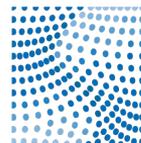


Combining Experimental and Observational Data for Causal Inference

Eric and Wendy Schmidt Center
Organisms – Clinical Trials Group
November 27, 2023



MIT EECS



**ERIC AND WENDY
SCHMIDT CENTER**
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I will walk through some mechanics of causal inference and mention some of our related work along the lines

- 🎯 Causal inference
 - objective
 - assumptions
- 🎯 Data
 - experimental and observational
 - strengths and weaknesses
- 🎯 Benchmarking results
 - are conclusions from two studies **compatible**?
- 🎯 Combining observational and experimental data
 - how to leverage their **complementary** strengths?

Causal inference: objective

- X »» baseline characteristics / covariates / features
- $A \in \{0, 1\}$ »» treatment assignment (e.g. placebo vs treatment)
- Y »» **observed** outcome
- Y^0, Y^1 »» **potential** outcomes under treatment $A = 0$ and $A = 1$
→ **only one** of them observed for every patient



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treatment effect (ATE)?

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= ?

$$\mathbf{E} [Y|A=1] - \mathbf{E}[Y|A=0]$$

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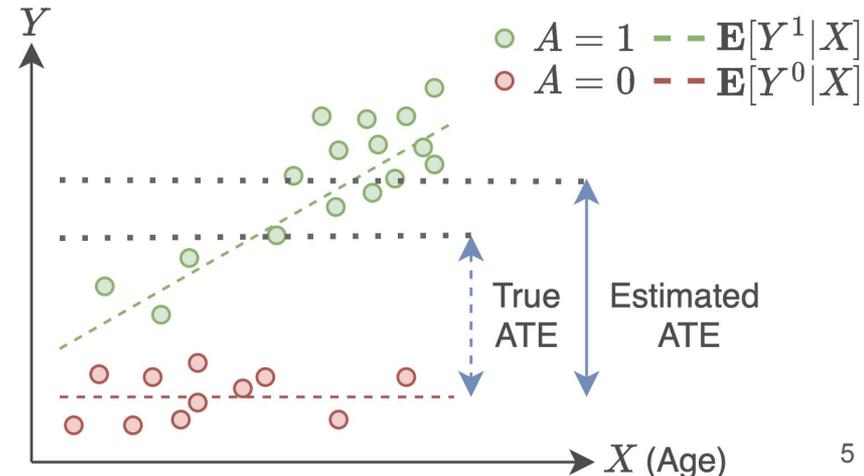
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Not if there is
⇒
confounding



- Assumptions for causal inference
 - $\mathbf{E}[Y^a \mid X] = \mathbf{E}[Y^a \mid X, A=a]$ (ignorability/no unmeasured confounding)
 - › we can attribute differences in outcomes to treatment **after fixing X**
 - $\mathbf{P}(A=a \mid X) > 0$ (positivity)
 - › for any baseline X, we have **some** data from both treatments
 - If $A=a$, then $Y = Y^a$ (consistency)
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$$\mathbf{E}[Y^a] = \mathbf{E}_X[\mathbf{E}[Y^a | X]]$$

Law of total expectation

$$= \mathbf{E}_X[\mathbf{E}[Y^a | X, A = a]]$$

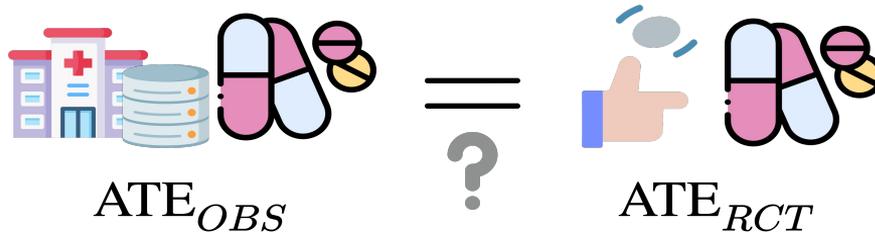
Ignorability + Positivity

$$= \mathbf{E}_X[\mathbf{E}[Y | X, A = a]]$$

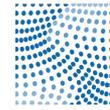
Consistency

- Assumptions for causal inference
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 - If $A=a$, then $Y = Y^a$ (consistency)
- When do these assumptions hold?
 - Always for **experimental data**
 - › RCTs satisfy them by design via randomized treatment assignments
 - › strong **internal validity**
 - Not always with **observational data**
 - › treatment assignments are not random
 - › do we control for all the confounders X ?
 - › how is positivity affected by including more features in X ?

