



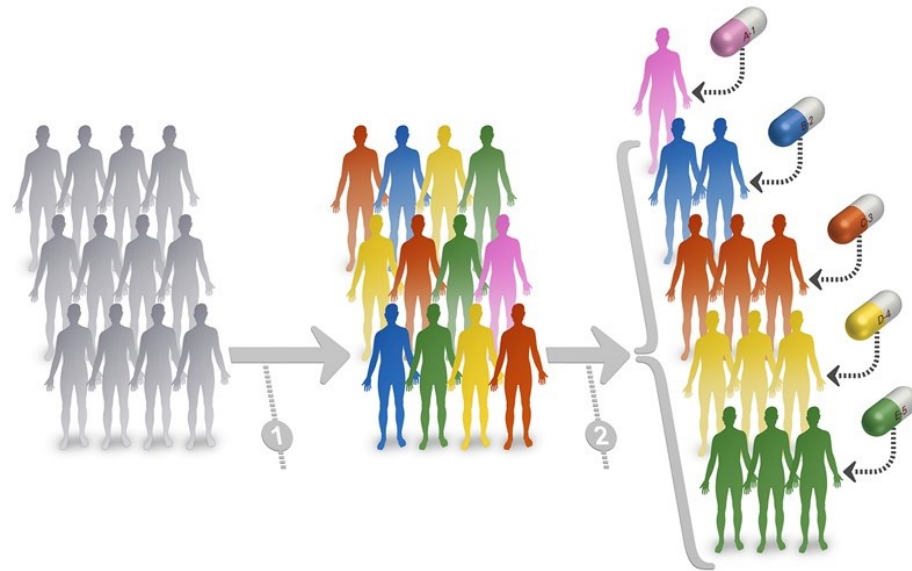
# When and How Can Machine Learning be Used for Treatment Recommendations in a Clinical Setting?

CHAIR-SU Workshop: The Learning Hospital  
March 2023

*Uri Shalit, Technion*

# Using patient data to personalize treatment

- One of the ultimate promises of big data in healthcare
- Especially important when there are no clear clinical guidelines



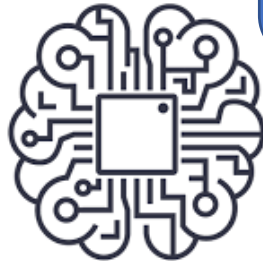


You have condition A. Treatment options are either  $T=0$  or  $T=1$




I'm not so sure what's best...

Recommend  $T=0$



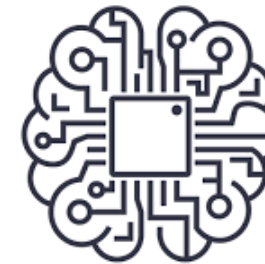
# Using patient data to personalize treatment: causal inference

- Decision making  requires **causal modeling**:  
Taking actions in the world
- Especially if model uses observational data
  - E.g. data collected from hospitals, clinics, and by patients themselves
  - Such data generally suffers from **confounding**
- No way to know if we are correct before deploying the system!
- How do we build confidence and avoid harm?



I'm not so sure what's best...

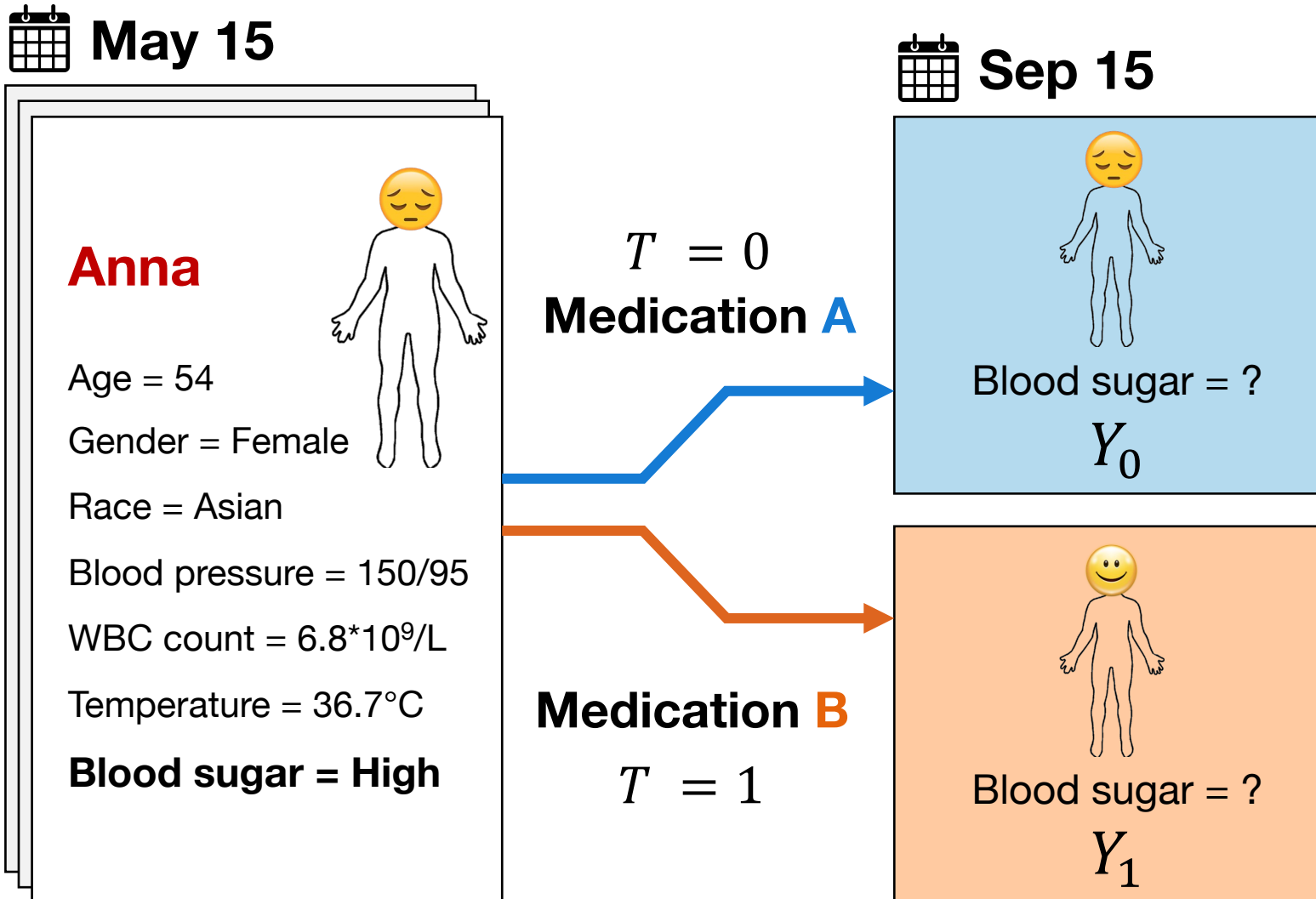
Recommend  
T=0



# This talk

- We propose a best-practices framework for using patient clinical data to build a treatment recommendation model
  - Responsibly
  - Not focused on a specific algorithm
- Three phases:
  - 1. Identification: can the data even do what I want it to do for me?**
  - 2. Estimation: what does the data tell me to do?**
  - 3. Validation: how much should I believe the model I just estimated?**

