# STATISTICS & ML WITH R

# Mapping causal & predictive approaches

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#### **MODULE 4 – LECTURE OUTLINE**

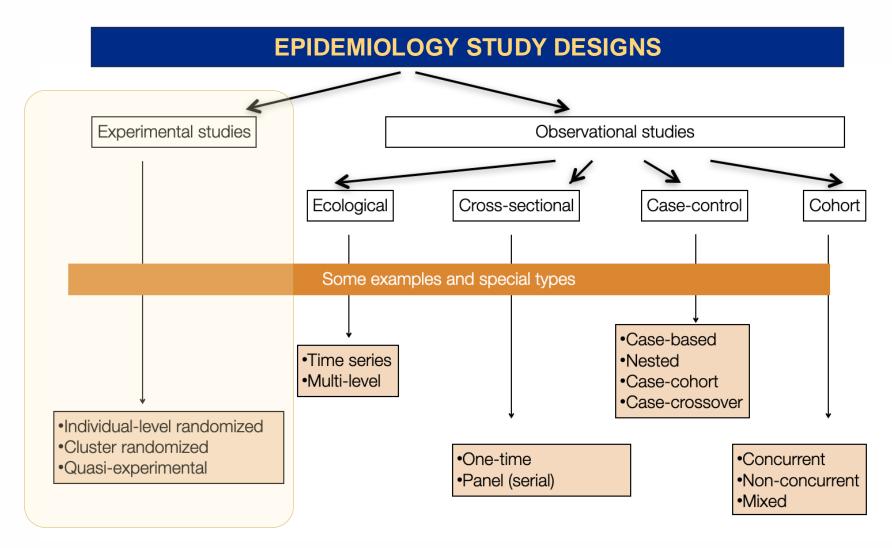
#### Mapping causal approaches

- Recall the essential features of experimental study designs
  - Learning the vocabulary of causal analysis
- Get a visual intuition of causal pathways, including challenging elements:
  - Collider variables
  - Confounder variables
  - Mediator variables
- Discuss the correct causal model to capture the association among exposure, outcome and other covariates, (including challenging ones)
- Define causal outcomes and choosing the appropriate "estimands":
  - ATE, ATT, or ATU?
- Devise statistical methods to estimate ATE, ATT, and ATU based on research question and (sub)population of interest

#### From *observational* to *experimental* studies

- "OBSERVATIONAL STUDIES" on variables of interest and their relationships have no controlled assignment of the treatment
  - We may find CORRELATION / ASSOCIATION, but it DOES NOT IMPLY CAUSATION!
     Why?
  - ... hidden variables may affect the relationship between the explanatory variable and the response variable
  - ...but often used (implicitly or not) to estimate causal effect of an exposure!
- "EXPERIMENTAL STUDIES" seek to uncover CAUSATION, so they are designed to provoke a response
  - Researchers assign the treatment to an experimental unit (or subject) and observing its effect
  - These studies use some ad hoc design principles and controlled independent variables

### Experimental and non-experimental study designs...



Source: https://bookdown.org/jbrophy115/bookdown-clinepi/design.html

## Different goals of statistical modeling (part 1/2)

- **1.** ASSOCIATION/CORRELATION → observational studies
  - aimed at summarizing or representing the data structure, without an underlying causal theory
  - may help form hypotheses for explanatory and predictive modeling
- 2. CAUSAL EXPLANATION → experimental studies
  - aimed at **testing "explanatory connection"** between <u>treatment and</u> outcome variables
  - prevalent in "causal theory-heavy" fields (economics, psychology, environmental science, etc.)
- Note:
  - ✓ The same modeling approach (e.g., fitting a regression model) can be used for different goals
  - ✓ While they shouldn't be confused, explanatory power and predictive accuracy are complementary goals: e.g., in bioinformatics (which has little theory and abundance of data), predictive models are pivotal in generating avenues for causal theory.
- **3. EMPIRICAL PREDICTION** → algorithmic machine learning and datamining modeling