- RCTs are **internally valid**
 - great! we can make causal inference for the **trial population**
- Are they **externally valid**?
 - External validity of randomised controlled trials: “to **whom** do the **results** of this **trial apply**?”, PM Rothwell, *The Lancet*, 2005



- strict exclusion criteria
 - › e.g. people with a history of cardiovascular disease
 - › what if those criteria also affects prognosis?



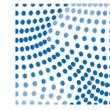
We use observational data to make causal inference when evidence from RCT is **not available** or **limited**

- $S \in \{0, 1\}$ » population index
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- Last week: **How to specify and emulate a target trial from the observational data?**
 - an intricate and laborious process
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- No unmeasured confounding assumption is **unverifiable**

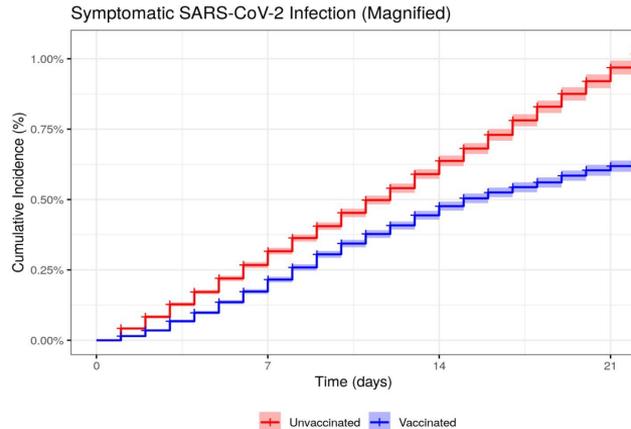


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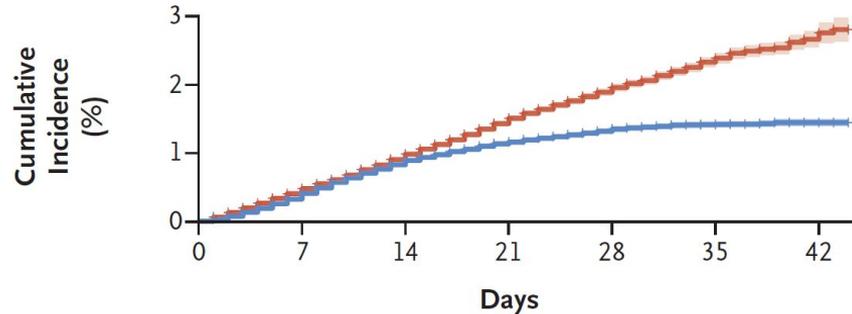
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 - $\mathbf{P}(A = a \mid X, S = 1) > 0$ (positivity)
 - If $A = a$ and $S = 1$, then $Y = Y^a$ (consistency)
- No unmeasured confounding assumption is **unverifiable**
- High dim. X makes propensity scores $\mathbf{P}(A \mid X)$ **extreme**
 - **high variance** in estimations
 - people use standardized weights and clip extreme values (e.g. >0.99)
 - reduces variance but introduces **bias**
 - › rely heavily on “extrapolation”

We use observational data to make causal inference when evidence from RCT is **not available** or **limited**

- We would like to get a sense of whether the **causal assumptions** indeed hold for the **observational study** (recall: internal validity)