# $Y_0, Y_1$ : potential outcomes (Rubin, Neyman)

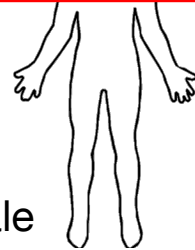


$Y_0, Y_1$ : potential outcomes  
(Rubin, Neyman)

$$CATE(x) \equiv \mathbb{E}[Y_1 - Y_0 | x]$$

Conditional Average Treatment Effect

**Anna**



Age = 54

Gender = Female

Race = Asian

Blood pressure = 150/95

WBC count =  $6.8 \cdot 10^9/L$

Temperature =  $36.7^\circ C$

**Blood sugar = High**

$x$

**Medication A**



Blood sugar = ?

$Y_0$

**Medication B**

$T = 1$

Blood sugar = ?

$Y_1$

$Y_0, Y_1$ : potential outcomes  
(Rubin, Neyman)

$$CATE(x) \equiv \mathbb{E}[Y_1 - Y_0 | x]$$

- We never directly observe CATE
- We only see either  $Y_1$  or  $Y_0$
- The choice is *not random*

Blood pressure = 150/95

WBC count =  $6.8 \cdot 10^9/L$

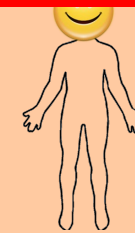
Temperature =  $36.7^\circ C$

**Blood sugar = High**

$x$

**Medication B**

$T = 1$



Blood sugar = ?

$Y_1$



# Individual-level treatment effects: CATE

- We wish to estimate the individual-level treatment effect, formally denoted **Conditional Average Treatment Effect (CATE)**

- In Rubin-Neyman potential outcome notation:

$$CATE(x) \equiv \mathbb{E}[Y_1 - Y_0 | x] = \mathbb{E}[Y_1 | x] - \mathbb{E}[Y_0 | x]$$

“**what if** we forced the patients with features  $x$  to receive treatment  $T = 1$ , vs. forced them to receive treatment  $T = 0$ ”

- We never directly observe  $CATE(x)$
- We can't provably know “**what if**”

# From CATE to recommendation

- $CATE(x) \equiv \mathbb{E}[Y_1 - Y_0 | x]$
- General idea:  
Estimate  $\widehat{CATE}(x)$  for incoming patient with features  $x$
- Present recommendation to doctor:
- $\widehat{CATE}(x) < 0 \rightarrow \text{recommend } T = 1$   
 $\widehat{CATE}(x) > 0 \rightarrow \text{recommend } T = 0$



# From CATE to recommendation

- $CATE(x) \equiv \mathbb{E}[Y_1 - Y_0 | x]$
- General idea:  
Estimate  $\widehat{CATE}(x)$  for incoming patient with features  $x$
- Present recommendation to doctor:
- $\widehat{CATE}(x) < 0 \rightarrow$  recommend  $T = 1$   
 $\widehat{CATE}(x) > 0 \rightarrow$  recommend  $T = 0$
- If uncertainty about  $\widehat{CATE}(x)$  is high  $\rightarrow$  **defer** recommendation



# Individual-level treatment effects: CATE

- $CATE(x) \equiv \mathbb{E}[Y_1 - Y_0|x]$
- $x$  is high-dimensional and practically unique to each unit
- Can (carefully) use machine learning based tools
  - Causal Forests (Wager & Athey 2015, 2018), Deep networks (Johansson, S, Sontag 2016, 2017, Parbhoo et al. 2018, Shi et al. 2019), Gaussian processes (Schulam & Saria 2018, Alaa & van der Schaar 2018), Meta-learning (Künzel et al. 2017, 2019, Nie & Wager 2017)
- **However:** These only work *under a strict set of **causal identification** conditions:*
  - no hidden confounders
  - common support between different treatments
  - no interference between units
- Most of these assumptions are not testable from data
- (Even supervised learning will not work unless the conditions hold)

# Major challenges

1

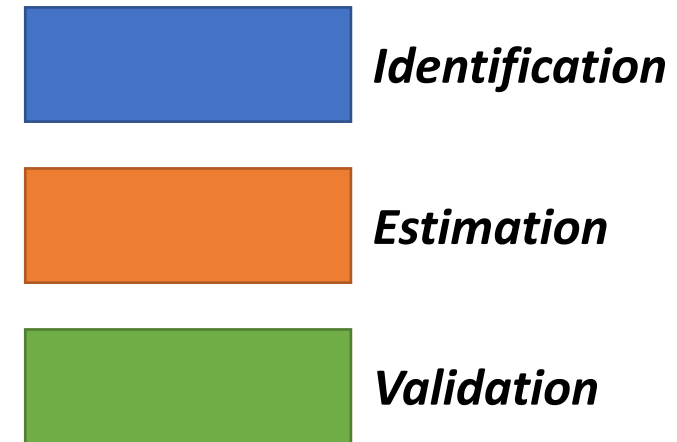
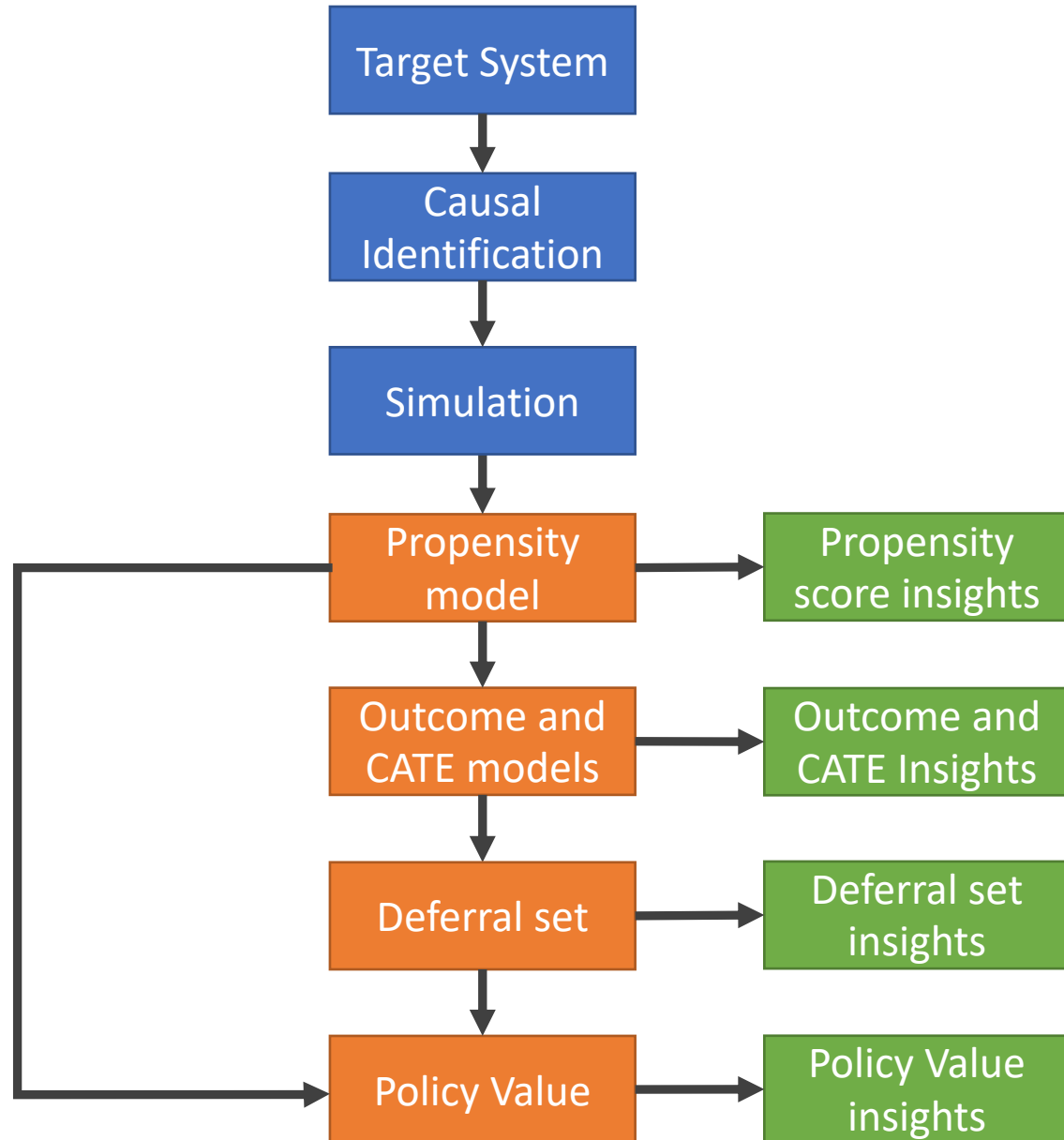
- All causal effect estimation methods rely on causal identification assumptions, some of which are hard to verify
- Prominently: “no unmeasured confounding”
- Some typical applications
  - Treatment assignment
  - Treatment dose
  - Important confounders: e.g. patient characteristics, physician model, e.g. imaging

