Generalization of trial's findings toward a target population: choice of a causal measure





Julie Josse. Senior Researcher Inria, Montpellier, France
Lead Inria-Inserm PreMeDICaL team: precision medicine by data integration & causal learning

Presentation Julie Josse: Stat and ML for bio-sciences

Academic background:

- ▷ Engineer & Assistant Prof. Agronomy Univ. Rennes (2007 2015)
- ▷ Visiting researcher + teaching Stanford Univ. (18 months)
- ▷ Professor at Ecole Polytechnique (IP Paris) (2016 2020)
- Senior Researcher at Inria Montpellier (Sept. 2020 -)
 Head of Premedical Inria-Inserm (National health research center) team

Research topics:

- Dimensionality reduction to visualize high dimensional heterogeneous data
- <u>Causal inference</u>: estimating treatment effect, combining RCT & observational data, personalized recommendation
- ▶ Medical collaborations: Critical care, Inst. Gustave Roussy, etc.

Implementations: R community

book R for Stat., Elected member R foundation, Founding of Rforwards to increase minority participation, packages: FactoMineR (4 500 download/day, > 5 million in total, >7000 citations), Rmisstastic, taskviews

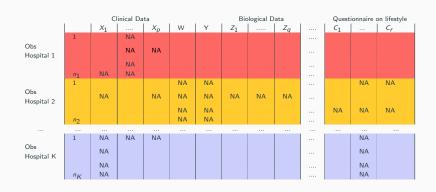
PreMeDICaL research axes

Personalized medicine by integration of different data sources

- ▷ relationship between data sources (relevance of each source?)
- $\, \triangleright \,$ solutions to handle complexe structure of missing values
- confidence in ML/AI algo. (uncertainty quantification conformal)
- federated learning, privacy

- ⇒ Translate research into clinically actionable solutions
- ⇒ Push methodological innovation up to patients, clinicians, regulators
- ⇒ Leverage ML, data, clinical expertise & existing recommendations

Data integration comes with methodological challenges



State-of-the-art ML/causal methods struggle with high dimensional multi-sources data with distributional shifts & missing data (systematic/sporadic)

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Personalized medicine by optimal prescription of treatment

- > causal inference techniques for (dynamic) policy learning
 - \Rightarrow who to treat and when
- ▷ leverage both randomized control trials (RCTs) and observational data
 - ⇒ launch a drug without running RCTs
 - ⇒ rethink evidence needed to bring treatments to the market faster
- ⇒ Translate research into clinically actionable solutions
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- ▶ 4000 new patients/ year

Center	Accident	Age	Sex	Weight	Lactactes	BP	TXA.	Y
Beaujon	fall	54	m	85	NM	180	treated	0
Pitie	gun	26	m	NR	NA	131	untreated	1
Beaujon	moto	63	m	80	3.9	145	treated	1
Pitie	moto	30	W	NR	lmp	107	untreated	0
HEGP	knife	16	m	98	2.5	118	treated	1

[:]

¹www.traumabase.eu - https://www.traumatrix.fr/

- ▶ 40000 patients
- ▶ 40 trauma centers
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 \Rightarrow Explain and Predict hemorrhagic shock given pre-hospital features.

Ex: logistic regression/ random forests with covariates with missing values

Clinical Trial: real-time testing of models in the ambulance via a mobile data collection application (ShockMatrix Google play)

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⇒ **Estimate causal effect**: Administration of the **treatment** *tranexamic acid* (*TXA*), given within 3 hours of the accident, on the **outcome** *28 days intra hospital mortality* for trauma brain patients

TXA decreases mortality for extra-cranial bleeding. Effect for intra-cranial bleeding? (detected by CT scan). TXA is one of the first treatments given

- ▶ 40000 patients
- ▶ 250 continuous and categorical variables
- ▶ 40 trauma centers
- ⊳ 4000 new patients/ year

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Causal inference with covariates with missing values (implemented in the R package grf). 2020. Mayer, I., Wager, S. & J.J. Doubly robust treatment effect estimation with incomplete confounders. *Annals Of Applied Statistics*.

CRASH3

- > 9000 individuals develp. countries
- No evidence for a TXA effect (positive effect for moderate injured patients)

Traumabase

- ▷ Observational sample
- \triangleright 8200 patients with brain trauma
- Slightly negative effect of TXA/ no evidence

²Lodi, Hernán et al. (2019). Effect Estimates in Randomized Trials and Observational Studies: Comparing Apples With Apples. *Am J Epidemiol*.

³ (2019). Effects of tranexamic acid on death in patients with acute trauma. brain injury. *Lancet*.

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- ▶ Multi-centric RCT 29 countries
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Is this expected? Is there a paradox?

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Is this expected? Is there a paradox?

- Treatment and outcome are not exactly the same²
- > Traumabase may suffer from unobserved confounding missing values
- Populations are different



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Leverage both RCT and observational data

RCT

- + No confounding
- Trial sample different from the population eligible for treatment

(big) Observational data

- Confounding
- Representative of the target population

We could use both to 4 . . .

- ▷validate observational methods, correct confounding bias

⁴Colnet, J.J. (2022). Causal inference for combining RCT & obs. studies. *Statistical Science*.

⁵Elias Bareinboim & Judea Pearl. (2016). Causal inference & the data-fusion problem. *PNAS*.

⁶Dahabreh, Haneuse, Robins, Robertson, Buchanan, Stuart, Hernán. (2021). Study Designs for Extending Causal Inferences From a RCT to a Target Population *American J. of Epidemiology*.

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The FDA has greenlighted the usage of the drug *Ibrance* to men with breast cancer, though clinical trials were performed only on women.

ightarrow Reduce drug approval times and costs for patients who could benefit

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Generalization task from a RCT to a target population

Two data sources:

- \triangleright A trial of size *n* with $p_R(x)$ the probability of observing individual with X = x,
- ▷ A sample of the target population of interest for e.g. a national cohort (resp. m and $p_T(x)$).