A Documented SARS-CoV-2 Infection

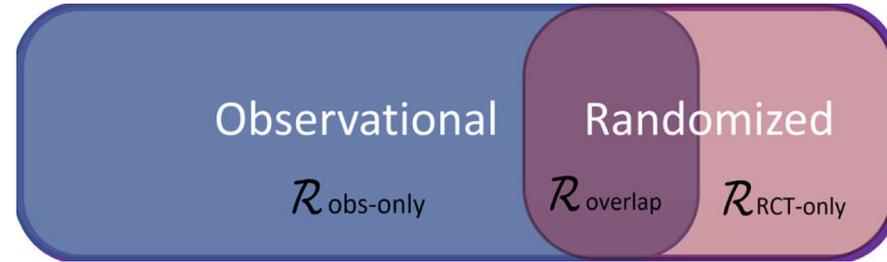


[1] Dagan+ BNT162b2 mRNA Covid-19 vaccine in a nationwide mass vaccination setting, *New England Journal of Medicine* (2021).

[2] @_MiguelHernan, URL: <https://twitter.com/MiguelHernan/status/1364700315044438023>, Twitter Thread.

Benchmarking observational data

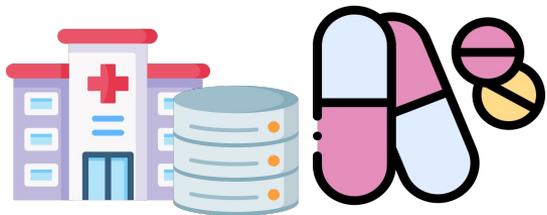
Benchmark an observational study to an RCT in “**overlap region**” before using it for “**no RCT region**”



Goal: Use experimental data as a form of validation — *falsify* assumptions of internal and external validity in observational studies

Compare average effects

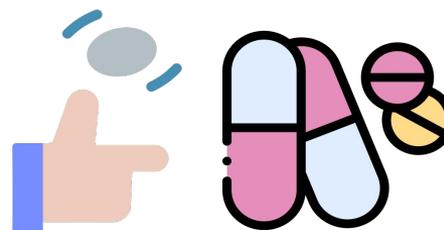
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ATE_{OBS}