How can we still build confidence and deploy a treatment recommendation system ?

- **There is no test set**
  - When our recommendation differs from what happened in practice → can't know for sure what would have happened had recommendation been used
- High stakes: even a pilot system might cause harm

2

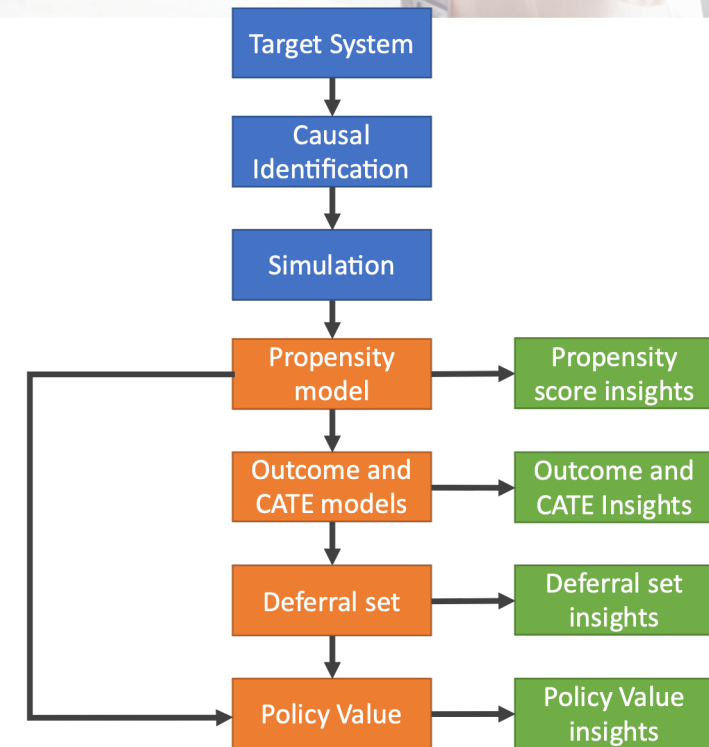
# Framework for robustly building causal decision support models



# Identification I: The Target System

(following Miguel Hernán's "Target Trial")

Define exactly the setting and context  
of the treatment recommendation system



# Identification I: The Target System

(following Miguel Hernán's "Target Trial")

Points for discussion with clinical partners:

1. Is treatment decision made by physicians at a **well-defined point in time**?
2. Is the set of possible actions small?
3. Are there clear clinical guidelines for decision?
4. Is there high variability in treatment decisions between physicians?
5. Are there well-defined and widely agreed upon outcomes?