Generalization task from a RCT to a target population

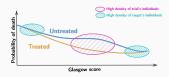
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Covariates distribution not the same in the RCT & target pop:

$$p_{\mathsf{R}}(x) \neq p_{\mathsf{T}}(x) \Rightarrow \underbrace{\tau_{\mathsf{R}} := \mathbb{E}_{\mathsf{R}}[Y(1) - Y(0)]}_{\mathsf{ATE in the RCT}} \neq \underbrace{\mathbb{E}_{\mathsf{T}}[Y(1) - Y(0)] := \tau_{\mathsf{T}}}_{\mathsf{Target ATE}}$$



Assumptions for ATE identifiability in generalization

Transportability (Ignorability on trial participation)⁷

$$\forall w \in \{0,1\}$$
 $\mathbb{E}_{\mathbb{R}}[Y(w) \mid X] = \mathbb{E}_{\mathbb{T}}[Y(w) \mid X = x]$

Corresponds to shifted prognostic variables

 $^{^7}$ Equivalent formulation with sampling mechanism S (S=1 trial eligibility & willingness to participate) in non-nested design, $\{Y(1),\,Y(0)\}\,\perp\,S\mid X$

 $^{^8}$ If this is too strong, we could generalize on a different target population: the target population for which eligibility criteria of the trial are ensured

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Overlap assumption⁸

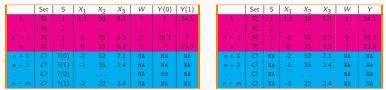
$$\forall x \in \mathbb{X}, p_{\mathbb{R}}(x) > 0 \text{ and } \operatorname{supp}(P_{\mathcal{T}}(X)) \subset \operatorname{supp}(P_{\mathcal{R}}(X))$$

The observational covariates support is included in the RCT's support. Every individual in the target population could have been selected into the trial

 $^{^{7}}$ Equivalent formulation with sampling mechanism S (S=1 trial eligibility & willingness to participate) in non-nested design, $\{Y(1),Y(0)\} \perp S \mid X$

 $^{^{8}}$ If this is too strong, we could generalize on a different target population: the target population for which eligibility criteria of the trial are ensured

Generalization of conditional outcome: identifiability

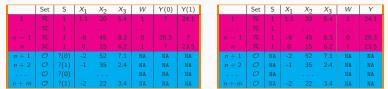


Data with observed treatment \ensuremath{W} and outcome \ensuremath{Y} only in the RCT.

Average Treatment Effect:
$$\tau_{\mathsf{T}} = \mathbb{E}_{\mathsf{T}}[Y_i(1) - Y_i(0)], \forall w \in \{0, 1\}$$

$$\begin{split} \mathbb{E}_{\mathsf{T}}\left[Y(1)\right] &= \mathbb{E}_{\mathsf{T}}\left[\mathbb{E}_{\mathsf{T}}\left[Y(1)\mid X\right]\right] \text{ Law of total expectation} \\ &= \mathbb{E}_{\mathsf{T}}\left[\mathbb{E}_{\mathsf{R}}\left[Y(1)\mid X\right]\right] \text{ Ignorability} \\ &= \mathbb{E}_{\mathsf{T}}\left[\mathbb{E}_{\mathsf{R}}[Y(1)\mid X=x,W=1]\right] \text{ Random treatment} \\ &= \mathbb{E}_{\mathsf{T}}\left[\mathbb{E}_{\mathsf{R}}\left[Y\mid X=x,W=1\right]\right] \text{ Consistency } Y=Y(1)W+(1-W)Y(0) \end{split}$$

Generalization of conditional outcome: identifiability



Data with observed treatment W and outcome Y only in the RCT.

Average Treatment Effect: $\tau_{\scriptscriptstyle T} = \mathbb{E}_{\scriptscriptstyle T}[Y_i(1) - Y_i(0)], \forall w \in \{0,1\}$

$$\mathbb{E}_{\mathsf{T}}\left[Y(1)\right] = \mathbb{E}_{\mathsf{T}}\left[\mathbb{E}_{\mathsf{T}}\left[Y(1) \mid X\right]\right] \text{ Law of total expectation}$$

$$= \mathbb{E}_{\mathsf{T}}\left[\mathbb{E}_{\mathsf{R}}\left[Y(1) \mid X\right]\right] \text{ Ignorability}$$

$$= \mathbb{E}_{\scriptscriptstyle\mathsf{T}}\left[\mathbb{E}_{\scriptscriptstyle\mathsf{R}}[Y(1)\mid X=x,W=1]
ight]$$
 Random treatment

$$= \mathbb{E}_{\mathsf{T}} \underbrace{\left[\mathbb{E}_{\mathsf{R}}[Y \mid X = x, W = 1]\right]}_{\mathsf{Consistency}} \mathsf{Consistency} Y = Y(1)W + (1 - W)Y(0)$$

Regression adjustment - plug-in gformula

$$\hat{ au}_{g,n,m} = rac{1}{m} \sum_{i \in \mathcal{T}} \left(\hat{\mu}_{1,n}(X_i) - \hat{\mu}_{0,n}(X_i)
ight)$$

Plug-in gformula: difference between conditional mean

Plug-in gformula

$$\hat{\tau}_{g,n,m} = \frac{1}{m} \sum_{i=n+1}^{n+m} (\hat{\mu}_{1,n}(X_i) - \hat{\mu}_{0,n}(X_i)),$$

$$\mu_w(x) = \mathbb{E}_{R}[Y \mid X = x, W = w]$$

			Covariates			Treat	Outcomes
	Set	S	X_1	X_2	X_3	W	Y
1	\mathcal{R}	1	1.1		9.4	1	
n + 1 n + 2	0	?	-1	35	7.1		
n+2	0	?	-2	52	2.4		
	0	?					
n + m	0	?	-2	22	3.4		

• Fit two models of the outcome (Y) on covariates (X) among trial participants (\mathcal{R}) for treated and for control to get $\widehat{\mu}_{1,n}$ & $\widehat{\mu}_{0,n}$

Plug-in gformula: difference between conditional mean

Plug-in gformula

$$\hat{\tau}_{g,n,m} = \frac{1}{m} \sum_{i=n+1}^{n+m} (\hat{\mu}_{1,n}(X_i) - \hat{\mu}_{0,n}(X_i)),$$

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			Covariates		Treat	Outcomes		
	Set	5	X_1	X_2	X_3	W	Y(0)	Y(1)
1	\mathcal{R}	1	1.1		9.4	1		24.1
n+1	0	?	-1	35	7.1		$\hat{\mu}_0(X_{n+1})$	$\hat{\mu}_1(X_{n+1})$
n+2	0	?	-2	52	2.4		$\hat{\mu}_0(X_{n+2})$	$\begin{array}{c} \hat{\mu}_1(X_{n+1}) \\ \hat{\mu}_1(X_{n+2}) \end{array}$
	0	?						
n + m	0	?	-2	22	3.4		$\hat{\mu}_0(X_{n+m})$	$\hat{\mu}_1(X_{n+m})$