# A framework for CAUSAL ANALYSIS

Key terminology

### The conceptual framework for causal analysis (1/3)

#### Fundamental vocabulary:

- Intervention decisions and actions that change the behaviors or situation of people/firms/other subjects (drug, vaccine, program participation)
  - TREATMENT = commonly used in experimental studies when researchers directly "assigns" the causal variable
  - EXPOSURE = commonly used observational studies when participants "naturally" experience the the **causal variable**
- Subjects = those that may be affected (at least in principle), in fact are
  - TREATED subjects
  - UNTREATED subjects
- Outcome = variable(s) that may be affected by the intervention
  - can be caused by exposure either directly or through an intermediate process
- Causation = causal processes that lead to the development of outcomes

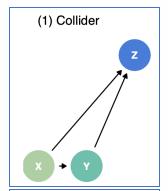
## The conceptual framework for causal analysis (2/3)

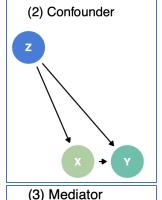
- - **Bias** = systematic error that can occur <u>at different stages</u> of the study: *data collection, analysis* or *interpretation* of the causal relationship exposure-outcome.
    - **Selection bias** = both the exposure and the outcome affect whether an individual is included in the sampled population
      - Sampling bias = some members of the intended population are less likely to be included than others
      - Attrition bias = participants who drop out of a study systematically differ from those who remain
      - Non-response bias = participants who refuse to participate in the study systematically differ from those who take part
    - Recall bias = a systematic difference in the ability of participant groups to accurately recall information
    - Information bias = there is misclassification or inaccurate measurement (e.g., patients underreporting smoking habits)
    - •

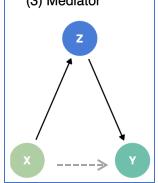
Check out this cool list of all types of bias: <a href="https://quantifyinghealth.com/list-of-biases/">https://quantifyinghealth.com/list-of-biases/</a>

#### The conceptual framework for causal analysis (3/3)

- Fundamental vocabulary ("tricky ones" ( ):
  - Collider = variable that is influenced by treatment and outcome (like a "common effect")
    - **EXAMPLE**: sleepiness (Z), with shift work (X) and apnea (Y)
    - Conditioning on or controlling for a collider in the causal model can create a distortion ("collider bias")
  - Confounder = variable that affect both treatment and outcome ("apparent" cause), but it is not in the causal pathway
    - **EXAMPLE**: smoking (Z), with exercise (X) and lung cancer (Y)
    - Most confounder variables involve some kind of selection (e.g., self-selection) that can be addressed stratifying subjects by it
  - Mediator = is a variable that is in the causal pathway and "explains" why treatment affects outcome (like a "mechanisms")
    - **EXAMPLE**: immune function (Z), with exercise (X) and lung cancer (Y)
    - Conditioning on or controlling for a mediator can be done to assess what part of the effect they play







#### **Estimands, Estimators, Estimates**

- The **estimand** is the **target outcome** of interest about the causal effect of a treatment in a population
  - EXAMPLEs: ATE (Average Treatment Effect), ATT (Average Treatment Effect on the Treated), or ATU (Average Treatment Effect on the Untreated)
- The **estimator** is the **statistical method** ("recipe") by which we approximate this *estimand* using data
  - EXAMPLEs: difference-in-means for ATE in a randomized controlled trial (RCT), or propensity scores matching (PSM) for ATT within observational data.
- The estimate is the numerical value we get when we plug our data into the estimator
  - **EXAMPLE**: we calculate an **ATE = 3.5 units** (e.g., a treatment improves test scores by 3.5 points on average in the entire population)