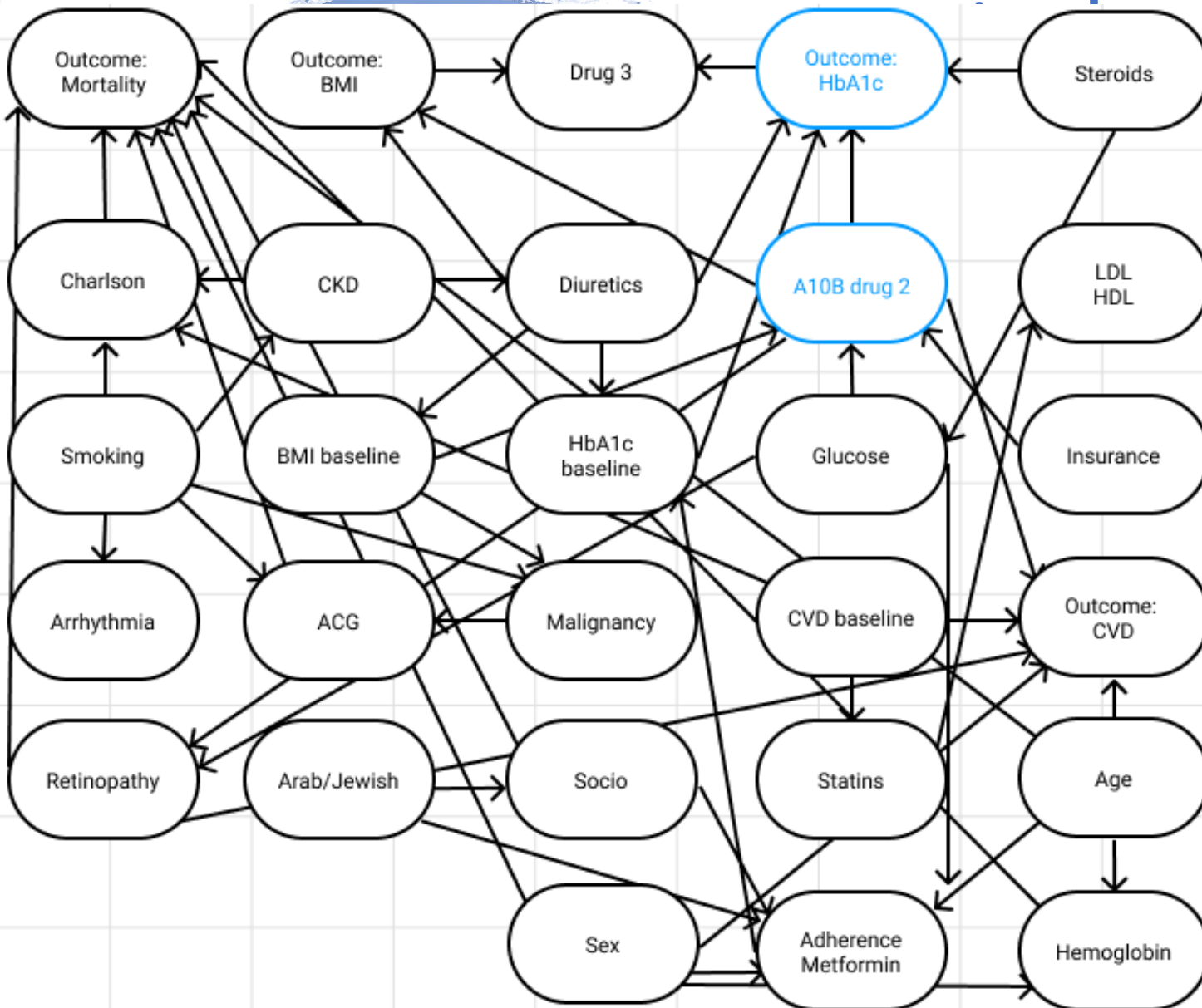
Help clarify discussion with clinicians about "AI assistants"





# Identification II:

data suitable for  
rig the target system?



- For observational data, have we measured all (most) known confounders?
- Do we have temporal separation of what data is recorded before/after the treatment assignment?
- Causal graphs built with domain experts can be useful here

First return point: no identification



First return point: no identification

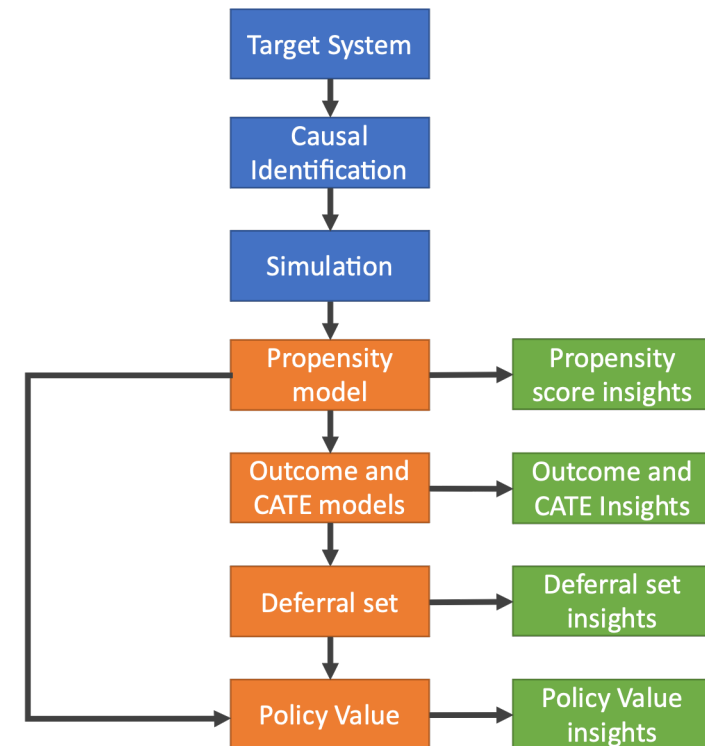


# Estimation I + II: CATE and propensity score

Many great methods for estimating:

- Propensity scores  $p(t = 1|x)$
- $CATE(x)$

Will not go into details here!



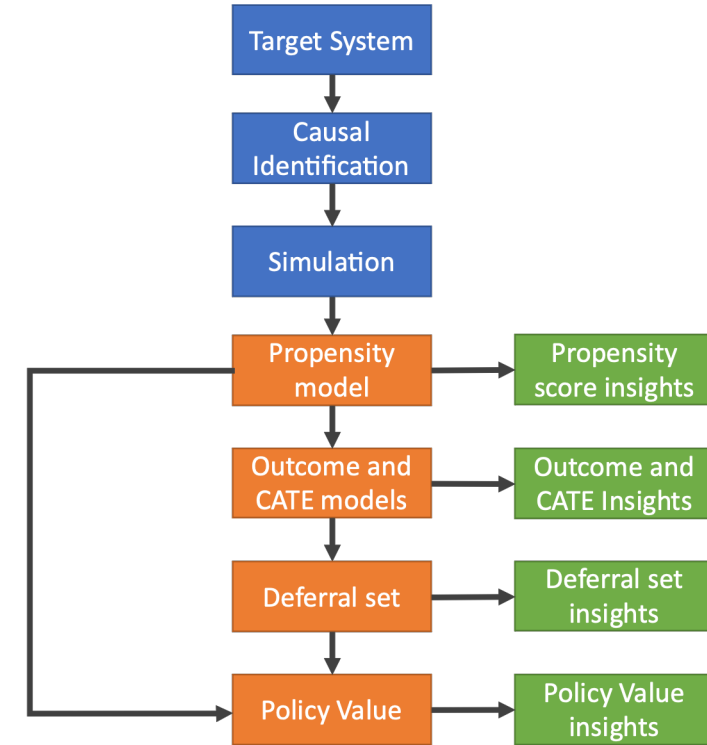
# Validation I + II: CATE and propensity score

- No traditional test set for CATE
- Still: should evaluate regression/classification models with (MSE, AUC, RMSE, etc.)

