S = 0] + \beta_{va}E[V | S = 0]$$

## 1 Outcome-model-based sensitivity analysis

- i. estimate  $\beta_a, \beta_{za}, \beta_{va}$  using trial data
- ii. obtain  $E[Z|S = 0]$  and specify range for  $E[V | S = 0]$
- iii. combine

## 2 Weighted-outcome-model-based sensitivity analysis

- . weight trial sample to resemble target population w.r.t.  $X, Z$
- i. estimate  $\beta_a, \beta_{za}, \beta_{va}$  using trial data
- ii. obtain  $E[Z|S = 0]$  and specify range for  $E[V | S = 0]$
- iii. combine

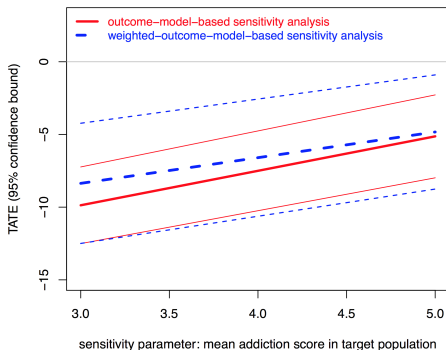


## Example of a $V$ case

Smoking cessation intervention for heavy smokers among attendants of alcohol/substance abuse treatment: SATE = 10 fewer cigarettes per day

- $Z$ : being African-American, baseline daily number of cigarettes
- $V$ : baseline addiction score;  $E[V | S = 1] = 4.05$

Target pop: people who seek alcohol/substance treatment who smoke heavily



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## For fully unobserved effect modification ( $U$ case)

$U$ : the *remaining composite effect modifier*

- captures all unobserved factors that modify treatment effects
- independent of observed covariates and effect modifiers

Unlike the  $V$  case, we cannot use

$$\text{TATE} = \beta_a + \beta_{za}[Z \mid S = 0] + \beta_{ua}[U \mid S = 0]$$

for sensitivity analysis





## For fully unobserved effect modification ( $U$ case)

We hope to use a bias formula instead:

$$\begin{aligned} \text{TATE} = \text{SATE} &+ \beta_{za}\{[Z | S = 0] - [Z | S = 1]\} + \\ &+ \beta_{ua}\{[U | S = 0] - [U | S = 1]\} \end{aligned}$$

Let  $\Delta_u = E[U|S = 0] - E[U|S = 1]$

$$\text{TATE} = \text{SATE} + \beta_{za}\{E[Z | S = 0] - E[Z | S = 1]\} + \beta_{ua}\Delta_u$$

$$\begin{aligned} \text{TATE} &= \text{wtd.ATE} + \beta_{za} \left\{ E[Z | S = 0] - \frac{\sum W_i(S_i = 1)Z_i}{\sum W_i(S_i = 1)} \right\} + \beta_{ua}\Delta_u \\ &\approx \text{wtd.ATE} + \beta_{ua}\Delta_u \end{aligned}$$



# For fully unobserved effect modification ( $U$ case)

## 1 Bias-formula-based sensitivity analysis

$$\text{TATE} = \text{SATE} + \beta_{za}\{E[Z | S = 0] - E[Z | S = 1]\} + \beta_{ua}\Delta_u$$

- i. estimate SATE,  $E[Z | S = 1]$  and  $\beta_{za}$  using trial data
- ii. obtain estimate for  $E[Z | S = 0]$ , specify ranges for  $\beta_{ua}$  and  $\Delta_u$
- iii. combine

## 2 Weighting-plus-bias-formula-based sensitivity analysis

$$\text{TATE} = \text{wtd.ATE} + \beta_{za} \left\{ E[Z | S = 0] - \frac{\sum W_i(S_i = 1)Z_i}{\sum W_i(S_i = 1)} \right\} + \beta_{ua}\Delta_u$$

- . weight trial sample to resemble target population w.r.t.  $X, Z$
- i. estimate wtd.SATE,  $\frac{\sum W_i(S_i=1)Z_i}{\sum W_i(S_i=1)}$  and  $\beta_{za}$  using trial data
  - ii. obtain estimate for  $E[Z | S = 0]$ , specify ranges for  $\beta_{ua}$  and  $\Delta_u$
  - iii. combine



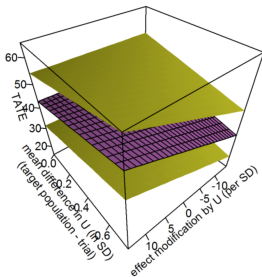
## Example of a $U$ case (data artificially altered)

Trial comparing a new antiretroviral therapy regimen to an old one:  
SATE = increase CD4 count by 36 cells/ml at 2 months post treatment

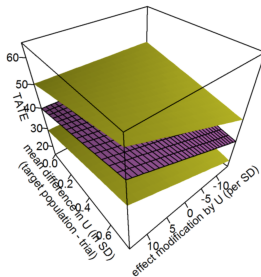
- $Z$ : being White and without severe immune suppression (interaction term coef  $\approx -15$ ), age (interaction term coef  $\approx 11$  per SD)
- concerned about  $U$ : specify  $\Delta_u = (0, 0.7)$  and  $\beta_{ua} = (-15, 15)$

Target population: people with HIV in the US

bias-formula-based method



weighting-plus-bias-formula-based method



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## Extension to Binary Outcomes ( $V$ case)

What if the primary outcome in the trial is not continuous, but binary?

Let  $P_a$  = probability of outcome occurring under treatment "a"

$$TATE = E \left[ \frac{P_{treat}/(1 - P_{treat})}{P_{control}/(1 - P_{control})} \mid S = 0 \right]$$

Assume the following outcome model:

$$\text{logit}(P(Y_i = 1)) = \beta_0 + \beta_a A_i + \beta_{za} Z_i A_i + \beta_{va} V_i A_i + f_{xzv}(X_i, Z_i, V_i)$$

Therefore,

$$TATE = E[\exp\{\beta_a + \beta_{za}Z + \beta_{va}V\} \mid S = 0]$$



## Extension to Binary Outcomes ( $V$ case)

but  $V$  is unmeasured in the population, so  $E[e^{\beta_{va}V} | S = 0]$  is unknown. Using MGFs and assuming  $Z$  and  $V$  to be Normal,

$$\text{TATE} = \lambda \times \exp \left\{ \beta_{va} E[V | S = 0] + \frac{1}{2} \beta_{va}^2 \text{var}(V | S = 0) \right\}$$

where  $\lambda = \exp \left\{ \beta_a + \beta_{za} E[Z | S = 0] + \frac{1}{2} \beta_{za}^2 \text{var}(Z | S = 0) \right\}$

Can then perform same sensitivity analyses for  $V$  case (outcome model or weighting)

- Must make additional assumptions about the distribution of  $V$  and  $Z$  in the population
- Can hold one unknown fixed, or vary both together



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# Conclusion

- Methods exist to improve upon generalizability of RCTs when drawing population-level inferences
- Population data availability is a primary limiting factor for satisfying assumption of no unobserved TEMs
- Proposed sensitivity analyses for generalizing average treatment effects when TEMs are both partially and fully unobserved
- While sensitivity analyses are helpful when some variables are not observed, they are not a panacea





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