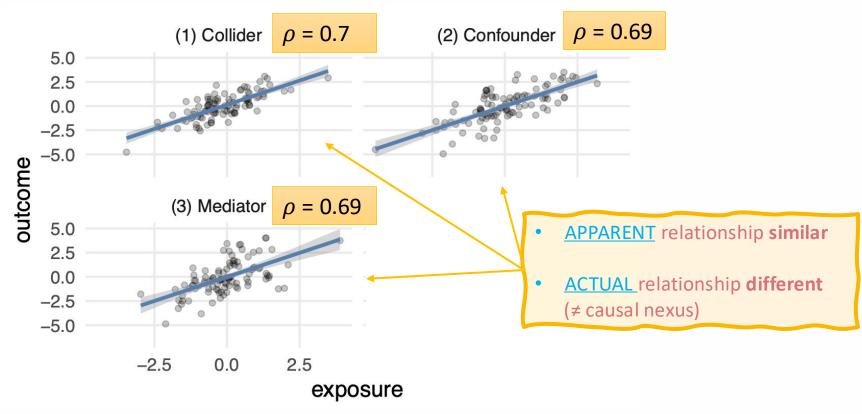
## Visualizing causal maps

Using "DAGs" in guiding statistical modeling

### Typical challenges in estimating causal effects: a.k.a.

### "correlation does not imply causation"

- Consider 3 distinct datasets: while their statistical summaries and visualizations are very similar, the true causal nexus differs!
- **Deciding the** correct model requires knowledge of the data-generating mechanism (i.e. the random assignment to exposure/not exposure in experiments)



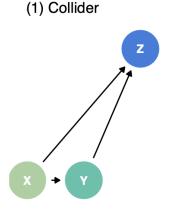
Source: Barrett, M., McGowan, L. D., & Gerke, T. (2024). Causal Inference in R. Retrieved from https://www.r-causal.org/

## Typical challenges in estimating causal effects: visual intuition

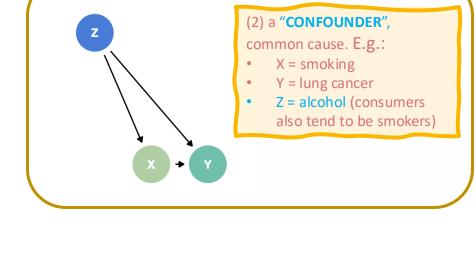
- Directed Acyclic Graphs (DAGs) can offer visual intuition of the causal nexus at play in the 3 datasets. Failure to adjust models to these situation leads to BIAS
  - X is some exposure of interest, Y an outcome, and Z a known, measured factor

(1) a "COLLIDER", common effect (that invertedly connects). E.g.:X = sodium intake

- Y = systolic blood pressure
- Z = urinary protein excretion



(3) Mediator

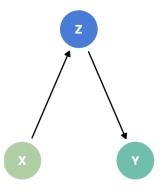


19

(2) Confounder

(3) a "MEDIATOR" is caused by X and then it causes Y. E.g.:

- X = screen time
- Y = obesity
- Z = physical exercise



Source: Barrett, M., McGowan, L. D., & Gerke, T. (2024). Causal Inference in R. Retrieved from https://www.r-causal.org/

#### How to deal with collider (common effect) when modeling?

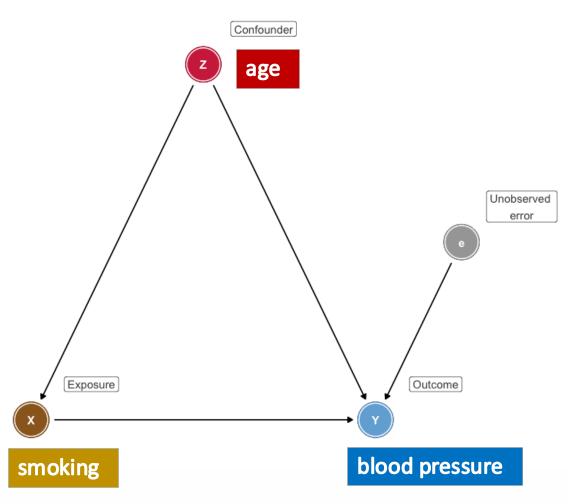
Causal map with COLLIDER (Z) Collider hospitalization Unobserved error Exposure Outcome smoking cancer

- We must NOT control for collider.
- Colliders CAN HIDE REAL CAUSE EFFECTS
  - i.e., it would distort the true relationship between the exposure and the outcome

(more in Lab. 4)

#### How to deal with confounder (common cause) when modeling?

Causal map with CONFOUNDER (Z)