No explanation for rejection



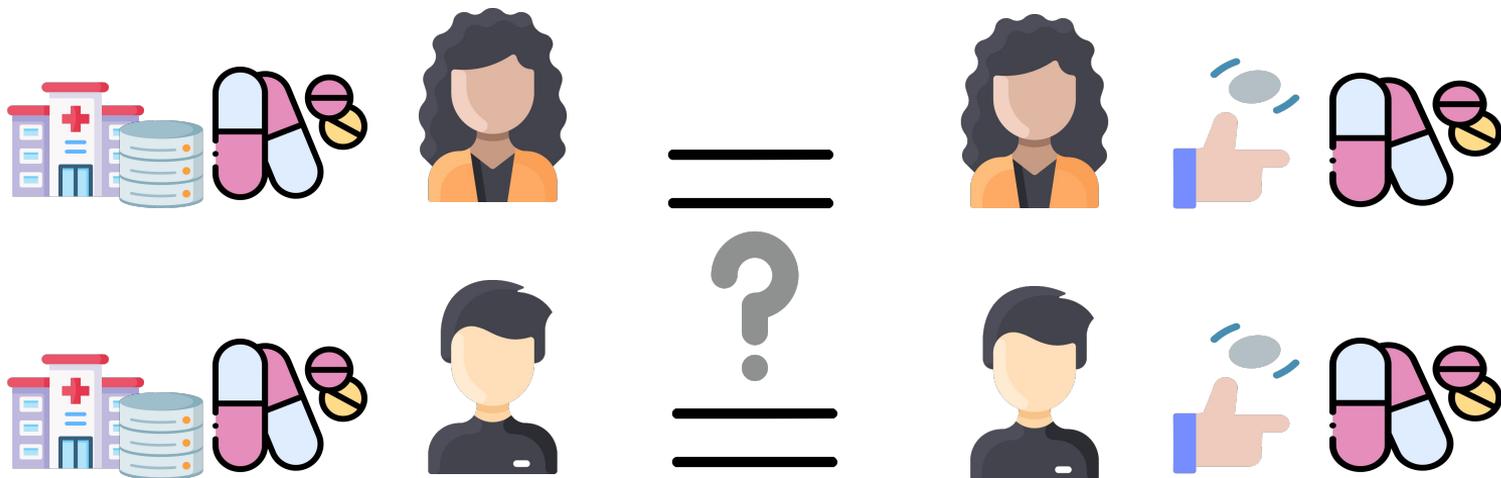
ATE_{RCT}



Could lead to false negatives

Compare group-level average effects

Goal: Use experimental data as a form of validation — *falsify* assumptions of internal and external validity in observational studies



Need a-priori specification of the subgroups;
would be nice to find these automatically

Compare conditional average effects

Goal: Use experimental data as a form of validation — *falsify* assumptions of internal and external validity in observational studies

S	X_0	...	X_1	A	Y	ψ_0	ψ_1	ψ
0	1		-.5	0	3	-20	0	20
0	1		.2	1	7	37	0	-37
⋮	⋮			⋮	⋮	⋮	⋮	
1	1		.3	1	9	0	46	46
1	1		-.4	0	11	0	-12	-12



CATE signal ψ_1 from
OBS ($S=1$)



CATE signal ψ_0 from
RCT ($S=0$)

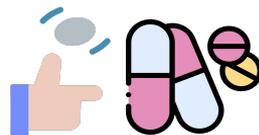
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CATE signal ψ_1 from
OBS ($S=1$)



CATE signal ψ_0 from
RCT ($S=0$)

(CATE in RCT)
$$E [Y^1 - Y^0 \mid X, S=1]$$

(CATE in OBS)
$$E [Y^1 - Y^0 \mid X, S=0]$$

Compare conditional average effects

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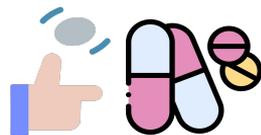


$$E[\psi_1 | X] = E[Y^1 - Y^0 | X, S=1]$$

(CATE in RCT)

CATE signal ψ_1 from
OBS ($S=1$)

Internal Validity of
RCT and OBS



$$E[\psi_0 | X] = E[Y^1 - Y^0 | X, S=0]$$

(CATE in OBS)

CATE signal ψ_0 from
RCT ($S=0$)

Compare conditional average effects

Goal: Use experimental data as a form of validation — *falsify* assumptions of internal and external validity in observational studies

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$$\mathbf{E} [\psi_1 | X] = \mathbf{E} [Y^1 - Y^0 | X, S=1]$$