- Compare different models
- Apply interpretation
- Check with clinicians
- Characterize

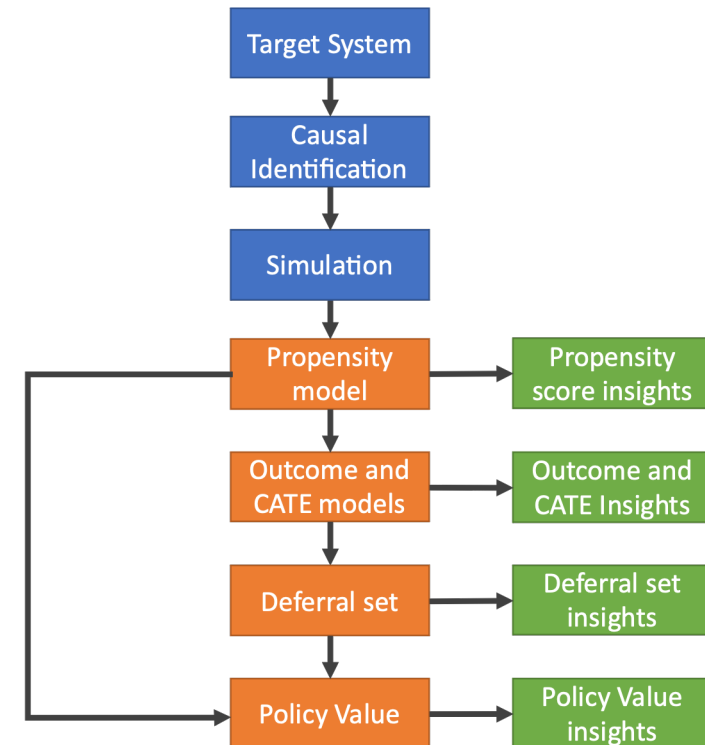


- Positive and negative sets:  
Who do we recommend receive each of the treatments?
- Clinician agreement and disagreement sets:  
Where do we think the clinicians were wrong?



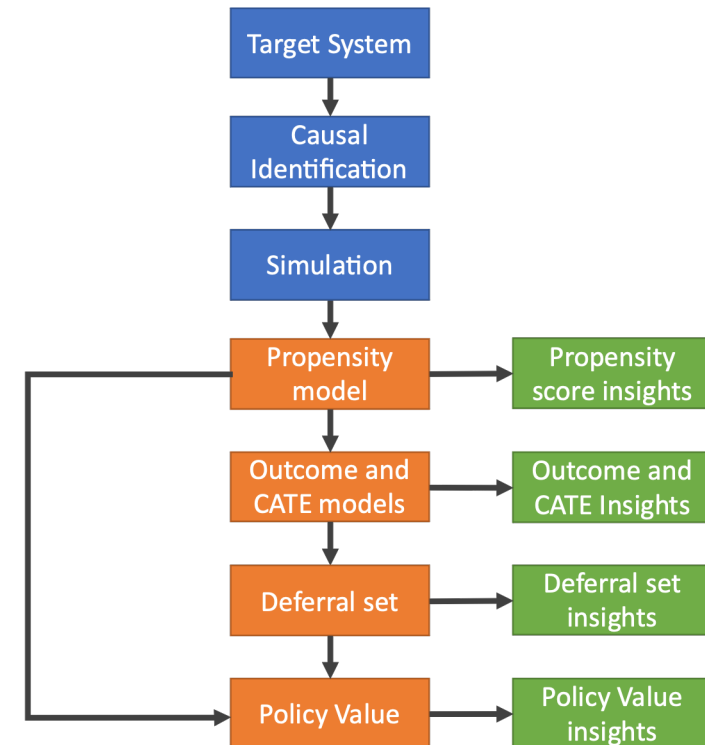
# Estimation III: Deferral Set

- Why we might doubt a given  $CATE(x)$  estimate:
  - Estimation error (finite samples, weak overlap, model misspecified, covariate shift)
  - Noise (measurement error, outcome stochasticity)
  - Causal error (**hidden confounding**)
- If in doubt, might wish to defer decision
- We work on modeling all sources of error



# Estimation IV: Policy Value

- Crucial metric for recommendations:  
“What would be the expected outcome if physicians treated as the model recommends”
- The **policy value** of the current “doctors’ policy” is simply the average outcome in the population
- A good recommendation would have a better policy value than the **doctors’ policy**
- Estimating policy values is itself a challenging causal problem
  - Don’t really know what would have happened in cases where our recommendations differ from the actual treatment in the data



# Preliminary results

- Analysis lead by graduate students Rom Gutman (Technion) and Shimon Sheiba (Technion, Cytoreason)





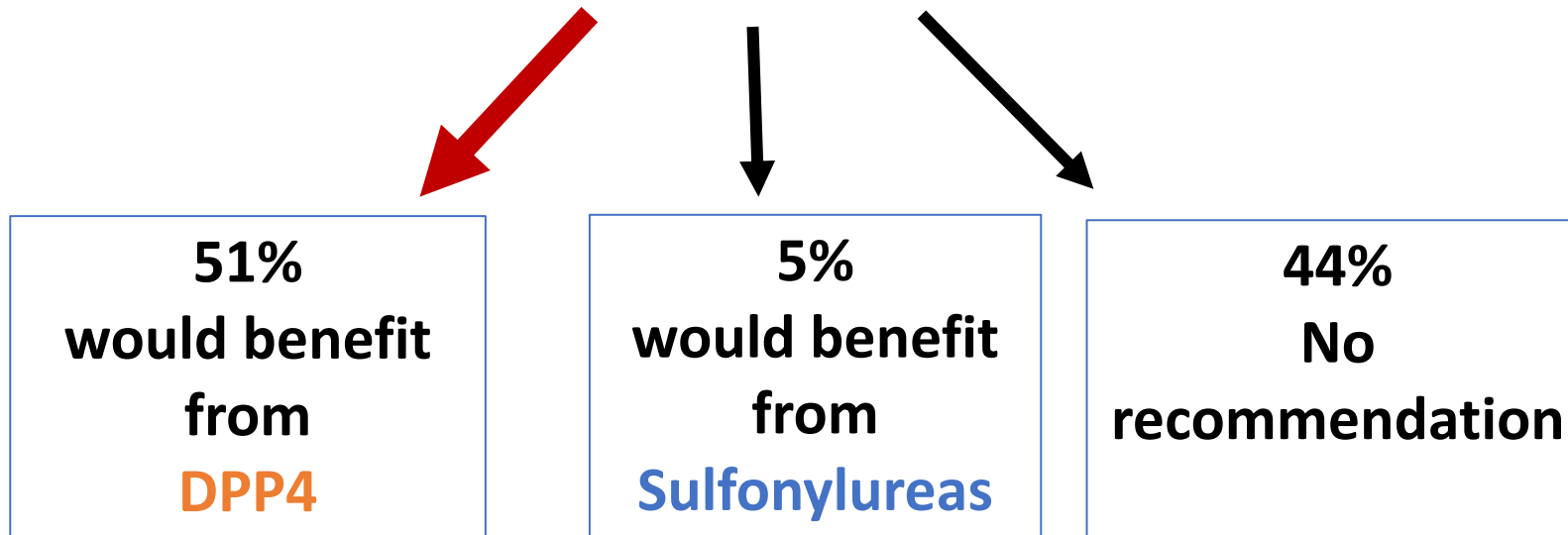
# Preliminary results 1: Chronic disease