- Fit two models of the outcome (Y) on covariates (X) among trial participants (\mathcal{R}) for treated and for control to get $\widehat{\mu}_{1,n}$ & $\widehat{\mu}_{0,n}$
- Apply these models to the covariates in the target pop, i.e., marginalize over the covariate distribution of the target pop, gives the expected outcomes
- Compute the differences between the expected outcomes on the target population $\widehat{\mu}_{1,n}(\cdot)$ $\widehat{\mu}_{0,n}(\cdot)$

Assumptions for identifiability with fewer covariates

Transportability (Ignorability on trial participation)⁹

$$\forall w \in \{0,1\}$$
 $\mathbb{E}_{\mathbb{R}}[Y(w) \mid X] = \mathbb{E}_{\mathbb{T}}[Y(w) \mid X = x]$

Corresponds to shifted prognostic variables

 $^{^9}$ Equivalent formulation with sampling mechanism S (S=1 trial eligibility & willingness to participate) in non-nested design, $(Y(1), Y(0)) \perp S \mid X$

 $^{^{10}}$ Equivalent formulation with sampling mechanism S: $(Y(1)-Y(0)) \perp \!\!\! \perp S \mid X$

Assumptions for identifiability with fewer covariates

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 $\mathbb{E}_{\mathbb{R}}[Y(w) \mid X] = \mathbb{E}_{\mathbb{T}}[Y(w) \mid X = x]$

Corresponds to shifted prognostic variables

Transportability of the CATE¹⁰

$$\underbrace{\tau_R(X)}_{\mathbb{E}_R[Y(1)-Y(0)|X]} = \underbrace{\tau_T(X)}_{\mathbb{E}_T[Y(1)-Y(0)|X]}$$

Corresponds to shifted treatment effect modifiers

⁹ Equivalent formulation with sampling mechanism S (S=1 trial eligibility & willingness to participate) in non-nested design, $(Y(1), Y(0)) \perp S \mid X$

Equivalent formulation with sampling mechanism S: $(Y(1) - Y(0)) \perp S \mid X$

Identifiability and estimation for generalization: weighting

Generalization of local effects¹¹

$$\begin{split} \tau_{\mathsf{T}} &= \mathbb{E}_{\mathsf{T}}[Y_i(1) - Y_i(0)] = \mathbb{E}_{\mathsf{T}}[\mathbb{E}_{\mathsf{T}}[Y_i(1) - Y_i(0)|X]] \\ &= \mathbb{E}_{\mathsf{T}}\left[\tau_{\mathsf{T}}(X)\right] = \mathbb{E}_{\mathsf{T}}\left[\tau_{\mathsf{R}}(X)\right] \text{ Transportability CATE} \\ &= \mathbb{E}_{\mathsf{R}}\left[\frac{p_{\mathsf{T}}(X)}{p_{\mathsf{R}}(X)}\tau_{\mathsf{R}}(X)\right] \end{split}$$

IPSW: inverse propensity sampling weighting¹²

$$\hat{\tau}_{IPSW,n,m} = \frac{1}{n} \sum_{i \in \mathcal{R}} \quad \frac{\hat{\rho}_{\mathsf{T}}(X_i)}{\hat{\rho}_{\mathsf{R}}(X_i)} \quad \left(\frac{WY}{e_{\mathsf{R}}(x)} - \frac{(1-W)Y}{1-e_{\mathsf{R}}(x)}\right),$$

$$e_R(x) = P(W = 1 \mid X = x) (= \pi = 0.5.)$$

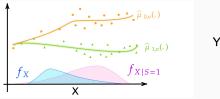
Re-weight, so that the trial follows the target sample's distribution: if proba to be in trial when old is small, then up-weight old in trial

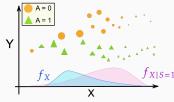
- ▶ Re-weighting can be found in the 2000's (standardization, Rothman & Greenland)
- > But the idea of relying on an external representative sample is recent

¹¹When the measure is collapsible

¹²Cole & Stuart. (2010). Generalizing evidence from RCT to target pop.. American J. of

Generalization estimators: illustrative schematics





The trial findings $\hat{\tau}_{R,n}$ would over-estimate the target treatment effect τ_T Left: the plug-in G-formula model the response using the RCT observation

Right: IPSW weight the RCT observations

 f_X $(f_{X|S=1})$ density of the target (resp. trial) pop., $\hat{\mu}_{w,n}(\cdot)$ fitted response surface with n trial obs.

Theorem - consistency¹³

Under assumptions, $\hat{\tau}_{\text{IPSW},n,m}$ and $\hat{\tau}_{g,n,m}$ converges toward τ_{T} in L^1 norm,

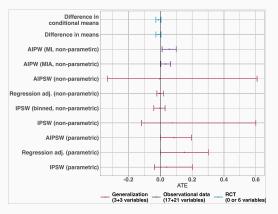
$$\hat{ au}_{\mathsf{IPSW},n,m} \xrightarrow[n,m o \infty]{L^1} au_\mathsf{T}$$

$$\hat{\tau}_{g,n,m} \xrightarrow[n,m\to\infty]{L^1} \tau_{\mathsf{T}}$$

¹³Colnet, J.J et al. 2022. Generalizing a causal effect: sensitivity analysis and missing covariates.
Journal of Causal Inference.

Generalization from Crash 3 trial to the Traumabase

Comparison of trials, observational data, and generalization estimates



Confidence intervals obtained with stratified bootstrap

x-axis: Estimat. of the ATE (\times 100), bootstrap CI

y-axis: Methods - parametric: logistic regression or non parametric forests for nuisances

 $\label{eq:missing walues handled with multiple imputation MI or missing incorportaed in attributes MIA for$

forests

• 1) Shifted effect modifiers not available in Traumabase¹⁴. Missing covariates in one/both sets: **sensitivity analysis**

				Covariate	S	Treat	Outcomes
	Set	5	X_1	X_2	<i>X</i> ₃	W	Y
1	\mathcal{R}	1	1.1	20	NA	1	24.1
	\mathcal{R}	1	-6	45	NA	0	26.3
n	\mathcal{R}	1	0	15	NA	1	23.5
n + 1	0	?	-1	35	7.1		
n + 2	0	?	-2	52	2.4		
	0	?					
n + m	0	?	-2	22	3.4		

Difficult to give sensitivity parameters, semi-parametric model (linear CATE), shift on means, sensitivity plot for one missing variable, etc.