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CATE signal ψ_1 from
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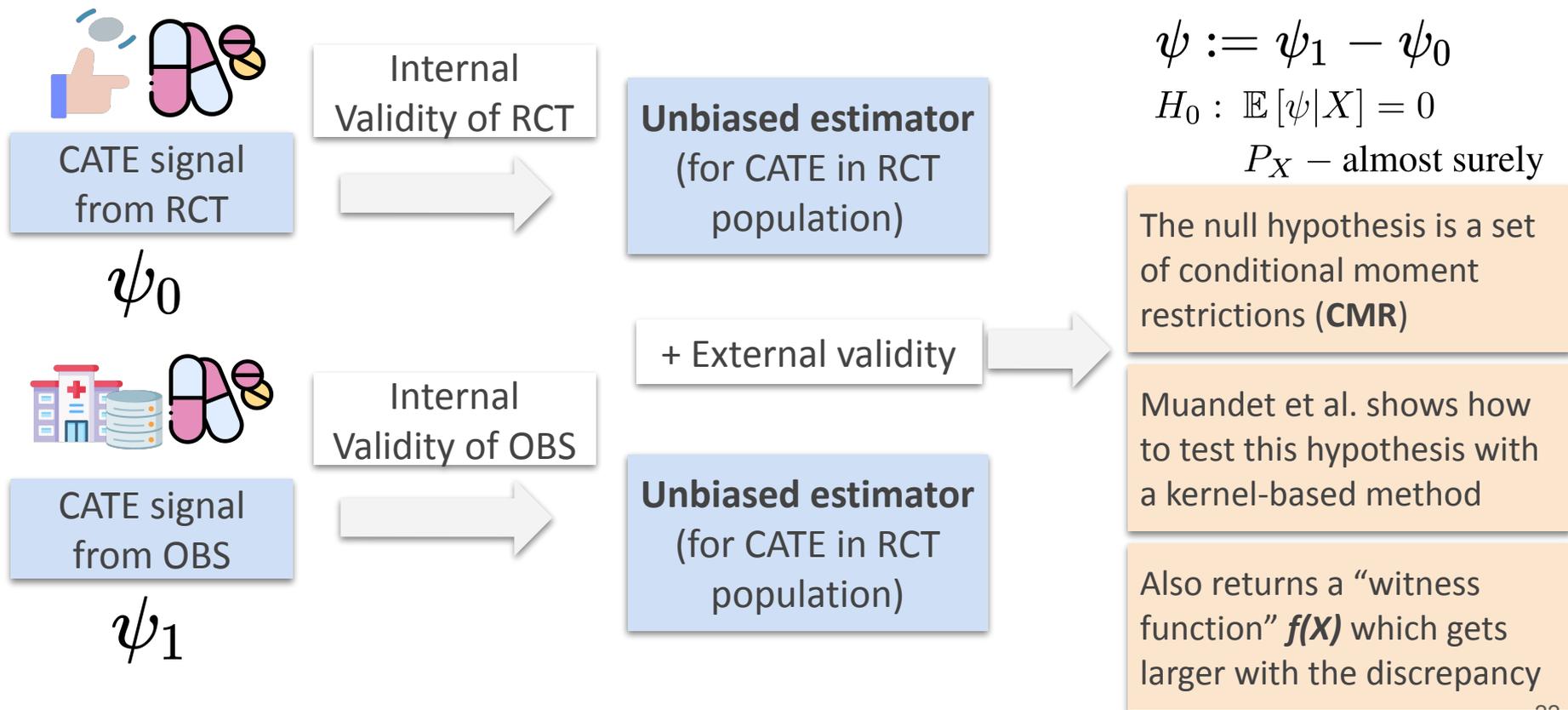
(CATE in OBS)

CATE signal ψ_0 from
RCT ($S=0$)

+ External validity

H₀: $\mathbf{E} [\psi | X] = 0$, $\psi := \psi_1 - \psi_0$
Our null hypothesis to test

Compare conditional average effects



$$\psi := \psi_1 - \psi_0$$

$$H_0 : \mathbb{E}[\psi|X] = 0$$

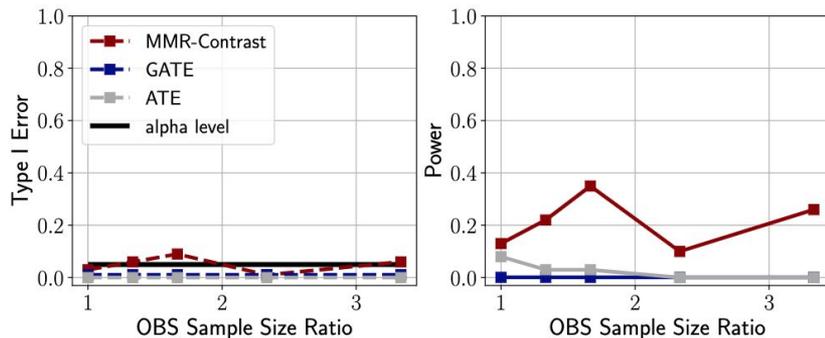
P_X – almost surely

The null hypothesis is a set of conditional moment restrictions (**CMR**)