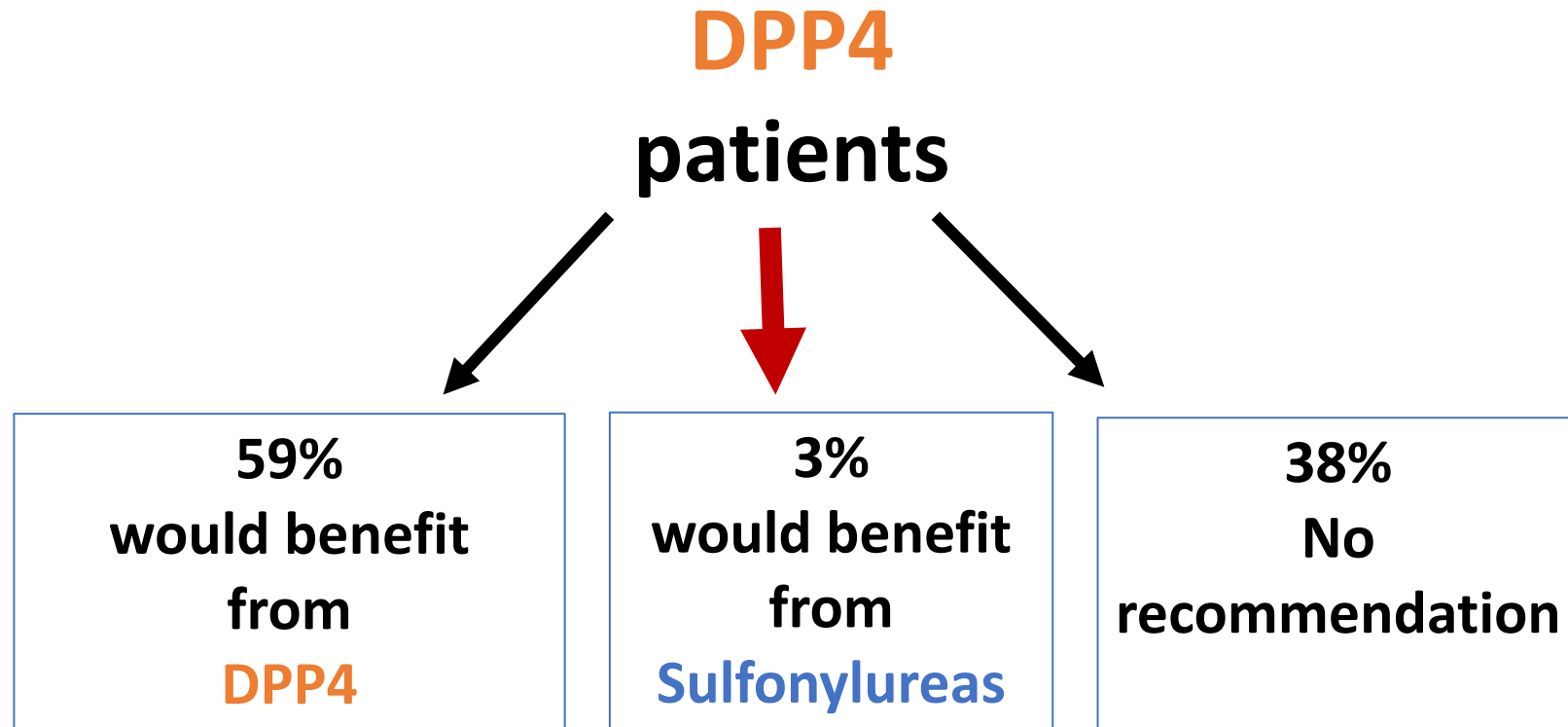
- Joint work with Clalit Research Institute (Prof. Ran Balicer)
- Investigating the effects of Sulfonylureas vs. DPP4 on type-II diabetes patients who have not responded to first-line therapy
- Goal: reduce blood sugar, measured in A1C
- More than 50,000 patients
- More than 200 covariates which are potential confounders: demographics, lab tests, diagnoses, medications, administrative and more

# Preliminary results – optimal care

## Sulfonylureas patients

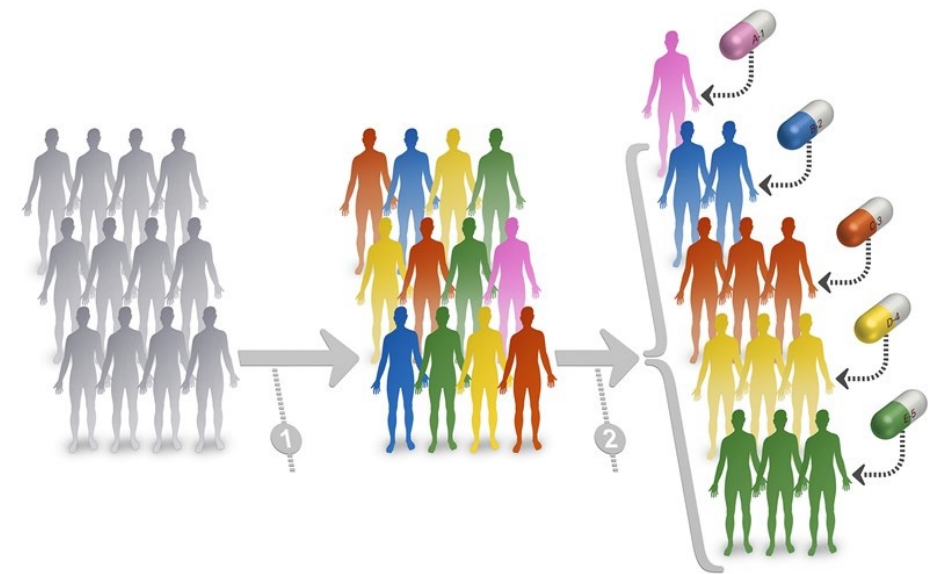


# Preliminary results – optimal care



# Is personalization worth it?

- Between Sulfonylureas and DPP4, the answer: no!
- We detect no significant difference between:
  - a) moving *everyone* to DPP4
  - b) personalized treatment
- Clinical trials later showed advantage of DPP4
- Newer medications are now in use



# Preliminary results: Acute disease

- The causal effects of diuretics on kidney function in hospitalized acute heart failure patients with kidney injury in Rambam Medical Center
- Clinical collaborators:  
Dr. Oren Caspi and Prof. Doron Aronson  
(Technion University & Rambam Health Care Campus)



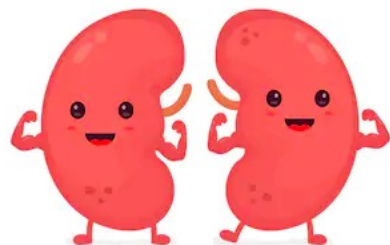
# Preliminary results: Heart failure with kidney injury

- Causal effects of *diuretics* on *kidney function* in hospitalized acute heart failure patients with kidney injury in Rambam Medical Center
- Physicians tell us:  
They have poor guidance how to prescribe diuretics and blood-pressure medications to these patients
- 2157 hospitalized heart failure patients with rise in serum creatinine, indicating kidney injury
- More than 200 covariates which are potential confounders:  
demographics, lab tests, diagnoses, medications, administrative and more
- Empirically: half of cohort had **increased diuretics or leave the same**, half had **decreased diuretics**