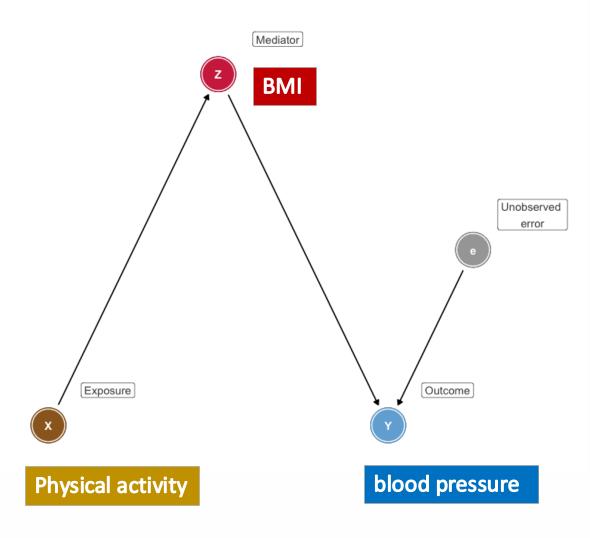
- We must control for a <u>confounder</u>, so we reduce bias. HOW?
- 1) At **design** stage:
  - Random assignment
  - Restriction (only participants of a certain confounder category)
  - Matching observations (confounder distributed evenly by exposure)
- 2) In **analysis** stage:
  - Stratifying sample in subgroups (by confounding)
  - Including term in regression
  - Inverse probability weighting (equalizing frequency of counfounder by exposure)
  - Instrumental variable estimation

Source: Excellent reading https://quantifyinghealth.com/control-confounding/

(more in Lab. 4)

#### How to deal with mediator (mechanism) when modeling?

Causal map with MEDIATOR (Z)



- We could control for the mediator, depending on which effect we focus on:
  - with UNADJUSTED MODEL we get only the <u>total</u> effect (direct + indirect) of X on Y
  - with ADJUSTED MODEL we separate the <u>direct</u> effect of X on Y (not mediated), and the <u>indirect</u> effect of M on Y (mediated)
- Normally both models are shown
- The Adjusted model enables to see the PROPORTION of the MEDIATOR mechanism in the causal path

(more in Lab. 4)

# Measuring causal outcomes of interest

Commonly used "estimands" (ATE, ATT, ATU) and how to select and interpret them correctly for making valid inferences

The explanation and examples below follow very closely these two sources:

- **1.** Noah Greifer, Elizabeth A. Stuart, <u>Choosing the Causal Estimand for Propensity Score Analysis of Observational Studies</u>
- 2. Andrew Heiss, Demystifying causal inference estimands: ATE, ATT, and ATU

### Defining estimands at the subject level

#### NOTATION:

- $Y^0$  and  $Y^1$  are the potential outcomes in the absence and presence of treatment for patient i in a study on a new drug on blood pressure,
  - $Y_i^0$  = patient's blood pressure with receives standard of care
  - $Y_i^1$  = patient's blood pressure with takes new drug
- ITE = Individual Treatment Effect (\*) = difference, for subject i, between potential outcome y if treated and if untreated

$$te_i = \delta_i = y_i^1 - y_i^0$$
 where treatment is  $T_i = \{0,1\}$ 

- (\*) ITE is *never* observable!!
- Hence, we will look at averages...
- ATE = Average Treatment Effect = average of ITE differences across subjects

$$E[te_i] = E[Y_i^1 - Y_i^0] = E[Y_i^1 | T_i = 1] - E[Y_i^0 | T_i = 0]$$

- (\*) The Avg of the differences = the difference of Averages!
- ATE can hide different distributions of ITEs (e.g., positives and negatives that cancel each outer out)
- Important to have a well-defined group or population

### Defining estimands at the subject level

 ATT (or ATET) = Average Treatment effect on the Treated = average treatment effect across all subjects that end up TREATED

$$E[\delta_i \mid T_i = 1] = E[Y_i^1 - Y_i^0 \mid T_i = 1] = E[Y_i^1 \mid T_i = 1] - E[Y_i^0 \mid T_i = 1]$$