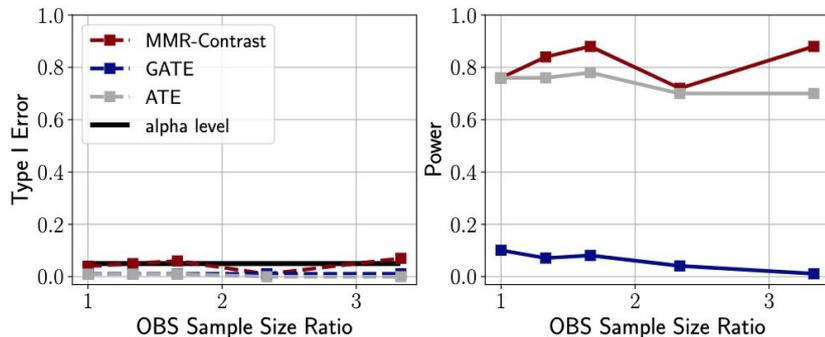
Muandet et al. shows how to test this hypothesis with a kernel-based method

Also returns a “witness function” $f(\mathbf{x})$ which gets larger with the discrepancy

IHDP and WHI experiments



(a) Low confounder strength ($\max(\gamma) = 1.$). (left) no unobserved confounders; (right): one confounder concealed



(b) High confounder strength ($\max(\gamma) = 2.75$). (left) no unobserved confounders; (right): one confounder concealed

<i>Selection Bias</i>	MMR-Contrast	ATE	GATE
$p = 0$	0.29	0.32	0.17
$p = 0.05$	0.67	0.58	0.40
$p = 0.10$	0.94	0.88	0.67
$p = 0.15$	1.0	0.98	0.91

Table 1: Rejection rate when introducing different amounts of selection bias into the observational data in WHI study. p stands for the strength of selection introduced in the the data (refer to Section 5 for details).

Compare conditional average effects



- **Censored** case?
 - we do not observe the outcome in some patients
 - › there is **censoring time**
 - › loss-to-follow-up, end of study, drop-out...
 - internal validity **not enough** for causal inference
 - › need **new assumptions** on censoring mechanism

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 - › need **new assumptions** on censoring mechanism
- **Independent censoring**
 - censoring time (C) is independent of time-to-event outcome (Y)
- **Global censoring**
 - they are dependent, but the same way in RCT and OBS
 - › e.g., similar drop-out due to adverse side effects

Compare conditional average effects



- Naively handling censored data does not work
 - easily bias the results
- We develop new signals ψ_0 and ψ_1
- Under independent censoring we can recover internal validity
 - $\mathbf{E}[\psi_1 | X] = \mathbf{E}[Y^1 - Y^0 | X, S=1]$ (can estimate CATE despite censoring)
 - › business as usual: test $\mathbf{H}_0: \mathbf{E}[\psi | X] = 0$
- Global censoring is more interesting, cannot estimate CATE
 - but still, internal + external validity implies: $\mathbf{E}[\psi | X] = 0$