# Preliminary results

## Heart Failure with kidney injury

- T=1: “Decrease diuretics”
  - Often improves kidney function
  - Might hurt cardiac function
- Must balance multiple outcomes
- *“Should we increase, keep or decrease diuretics for this patient?”*



# RESULTS: Held-out cohort (n=530)

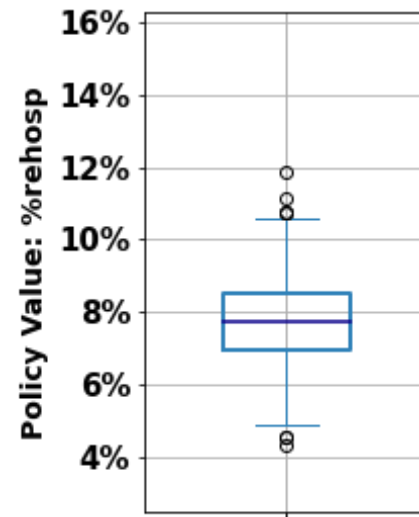
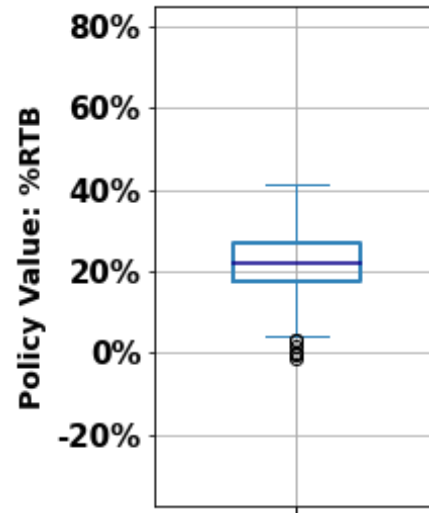
Compare outcomes under  
current practice vs. proposed  
Causal Machine Learning Model

Top: kidney function (higher=better)  
%RTB = %Return-to-baseline creatinine

Bottom: rehospitalization  
(indicator of cardiac function, lower=better)

- Our recommendations are better than current practice for %RTB ( $p=0.015$ , median 41% vs. 22%) and somewhat better for rehos. ( $p=0.048$ , median 6.5% vs. 7.7%)

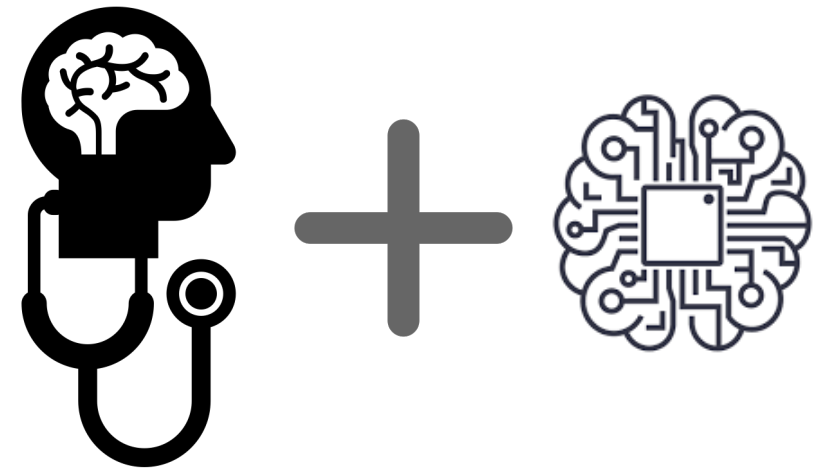
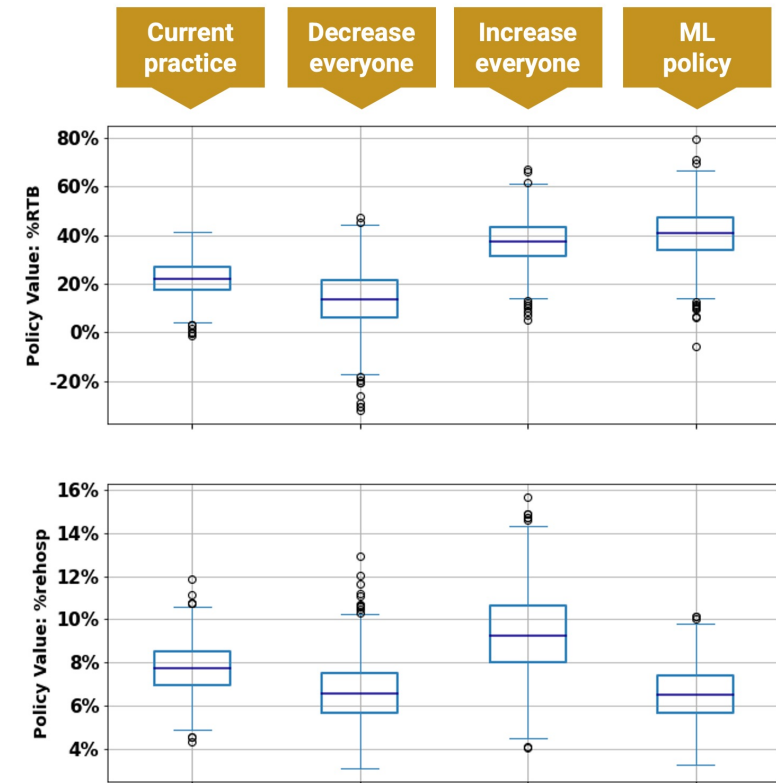
Current  
practice