- This refers to the avg of the differences conditionally on the fact that both groups "received" the treatment (" $|T_i=1"$ )
- $[Y_i^0 | T_i = 1]$  is essentially the counterfactual for  $Y_i^1$  in a 'parallel universe' where **exactly** the same people who were treated in this universe would not get the treatment
- ATU = Average Treatment effect on the Untreated = average treatment effect across all subjects who were NOT TREATED

$$E[\delta_i \mid T_i = 0] = E[Y_i^1 - Y_i^0 \mid T_i = 0] = E[Y_i^1 \mid T_i = 0] - E[Y_i^0 \mid T_i = 0]$$

- This time we seek the Avg of the differences (" $|T_i=1 \>$ ") conditionally on the fact that both groups were "assigned" to the treatment
- $[Y_i^1 \mid T_i = 0]$  is essentially the counterfactual for  $Y_i^0$  in a 'parallel universe' where **exactly the same people** who were NOT treated in this universe would get the treatment

### By the way!

- Treatment is a binary random variable  $T_i = \{0,1\}$
- Outcome of interest is  $Y_i = \begin{cases} Y_i^0 & (if \ T_i = 0) \\ Y_i^1 & (if \ T_i = 1) \end{cases}$
- ATE = Average Treatment Effect = average of ITE differences across subjects
- ATT/ATET = Average Treatment effect on the Treated = average treatment effect across all subjects that end up TREATED

	EXAMPLE: Does hospitalization (T) increase health (Y)?		
ATE =	$E[Y_i^1   T_i = 1] - E[Y_i^0   T_i = 0]$	Avg health of hospitalized group – avg health of <b>NOT</b> hospitalized group	
ATT +	$E[Y_i^1 \mid T_i = 1] - E[Y_i^0 \mid T_i = 1]$	Avg health of treated group – [counterfactual] avg health $E[Y_i^0]$ of treated group IF NOT hospitalized	
+ Selection bias	$\begin{aligned} \mathbf{E}[\mathbf{Y_i^0} \mid \mathbf{T_i} = 1] &- \mathbf{E}[\mathbf{Y_i^0} \mid \mathbf{T_i} = 0] \\ \text{(hospitalized have worse } \mathbf{Y_i^0} \text{ than } \\ \text{non hospitalized)} \end{aligned}$	Difference in [counterfactual] avg health $E[Y_i^0]$ of treated group IF NOT hospitalized - those who were NOT hospitalized	

## EXE. potential causal outcomes, ITE ( $\delta_i$ ), depends on patients' characteristics)

	Confounder	Treatment		Unobservab	le	Realized	
	Age			outcomes /	ITE	Outcome	
ID	$Z_i$	$X_i$	$Y_i^1$	$Y_i^0$	$Y_i^1-Y_i^0$	$Y_i$	20 + 5 + 5 + 5
1	Old	1	80	60	20	80	$ATT = \frac{20 + 3 + 3 + 3}{4}$
2	Old	1	75	70	5	75	7
3	Old	1	85	80	5	85	
4	Old	0	70	60	10	60	
5	Young	1	75	70	5	75	10 + 0 - 10 +
6	Young	0	80	80	0	80	<b>ATU</b> = 4
7	Young	0	90	100	-10	100	
8	Young	0	85	80	5	,′ 80	
					1		

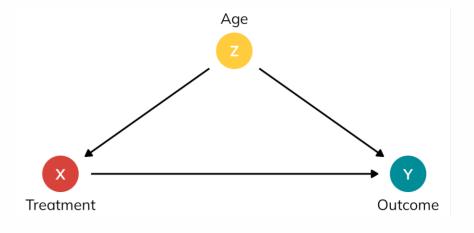
ATE = 
$$\frac{20 + 5 + 5 + 5 + 10 + 0 + -10 + 5}{8} = 5$$
  
ATE =  $\left(\frac{4}{8} \times 8.75\right) + \left(\frac{4}{8} \times 1.25\right) = 4.375 + 0.625 = 5$  (ATE decomposition)