¹⁴Colnet, J.J, et al. 2022. Generalizing a causal effect: sensitivity analysis. J. of Causal Inference.

¹⁵Mayer, J.J. 2021. Generalizing effects with incomplete covariates *Biometrical Journal*.

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• 2) Missing values: Missing values in both RCT and Obs data 15

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Difficult to give sensitivity parameters, semi-parametric model (linear CATE), shift on means, sensitivity plot for one missing variable, etc.

- 2) Missing values: Missing values in both RCT and Obs data¹⁵
- 3) Effect of **finite sample**? Which covariate to include? Would adding prognostic variables reduce the variance as in the classical case?¹⁶

¹⁴Colnet, J.J, et al. 2022. Generalizing a causal effect: sensitivity analysis. *J. of Causal Inference*.

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¹⁶Colnet, **J.J** et al. 2022. Reweighting the RCT for generalization: finite sample analysis and variable selection. *In revision JRSSC*.

Reweighting the RCT: finite sample & asymptotic analysis¹⁷

$$\hat{ au}_{\pi,n,m} = rac{1}{n} \sum_{i \in \mathcal{R}} rac{ \hat{eta}_{\mathsf{T}}(m{X}_i) }{ \hat{eta}_{\mathsf{R}}(m{X}_i) } \quad Y_i \left(rac{W_i}{\pi} - rac{1 - W_i}{1 - \pi}
ight) \,,$$
Estimated with \mathcal{R}

Asymptotic properties - completely estimated

Letting
$$\lim_{n,m\to\infty} m/n = \lambda \in [0,\infty]$$

$$\lim_{n,m\to\infty} \min(n,m) \operatorname{\sf Var}\left[\hat{\tau}_{\pi,n,m}\right] = \min(1,\lambda) \left(\frac{\operatorname{\sf Var}\left[\tau(X)\right]}{\lambda} + V_{\mathsf{so}}\right)$$

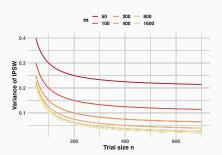
Variance depends on the size of the two data sets, n and m

- ▶ If target >> trial, $m/n \to \infty$, (i.e., $\lambda = \infty$): asymptotic variance = Semi-Oracle's one and depends on the ratio of proba.
- ▷ If trial >> target $m/n \to 0$, (i.e., $\lambda = 0$): asymptotic variance depends on $Var[\tau(X)]$.

¹⁷Colnet, J.J et al. 2022. Reweighting the RCT for generalization: finite sample analysis and variable selection. *In revision JRSSC*.

Reweighting the RCT: finite sample & asymptotic analysis¹⁷

$$\hat{\tau}_{\pi,n,m} = \frac{1}{n} \sum_{i \in \mathcal{R}} \frac{\frac{\hat{\rho}_{\mathsf{T}}(X_i)}{\hat{\rho}_{\mathsf{R}}(X_i)}}{\frac{\hat{\rho}_{\mathsf{R}}(X_i)}{\hat{\rho}_{\mathsf{R}}(X_i)}} \quad Y_i \left(\frac{W_i}{\pi} - \frac{1 - W_i}{1 - \pi}\right) \,,$$
Estimated with \mathcal{R}

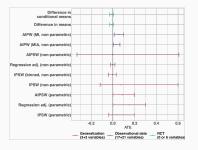


- Adding shifted and independent covariates: loss in variance
- Adding non-shifted treatment effect modifiers: gain in variance

 $^{^{17}}$ Colnet, J.J et al. 2022. Reweighting the RCT for generalization: finite sample analysis and variable selection. *In revision JRSSC*.

Generalization from Crash 3 trial to the Traumabase

Comparison of trials, observational data, and generalization estimates



Confidence intervals obtained with stratified bootstrap

- ▷ 1) Shifted effect modifiers not available in Traumabase¹⁸
- 2) Missing values in both RCT and Obs data¹⁹
- → 3) Effect of finite sample? Which covariate to include? Would adding prognostic variables reduce the variance as in the classical case?²⁰
- > 4) Clinicians rather interested in the ratio than the difference

¹⁸Colnet, J.J. 2022. Generalizing a causal effect: sensitivity analysis, *Journal of Causal Inference*.

¹⁹Mayer, J.J. 2021. Generalizing effects with incomplete covariates *Biometrical Journal*.

²⁰Colnet, J.J. 2022. Reweighting the RCT for generalization: finite sample analysis & variable selection. Submitted

Risk ratio, odds ratio, risk difference

Which causal measure is easier to generalize?



Comparing two average situations

Binary outcome: $\mathbb{P}[Y(w) = 1] = \mathbb{E}[Y(w)]$ and $\mathbb{P}[Y(w) = 0] = 1 - \mathbb{E}[Y(w)]$.

Absolute measures

$$au^{ extsf{RD}} := \mathbb{E}\left[Y(1)\right] - \mathbb{E}\left[Y(0)\right], \qquad au^{ extsf{NNT}} := au_{ extsf{RD}}^{-1}.$$

• Number Needed to Treat (NNT): how many individuals should be treated to observe one individual answering positively to treatment.

Relative measures

$$\tau^{\text{RR}} := \frac{\mathbb{E}[Y(1)]}{\mathbb{E}[Y(0)]}, \quad \tau^{\text{SR}} := \frac{\mathbb{P}[Y(1) = 0]}{\mathbb{P}[Y(0) = 0]} = \frac{1 - \mathbb{E}[Y(1)]}{1 - \mathbb{E}[Y(0)]},$$
$$\tau^{\text{OR}} := \frac{\mathbb{P}[Y(1) = 1]}{\mathbb{P}[Y(1) = 0]} \left(\frac{\mathbb{P}[Y(0) = 1]}{\mathbb{P}[Y(0) = 0]}\right)^{-1}$$

- A null effect now corresponds to a Risk Ratio of 1
- Survival Ratio (SR) corresponds to the RR with swapped labels Y
- \bullet RR is not symmetric to the choice of outcome 0 and 1 –e.g. counting the living or the dead while Odds Ratio (OR) is