Compare conditional average effects

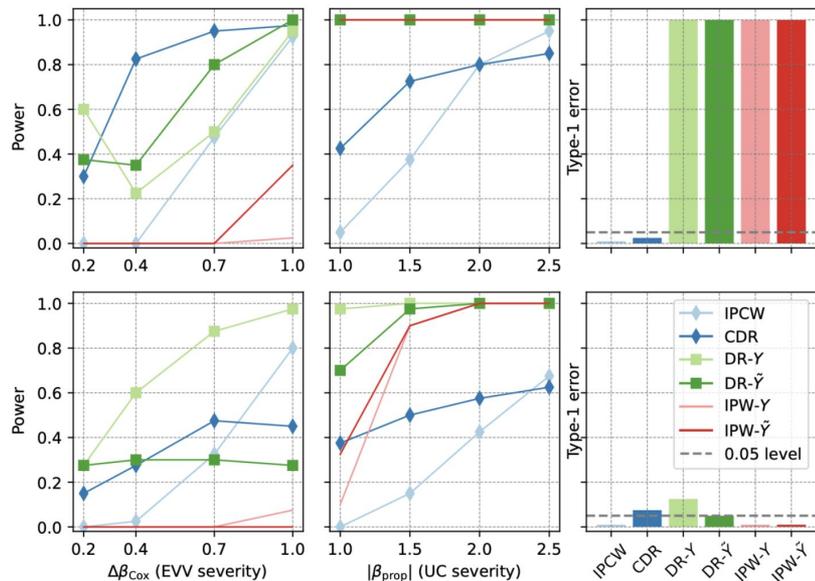
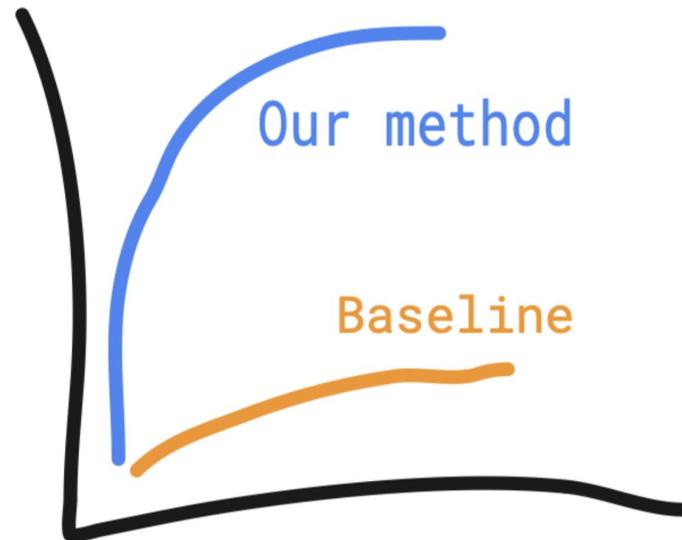


Figure 2: *Top row:* Conditionally independent censoring results. *Bottom row:* Global censoring results. EVV = External validity violation (A2). UC = Unobserved confounding (A1). OS size is $n_1 = 2955$.

(Under review)



Combining RCT + Observational Data



Science

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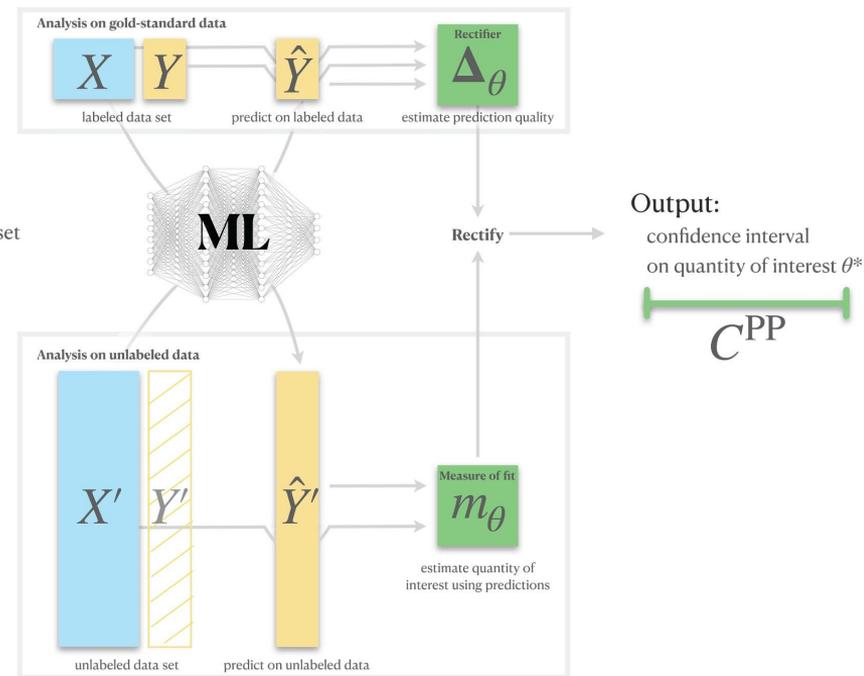
RESEARCH ARTICLE | MACHINE LEARNING