# Revisiting the confounder seen in DAG visualization

Accounting for age for an accurate estimate of ATE

#### Acknowledging a confounder variable

			<i>_</i> -	
	1	Confounder	Treatment	Realized
	\ -1 -1	Age	Treated	Outcome
ID	1	$Z_i$	$X_i$	$Y_i$
1	\   	Old	1	80
2	١	Old	1	75
3	 	Old	1	85
4	1	Old	0	60
5	1	Young	1	75
6	 	Young	0	80
7	(   	Young	0	100
8	1	Young	,′ 0	80



- So far, we ignored it, but:
- AGE seems to behave as a confounder:
  - → it is highly correlated with treatment status
  - → it also affects the ultimate value of the outcome
- Hence, we need to account for it statistically

#### How to deal with confounder variable?

- Recall that we listed different ways to control for a confounder (to reduce bias in estimands)
- Here, we illustrate 2 of them (feasible at analysis stage):
- 1) At **design** stage:
  - Random assignment
  - Restriction (only participants of a certain confounder category)
  - Matching observations (confounder distributed evenly by exposure)
- 2) In **analysis** stage:
  - ✓ Stratifying sample in subgroups (by confounding)
  - ✓ Including confounder variable as term in regression
  - Inverse probability weighting (equalizing frequency of counfounder by exposure)
  - Instrumental variable estimation

## 1) Stratification to deal with confounder (i.e. combining the weighted averages for old and young people)

	Confounder	Treatment	Realized
,	Age	Treated	Outcome
ID \	$Z_i$	$X_i$	$Y_i$
1	Old	1	80
2	Old	1 1 1 1	75
3	Old	1 1	85
4	Old	0	60
5	Young	1	75
6	Young	0	80
7	Young	0	100
8	Young	0	80

Effect<sub>old</sub> = 
$$\overline{Y}_{treated} - \overline{Y}_{untreated} =$$

$$= \frac{80 + 75 + 85}{3} - \frac{60}{1} = 20$$

Effect<sub>young</sub> = 
$$\overline{Y}_{treated} - \overline{Y}_{untreated} =$$

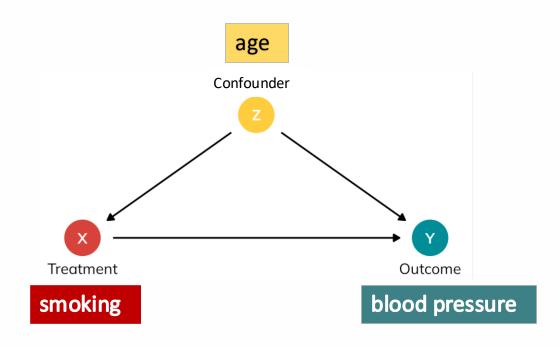
$$= \frac{75}{1} - \frac{80 + 100 + 80}{3} = -11.667$$

$$\mathbf{ATE}_{stratified} = \pi_{old} \mathbf{Effect}_{old} + \pi_{young} \mathbf{Effect}_{young} = \left(\frac{4}{8} \times 20\right) + \left(\frac{4}{8} \times -11.667\right) = 4.1667$$

After stratification based on the confounder we get a very close *approximation of the ATE* (=5)

#### 2) Introducing the confounder as term in the regression model

- Let's consider an example (to be discussed in the Lab) based on the NHANES dataset,
- where:
  - Z = Age = confounder
  - X = SmokeNow
  - Y = BPSysAve (blood pressure)
- In the Lab, we will fit a regression model for the outcome and compare the results WITH and WITHOUT the confounder in the model



```
# Unadjusted linear model
model_unadjusted <- lm(BPSysAve ~ SmokeNow, data = nhanes_conf)

# Adjusted model
model_adjusted <- lm(BPSysAve ~ SmokeNow + Age, data = nhanes_conf)</pre>
```

## 2) Introducing the confounder as term in the regression model reduces bias and isolates the true effect of the X on Y

- In the Adjusted Model (with Age), the estimate of the causal effect of smoking (SmokeNowYes) on blood pressure (BPSysAve) is more accurate
  - the regression "adjusts" for the influence of Z in the causal path X → Y
  - this prevents falsely attributing to Smoking (X) an effect that might actually result from X

NO Confounder	Confounder
124.384**	100.997**
[123.537, 125.231]	[98.863, 103.131]
s.e. = 0.432	s.e. = 1.089
-4.264**	0.684
[-5.524, -3.004]	[-0.554, 1.922]
s.e. = 0.643	s.e. = 0.631
	0.432**
	[0.395, 0.468]
	s.e. = 0.019
3108	3108
0.014	0.158
0.01	
	124.384**  [123.537, 125.231]  s.e. = 0.432  -4.264**  [-5.524, -3.004]  s.e. = 0.643  3108  0.014

#### Ways to deal with confounders must be carefully pondered

All these solutions have pros and cons that must be considered...