Different treatment measures give different impressions

RCT from Cook and Sackett (1995)

- Y = 1 stroke in 5 years and Y = 0 no stroke
- X=1 low baseline risk $\mathbb{P}\left[Y(0)=1\mid X=0\right]\geq \mathbb{P}\left[Y(0)=1\mid X=1\right]$

	$ au_{RD}$	$ au_{RR}$	$ au_{SR}$	$ au_{NNT}$	$ au_{OR}$
All (P _R)	-0.0452	0.6	1.05	22	0.57
X = 1	-0.006	0.6	1.01	167	0.6
X = 0	-0.08	0.6	1.1	13	0.545

- RD: treatment reduces by 0.045 the probability to suffer from a stroke
- ullet RR: the treated has 0.6 imes the risk of having a stroke comp. with the control
- SR: there is an increased chance of not having a stroke when treated compared to the control by a factor 1.05.
- NNT: one has to treat 22 people to prevent one additional stroke
- \bullet OR \approx RR in a stratum where prevalence of the outcome is low
- RD is heterogeneous with X while RR is homogeneous with X
- ⇒ Heterogeneity's property defined w.r.t. (i) covariates & (ii) a measure

The age-old question of how to report effects



"We wish to decide whether we shall count the failures or the successes and whether we shall make relative or absolute comparisons"

- Mindel C. Sheps, New England Journal of Medicine, in 1958

The choice of the measure is still actively discussed

e.g. Spiegelman and VanderWeele, 2017; Baker and Jackson, 2018; Feng et al., 2019; Doi et al., 2022; Xiao et al., 2021, 2022; Huitfeldt et al., 2021; Lapointe-Shaw et al., 2022; Liu et al., 2022 ...

- CONSORT guidelines recommend to report all of them

A desirable property collapsibility

i.e. population's effect is equal to a weighted sum of local effects



Simpson paradox:

(a) Overall population, $\tau_{\rm OR} \approx 0.26$

	Y=0	Y=1
A=1	1005	95
A=0	1074	26

(b) $au_{
m OR|F=1} pprox 0.167$ and $au_{
m OR|F=0} pprox 0.166$

F= 1	Y=0	Y=1
A=1	40	60
A=0	80	20

F=0	Y=0	Y=1
A=1	965	35
A=0	994	6

Marginal effect larger than subgroups' effects Unfortunately, not all measures are collapsible

Writting marginal effect as a weighed sum of conditional effects

Direct collapsibility

$$\mathbb{E}\left[\tau(X)\right] = \tau$$

⇒ RD directly collapsible:

$$\tau_{\scriptscriptstyle \rm R}^{\scriptscriptstyle \rm RD} = {\color{red} p_{\scriptscriptstyle \rm R}(X=1)} \times \tau_{\scriptscriptstyle \rm R}^{\scriptscriptstyle \rm RD}(X=1) + {\color{red} p_{\scriptscriptstyle \rm R}(X=0)} \times \tau_{\scriptscriptstyle \rm R}^{\scriptscriptstyle \rm RD}(X=0)$$

Ex: $0.0452 = -0.47 \times 0.006 - 0.53 \times 0.08$

Weights are equal to the population's proportions

Useful for generalization! (replacing p_R by p_T)

Writting marginal effect as a weighed sum of conditional effects

Direct collapsibility

$$\mathbb{E}\left[\tau(X)\right] = \tau$$

 \Rightarrow RD directly collapsible:

$$au_{ extsf{R}}^{ extsf{RD}} = extsf{p}_{ extsf{R}}(X=1) imes au_{ extsf{R}}^{ extsf{RD}}(X=1) + extsf{p}_{ extsf{R}}(X=0) imes au_{ extsf{R}}^{ extsf{RD}}(X=0)$$

Ex: $0.0452 = -0.47 \times 0.006 - 0.53 \times 0.08$

Weights are equal to the population's proportions Useful for generalization! (replacing p_R by p_T)

Collapsibility (require knowing Y(0))

$$\mathbb{E}\left[w(X, P(X, Y(0))) \tau(X)\right] = \tau$$

⇒ RR collapsible:

$$\mathbb{E}\left[\tau_{RR}(X)\frac{\mathbb{E}\left[Y(0)\mid X\right]}{\mathbb{E}\left[Y(0)\right]}\right] = \tau_{RR}$$

Measures' properties

Direct collapsibility

$$\mathbb{E}\left[\tau(X)\right]=\tau$$

Collapsibility (require knowing Y(0))

$$\mathbb{E}\left[w(X, P(X, Y(0))) \tau(X)\right] = \tau \quad \text{with } w \ge 0, \ \mathbb{E}\left[w(X, P(X, Y(0)))\right] = 1$$

Logic respecting

$$au \in \left[\min_{x}(au(x)), \max_{x}(au(x))\right].$$

Measure	Collapsible	Logic-respecting
Risk Difference (RD)	Yes	Yes
Number Neeeded to Treat (NNT)	No	Yes
Risk Ratio (RR)	Yes	Yes
Survival Ratio (SR)	Yes	Yes
Odds Ratio (OR)	No	No

Non parametric generative models

Continuous outcome

Assuming that
$$\mathbb{E}\big[|Y(1)|\, \big|X\big] < \infty$$
 and $\mathbb{E}\big[|Y(0)|\, \big|X\big] < \infty$

$$Y(0) = f(0,X) + \varepsilon_0$$
, with $f(0,X) = \mathbb{E}[Y(0) \mid X]$
 $Y(1) = f(1,X) + \varepsilon_1$, with $f(1,X) = \mathbb{E}[Y(1) \mid X]$

$$r(1) = r(1,\lambda) + \varepsilon_1, \text{ with } r(1,\lambda) = \mathbb{E}[r(1)|\lambda]$$

$$Y(w) = \underbrace{f(0,X)}_{:=b(X)} + w \underbrace{(f(1,X) - f(0,X))}_{:=m(X)} + \underbrace{w\varepsilon_1 + (1-w)\varepsilon_0}_{:=\varepsilon_w}.$$

Additive decomposition

$$Y(w) = b(X) + w m(X) + \varepsilon_w,$$

Baseline
$$b(X) := \mathbb{E}[Y(0) \mid X]$$

Modification
$$m(X) := \mathbb{E}[Y(1) - Y(0) \mid X]$$

$$\mathbb{E}\left[\varepsilon_{w}\mid X\right]=0$$

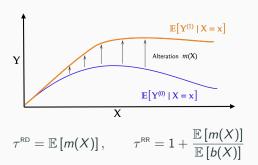
Spirit of Robinson's decomposition (1998), also in Nie et al. (2020)