Prediction-powered inference

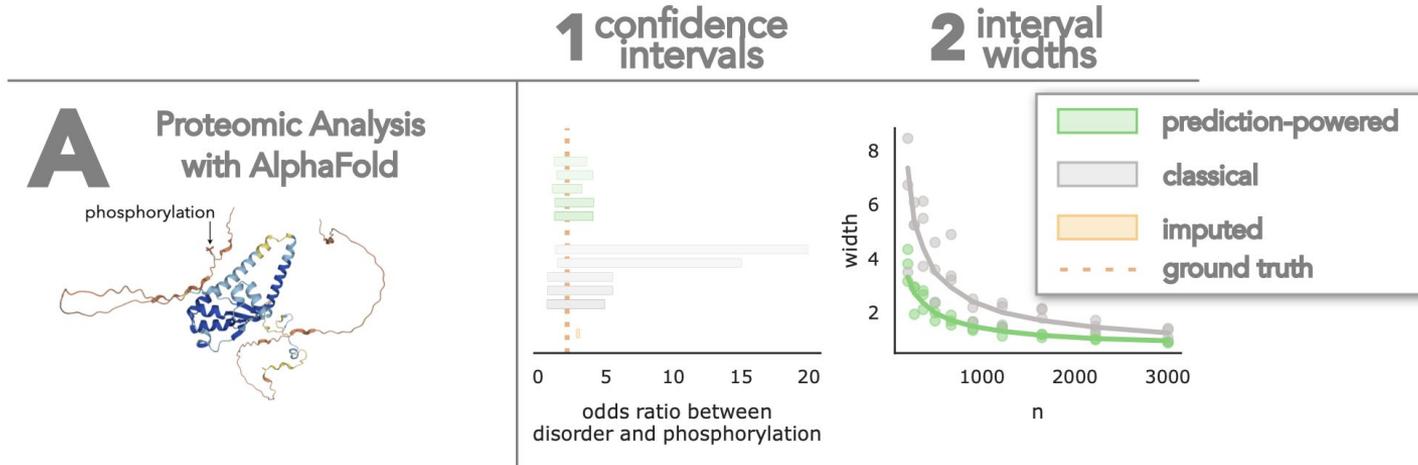
ANASTASIOS N. ANGELOPOULOS, STEPHEN BATES, CLARA FANNJIANG, MICHAEL I. JORDAN, AND TIJANA ZRNIC

- Gold-standard (**labeled**) small data
- **Unlabeled** huge data
- A predictive ML model f
 - cannot use it at face value
- Measure its **error** on labeled data
- Impute the unlabeled data
 - correct for the error learned
 - **interval imputations** rather than point imputations

Inputs:
gold-standard data set
unlabeled data set
ML algorithm



Combining RCT + Observational Data



- Odds ratio (**OR**) between post-translational modifications (**PTM**) and intrinsically disordered regions (**IDR**)
- Use AlphaFold to predict protein structures
- Quantify its error on a small labeled (**IDR**) dataset (**for OR**)
- Correct for the error in the imputations for the unlabeled proteins



(Work in progress)

- Recall the problem with RCTs: **limited external validity**
 - consider trial ($S = 0$) and target ($S = 2$) populations
 - $\mathbf{E}[Y^1 | S = 0] \neq \mathbf{E}[Y^1 | S = 2]$
 - confounding! trial participants are systematically different than our target population in prognostic factors

$$\begin{aligned}\mathbf{E}[Y^1 | S = 2] &= \mathbf{E}_{P_{X|S=2}}[\mathbf{E}[Y^1 | X, S = 2]] \\ &= \mathbf{E}_{P_{X|S=2}}[\mathbf{E}[Y^1 | X, S = 0]] \\ &= \mathbf{E}_{P_{X|S=2}}[\mathbf{E}[Y | X, S = 0, A = 1]]\end{aligned}$$