#### WHAT CAN BE DONE AT DATA COLLECTION?

- RANDOM ASSIGNMENT to treatment exposure
- **RESTRICTION** of participants of a certain confounder category
- MATCHING by distributing the confounder evenly (between exposed and unexposed)

#### WHAT CAN BE DONE DURING ANALYSIS?

- **STRATIFICATION** (as we saw by Age) by estimating the relationship outcome within different subsets of the confounder
- REGRESSION (as we saw with Age) can scale to many confounders
- INVERSE PROBABILITY WEIGHTING to balance confounder-weighted distributions and achieve comparability between treated and untreated groups

# What about the other estimands? (ATT, ATU)

## Choosing the estimands and the proper statistical method to estimate the effect

- In a randomized trial, the treated and untreated groups will, on average, have the same distributions of patient characteristics, so the ATT, ATU, and ATE will be the same
- Without randomization, however, the treatment groups can have quite different distributions of characteristics, ATT, ATU, and ATE will differ when these characteristics also relate to the treatment effect
  - So, when using observational data: for whom should the treatment effect be estimated?

#### Choosing the estimands based on the research question

<u>BEFORE</u> analyzing a dataset, let's consider which question we are asking, and about which target population group,

<u>THEN</u> choose a statistical method that corresponds to the chosen estimand.

Estimands	Target Population	Example research question and research/policy addressed
ATT	Treated patients	Examining an intervention that would only reach those currently receiving it: - e.g. decision to replace / withhold a treatment for currently treated patients
ATU	Untreated patients (control)	How would untreated patients respond to a new potential treatment/exposure? - e.g. decision to extend a medical practice (drug prescription/vaccine) to a group that would not otherwise receive it
ATE	Full sample / population	Should a specific policy be applied to all eligible patients? How would the outcome be on average? - e.g. regulating a system-wide policy for a previously unregulated practice - useful when treatment decisions are not well informed (ATE does not depend on current treatment assignment) - NOT OK when patients' benefit depend on clinical judgment

#### Recapping key points of the lecture

- Capturing the causal nexus between a treatment and an outcome may be tricky due to:
  - constraints in study design, sampling, or data collection process
  - repeated measures over time
  - effects of confounder, collider or mediator/mechanism variables
  - etc.
- Visual causal maps may help guiding the analysis, by summarizing how variables affect each other
  - e.g. DAGs (Directed Acyclic Graphs)
- Another key step is deciding which estimand(s) we are seeking with reference to the specific target population:
  - ATE if a treatment targeted to the general population (e.g. a universal policy)
  - ATT if a treatment targeted subjects already exposed (e.g. ≠ drug for treated patients)
  - ATU if a new treatment apply to currently untreated patients (e.g. new drug)
- Each estimand implies its own assumptions, interpretation, and statistical methods
- coming up next...

# Shifting emphasis on empirical outcome prediction

Introduction to Machine Learning (ML) models

# A conceptual framework to understand different types of statistical modeling (part 2/2)

- **1.** association/correlation → observational studies
- **2.** causal explanation  $\rightarrow$  experimental studies
- **3. empirical prediction** → algorithmic machine learning and data-mining modeling
  - aimed at predicting new or future observations (without necessarily explaining how)
  - relies on big data
  - prevalent in fields like natural language processing, bioinformatics, etc.. In epidemiology, there is more of a mix <u>causal explanation & empirical</u> prediction

#### NOTES:

✓ "Prediction" does not necessarily refer to future events, but rather
to future datasets that were previously unseen to the algorithm

## ...stay tuned for next chapter on ML