Nonparametric generative model, continuous outcome

$$Y(w) = b(X) + w m(X) + \varepsilon_w,$$

Baseline $b(X) := \mathbb{E}[Y(0) \mid X]$

Modification $m(X) := \mathbb{E}[Y(1) - Y(0) \mid X]$



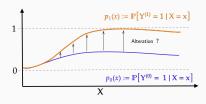
- With RR, baseline b(X) is entangled with effect m(X)
- RD independent of baseline

Ex. of the Russian Roulette: Harmful "homogeneous" treatment effect

Nonparametric generative model, binary outcome

$$\mathbb{P}[Y(w) = 1 \mid X = x] = b(x) + w \underbrace{(1 - b(x))}_{\text{Entanglement}} \frac{1}{6}.$$

Baseline
$$b(X) := \mathbb{P}[Y(0) = 1 \mid X = x]$$

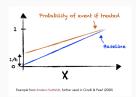


Ex. of the Russian Roulette: Harmful "homogeneous" treatment effect

Nonparametric generative model, binary outcome

$$\mathbb{P}\left[Y(w) = 1 \mid X = x\right] = b(x) + w \underbrace{\left(1 - b\left(x\right)\right)}_{\text{Entanglement}} \frac{1}{6}.$$

Baseline
$$b(X) := \mathbb{P}[Y(0) = 1 \mid X = x]$$



$$\tau^{\text{\tiny RD}} = \frac{1}{6} \left(1 - \mathbb{E} \left[b(x) \right] \right), \quad \lim_{\mathbb{E} \left[b(x) \right] \to 1} \tau_{\text{\tiny RD}} = 0. \quad \tau_{\text{\tiny SR}} = 1 - \frac{1}{6}$$

- With RD, baseline b(X) is entangled with effect 1/6
- SR independent of baseline

Heterogeneous treatment effect

Nonparametric generative model, binary outcome

$$m_g(x) := \mathbb{P}[Y(1) = 0 \mid Y(0) = 1, X = x], \quad m_b(x) := \mathbb{P}[Y(1) = 1 \mid Y(0) = 0, X = x]$$

$$\mathbb{P}[Y(w) = 1 \mid X = x] = b(x) + w (1 - b(x)) m_b(x) - w b(x) m_g(x),$$

A beneficial effect $(m_b(x) = 0)$ is visible on a high baseline $(b(x) \approx 1)$.

A deleterious effect $(m_g(x)=0)$ is visible on low baseline $(1-b(x)\approx 1)$

Heterogeneous treatment effect

Nonparametric generative model, binary outcome

$$m_g(x) := \mathbb{P}[Y(1) = 0 \mid Y(0) = 1, X = x], \quad m_b(x) := \mathbb{P}[Y(1) = 1 \mid Y(0) = 0, X = x]$$

$$\mathbb{P}\left[Y(w)=1\mid X=x\right]=b(x)\underbrace{+w\left(1-b\left(x\right)\right)m_{b}\left(x\right)}_{\nearrow}\underbrace{-w\,b\left(x\right)m_{g}\left(x\right)}_{\nearrow},$$

A beneficial effect $(m_b(x) = 0)$ is visible on a high baseline $(b(x) \approx 1)$.

A deleterious effect $(m_{\rm g}(x)=0)$ is visible on low baseline (1-b(x)pprox 1)

Disantanglement of treatment effect and baseline

Assuming that the treatment is beneficial (i.e. $\forall x, m_b(x) = 0$),

$$\tau_{RR}(x) = 1 - m_g(x).$$

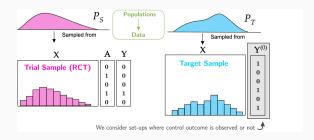
Computing group effects on X affecting b(.): constant effect

Harmful effect: conditional SR (like the Russian Roulette)

Beneficial effect: conditional RR

Back to generalizability

Aim: transport trial finding's to a target population, using the trial data and a sample of the target population

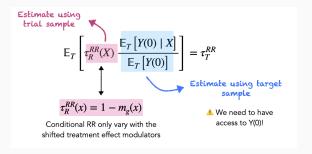


Back to generalizability

Generalizing	Conditional Outcome	Local effects
Assumption	$\mathbb{E}_{\mathbb{R}}[Y(w) \mid X] = \mathbb{E}_{\mathbb{T}}[Y(w) \mid X = x]$	$ au_R(X) = au_{T}(X)$
Unformal	All shifted prognostic covariates	All shifted effect modifiers
Identification	$\mathbb{E}_{T}\left[Y(w)\right] = \mathbb{E}_{T}\left[\mathbb{E}_{R}\left[Y(w) \mid X\right]\right]$	$\mathbb{E}_{\mathbb{R}}\left[\frac{p_{T}(X)}{p_{R}(X)}g_{T}(Y(0),X)\tau_{R}(X)\right]$

- Depending on the assumptions, either conditional outcome or local treatment effect can be generalised
- ullet Generalizing local effects only for collapsible measure, information on $Y^{(0)}$ with weights required

Generalize local effect: Y binary and a beneficial effect



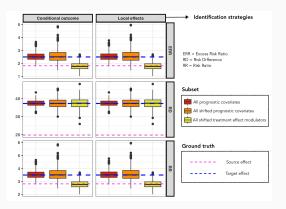
Different covariates sets are required to retrieve the target population effect depending on

- (i) the causal measure
- (ii) the nature of the outcome (continuous, binary),
- (iii) the method to generalize (conditional outcome or local effect).

Simulations: continuous outcome

$$Y = b(X_1, X_2, X_3, X_4, X_5, X_6) + W m(X_1, X_2, X_5) + \varepsilon.$$

 $X_1, \ldots X_4$ shifted. Linear setting

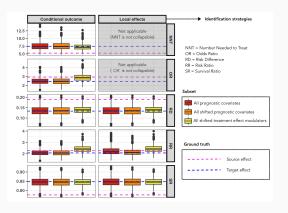


- RD can be generalized with only X_1, X_2
- Generalizing conditional outcome & local effects similar
- Adding prognostic variables improve precisions

Simulations: binary outcome

$$\mathbb{P}[Y(w) = 1 \mid X = x] = b(X_1, X_2, X_3) + a(1 - b(X_1, X_2, X_3)) \ m_b(X_2, X_3),$$

 $X_1 = \text{lifestyle}, \ X_2 = \text{stress}, \ X_3 = \text{gender}; \ \text{gender not shifted}$



• SR can generalize with Stress only when generalizing local effects

Conclusion: many medical and statistical challenges

Some measures are easier to generalize (i.e. needs less covariates to adjust on): less sensitive to a population's shift

Summary

- Collapsibility is an important property
- ▶ Generalize by conditional outcomes requires shifted prognostic var.
- ▶ Generalize by local effects requires shifted treatment effect modifiers
- ▷ Continuous outcome: RD depends on the modification
- ▷ Binary outcome: Conditional SR for harmful, RR for beneficial effect

²¹https://cran.r-project.org/web/views/CausalInference.html

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Summary

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- ▶ Generalize by conditional outcomes requires shifted prognostic var.
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On going

- ▷ Estimation strategies to estimate/generalize the RR
- ▷ Insights from/Impact for Meta Analyses
- Usefulness of the model both beneficial and harmful...
- ▶ Methods used are methods implemented. Taskview on causal inference ²¹
- > Federated causal inference

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- Usefulness of the model both beneficial and harmful...
- ▶ Methods used are methods implemented. Taskview on causal inference ²¹
- > Federated causal inference
 - > Logistic (non parametric) regression model Constant conditional OR

²¹https://cran.r-project.org/web/views/CausalInference.html

Collaborators





Bénédicte Colnet (DSS), Erwan Scornet (LPSM),



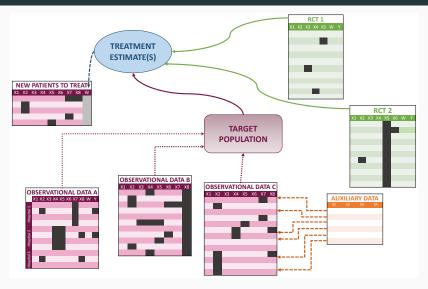




Imke Mayer (Owkin), Tobias Gauss (Traumabase), G. Varoquaux (Inria)

- Risk ratio, odds ratio, risk difference... Which causal measure is easier to generalize? Submitted
- Mayer, et al. Doubly robust treatment effect estimation with missing attributes. Annals of Applied Statistics, 14(3), 2020.
- Colnet, Mayer et al. Causal inference methods for combining randomized trials and observational studies: a review. *Statitsical Science*. 2023.
- Mayer, Josse. Generalizing treatment effects with incomplete covariates: identifying assumptions and multiple imputation algorithms. Biometrical Journal. 2022.
- Colnet et al. Causal effect on a target population: a sensitivity analysis to handle missing covariates. Journal of Causal Inference. 2022.
- Colnet et al. Reweighting the RCT for generalization: finite sample analysis and variable selection. *Submitted*. 2022.

Real-world data can help strengthen clinical evidence



black correspond to sporadically & systematic missing covariates

$$P(Y^{(0)} = 1) X = x = m(x) + \frac{1 - m(x)}{6}$$

$$P(Y^{(0)} = 1) X = x = m(x) + \frac{1 - m(x)}{6} + \frac{5}{6} m(x)$$

$$P(Y^{(0)} = 1) X = x = m(x) + \frac{1 - m(x)}{6} + \frac{5}{6} m(x)$$

$$= m(x) + \frac{1 - m(x)}{6}$$

Resources

<u>R-miss-tastic</u> https://rmisstastic.netlify.com/R-miss-tastic

J., I. Mayer, N. Tierney & N. Vialaneix

Project funded by the R consortium (Infrastructure Steering Committee) 22

Aim: a reference platform on the theme of missing data management

- ▷ list existing packages
- ▷ available literature
- > analysis workflows on data
- \Rightarrow Federate the community
- \Rightarrow Contribute!

²²https://www.r-consortium.org/projects/call-for-proposals

Appendix

Semi-synthetic simulation

- ▷ All the results are illustrated on semi-synthetic simulations;
- ▷ Build from two large clinical data bases, reflecting a real-world situation
 - \diamond CRASH3 \sim 9 000 individuals.
 - ♦ Traumabase ~ 30 000 individuals.
- ▶ The outcome is the only synthetic part,

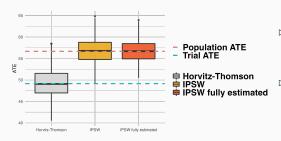
$$Y := f(GCS, Gender) + A \tau(TTT, Blood Pressure) + \epsilon_{TTT},$$

More precisely,

$$Y = 10 - { t Glasgow} + ({ t if Girl:} - 5 { t else:} 0) \ + A \left(15(6 - { t TTT}) + 3*({ t Systolic.blood.pressure} - 1)^2 \right) + arepsilon_{ t TTT},$$

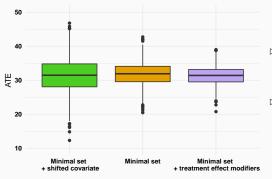
where $\varepsilon_{\rm TTT}$ is a random Gaussian noise with a standard deviation depending on the value of the covariate TTT.

Results from the semi-synthetic simulation



- \triangleright Reduced variance for IPSW fully estimated (π is also estimated).
- ➤ The re-weighted trial has not necessarily a larger variance.

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Papier:

Using Big Data to Emulate a Target Trial When a Randomized Trial Is Not Available

Miguel A. Hernán* and James M. Robins *

Correspondence to Dr. Miguel A. Hernán, Department of Epidemiology, 677 Huntington Avenue, Boston, MA 02115 (e-mail: miguel_hernan@post.harvard.edu). Initially submitted December 9, 2014; accepted for publication September 8, 2015.

Protocol component	CRASH2	CRASH3	Traumabase
Eligibility criteria	Adult trauma patients with, or at risk of, significant bleeding who were within 8 hours of injury.	Adults with TBI who were within 3 h of injury, had a Glasgow Coma Scale (GCS) score of 12 or lower or any intracranial bleeding on CT scan, and no major extracranial bleeding	Patients of age ≥ 16 years. Until 2019, the treatment TXA was only recommended for patients with hemorrhage (external or internal) and not specifically for TBI (which is not necessarily caused by intracranial hemorrhage)
Treatment strategies	Tranexamic acid (loading dose 1 g over 10 minutes then infusion of 1 g over 8 hours)	TXA (loading dose 1 g over 10 min then infusion of 1 g over 8 h) in less than 3 hours after injury	TXA is given before/at entry in the hospital in case of extracranial bleeding, time and dose are not registered
Assignment procedures	Participants are randomized (1:1)	Participants are randomized (1:1)	Observational study
Follow-up period	28 days after accident	28 days after accident	28 days after accident
Outcome	Death in hospital within 4 weeks of injury	Head injury-related death in hospital within 28 days of injury	Head injury-related death within 28 days of injury
Causal contrasts of interest	Intention-to-treat effect	Intention-to-treat effect	Intention-to-treat effect

TBI is a complex condition caused by severe trauma (e.g. in an accident) and it can be assumed that the severity of TBI is determined/fixed early after the accident i.e., the severity of TBI is "hard coded" once TBI occurs and the following treatments won't affect the severity, they only affect the chances of survival or other functional outcomes conditionally on the TBI severity.

In theory TBI severity could be assessed quickly after its occurrence (at the scene of the accident). But it is assessed and treated only at the hospital, once the patient is stable enough (i.e. without ongoing strong hemorrhage) to be placed in a CT scan.

One of the limit when combining data or comparing evidence is that TBI is not assessed in the same way in the different data set

Causal inference²³

Causal inference (simplest) question



Assume a policy/intervention/treatment W causes an outcome Y Aim: estimate the effect as acurately as possible (bias & variance)

²³Taskview to organize all packages on causal inference.

Causal inference²³

Potential Outcome framework (Neyman, 1923; Rubin, 1974)

treatment

$$ho$$
 n iid sample $(\underbrace{X_i}_{\text{covariates}}, \underbrace{W_i}_{\text{vi}}, \underbrace{Y_i(1), Y_i(0)}_{\text{potential outcomes}}) \in \mathbb{R}^d \times \{0, 1\} \times \mathbb{R} \times \mathbb{R}$

ightharpoonup Individual **causal effect** of the binary treatment: $\Delta_i = Y_i(1) - Y_i(0)$

Problem: Δ_i never observed (only observe one outcome/indiv)

Covariates		Treatment	Outcome(s)		
X_1	X_2	X_3	W	Y(0)	Y(1)
1.1	20	F	1	?	200
-6	45	F	0	10	?
0	15	М	1	?	150
-2	52	М	0	100	?

²³Taskview to organize all packages on causal inference.

Causal inference²³

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$$ho$$
 n iid sample $(\underbrace{X_i}_{\text{covariates}}, \underbrace{W_i}_{\text{potential outcomes}}, \underbrace{Y_i(1), Y_i(0)}_{\text{potential outcomes}}) \in \mathbb{R}^d \times \{0, 1\} \times \mathbb{R} \times \mathbb{R}$

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1.1	20	F	1	?	200
-6	45	F	0	10	?
0	15	М	1	?	150
-2	52	M	0	100	?

Average Treatment Effect (ATE): $\tau = \mathbb{E}[\Delta_i] = \mathbb{E}[Y_i(1) - Y_i(0)]$

The ATE is the difference of the average outcome had everyone gotten treated and the average outcome had nobody gotten treatment

²³Taskview to organize all packages on causal inference.

Randomized Controlled Trial

Identifiability assumptions

$$P(i) = W_i Y_i(1) + (1 - W_i) Y_i(0) \qquad \text{(consistency)}$$

$$P(i) = W_i \perp \{Y_i(0), Y_i(1), X_i\} \qquad \text{(random treatment assignment)}$$

Flip a coin to assign the treatment

We can check that
$$\tau = \mathbb{E}[\Delta_i] = \mathbb{E}[Y_i(1)] - \mathbb{E}[Y_i(0)]$$

$$= \mathbb{E}[Y_i(1)|W_i = 1] - \mathbb{E}[Y_i(0)|W_i = 0]$$

$$= \mathbb{E}[Y_i|W_i = 1] - \mathbb{E}[Y_i|W_i = 0]$$

 \Rightarrow Although Δ_i never observe, τ is identifiable and can be estimated

Difference-in-means estimator

$$\hat{\tau}_{DM} = \frac{1}{n_1} \sum_{W_i=1} Y_i - \frac{1}{n_0} \sum_{W_i=0} Y_i$$
, where $n_w = \sum_{i=1}^n \mathbb{1}_{W_i=w}$

$$\hat{\tau}_{DM}$$
 unbiased and \sqrt{n} -consistent $\sqrt{n}(\hat{\tau}_{DM} - \tau) \xrightarrow[n \to \infty]{d} \mathcal{N}(0, V_{DM})$

Randomized Controlled Trial

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Covariates		Treatment	Outcome(s)		
X_1	X_2	X_3	W	Y(0)	Y(1)
1.1	20	F	1	?	200
-6	45	F	0	10	?
0	15	M	1	?	150
-2	52	M	0	100	?

$$\hat{\tau}_{DM} = \frac{1}{n_1} \sum_{W_i=1} Y_i - \frac{1}{n_2} \sum_{W_i=0} Y_i;$$
 ATE = mean(red)-mean(blue)

Randomized Controlled Trial (RCT)

- same covariate distributions of treated and control groups
 - \Rightarrow High **internal** validity

Randomized Controlled Trial (RCT)

- - ⇒ High internal validity
- expensive, long, ethical limitations
- small sample size: restrictive inclusion criteria
 - \Rightarrow No personalized medicine
- trial sample different from the population eligible for treatment
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Observational data

- ▷ low cost
- ▷ large amounts of data (registries, biobanks, EHR, claims)
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- representative of the target populations
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Randomized Controlled Trial (RCT)

- pold standard (allocation ⊕)
- Same covariate distributions of
 - treated and control groups

 ⇒ High internal validity
- ▷ expensive, long, ethical limitations
- - ⇒ No personalized medicine
- trial sample different from the population eligible for treatment
 - ⇒ Low external validity

Observational data

- ▷ "big data": low quality
- ▷ lack of a controlled design opens the door to confounding bias
 - ⇒ Low **internal** validity
- ▷ low cost
- ▷ large amounts of data (registries, biobanks, EHR, claims)
 - \Rightarrow patient's heterogeneity
- representative of the target populations
 - ⇒ High external validity