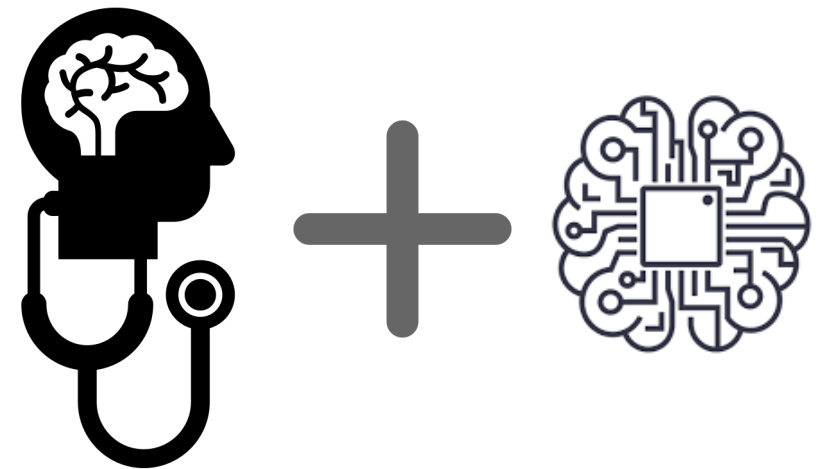
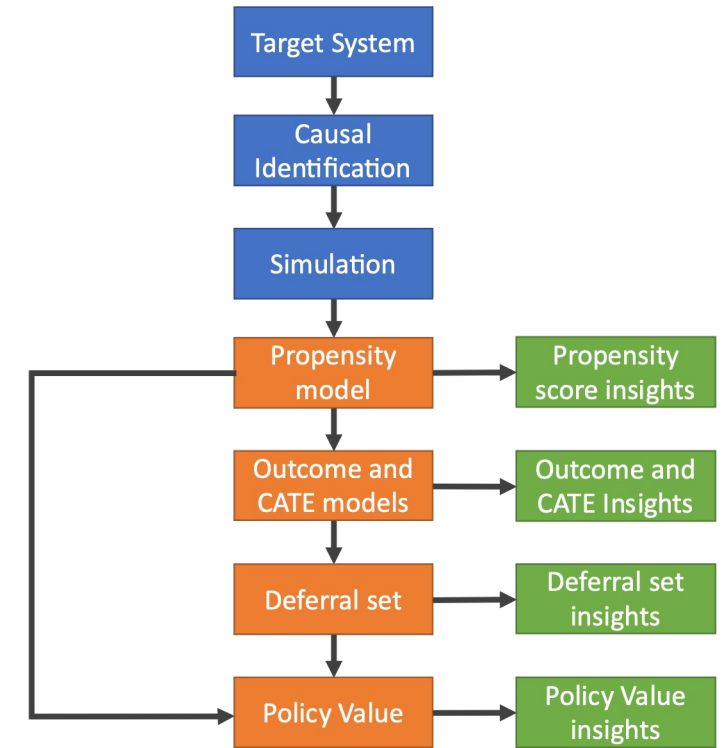
# Should we believe this?

- We don't **really** know what would have happened if our recommendations would have been followed
- The only way to truly know if our recommendations are useful: Run an experiment testing the system with real physicians and patients



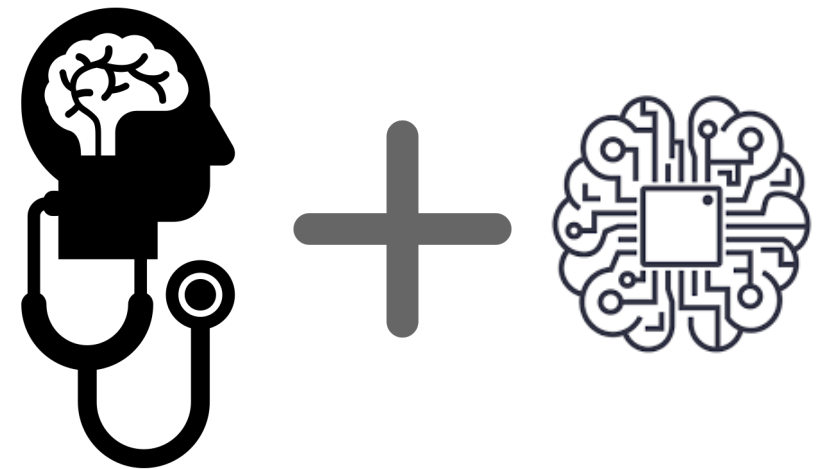
# Should we believe this?

- We don't **really** know what would have happened if our recommendations would have been followed
- The only way to truly know if our recommendations are useful: Run an experiment testing the system with real physicians and patients
- The goal of our framework is to come in the best possible safe shape towards such a trial
- **Currently planning the trial**



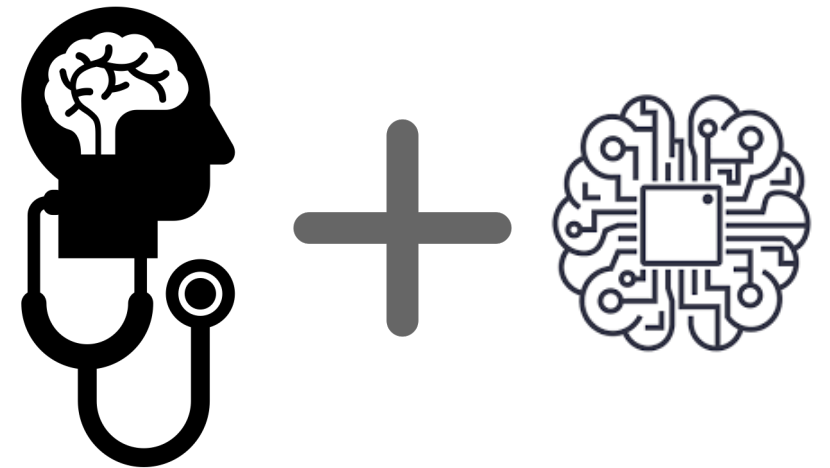
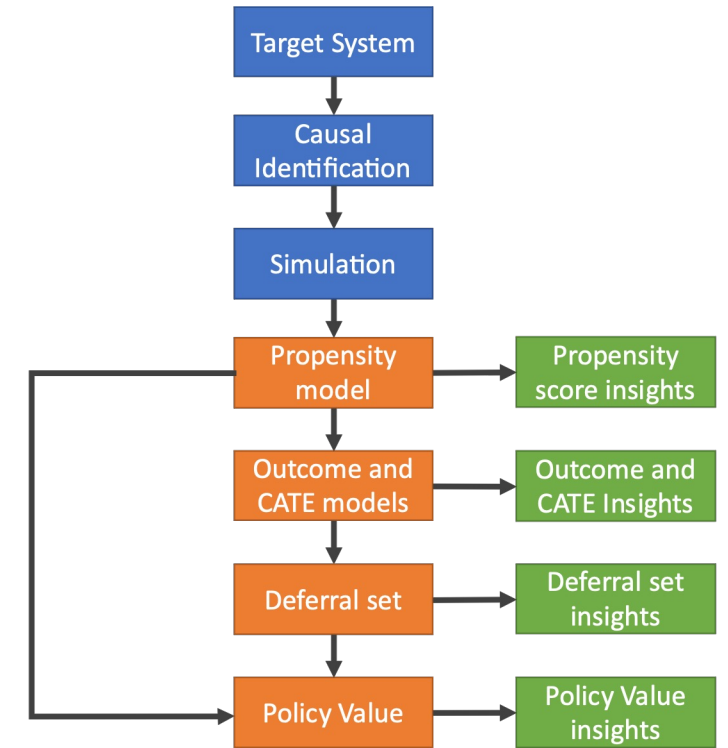
# Planning an “AI@hospital trial”

- Rambam Healthcare Campus:
  - 1,000 bed tertiary hospital
  - serving a population of 2,000,000 people
- TERA: Technion-Rambam Initiative in Medical AI
  - Jointly funded center
  - Money, clinician time, and space for joint research
  - Expedited access to data (regulatory and technical)
  - Support for deployment at bedside
- In practice
  - Strong clinical-computational collaboration on a personal level is key
  - Joint commitment to goal and willingness to invest time & energy in it (especially clinician’s time!)



# Planning an “AI@hospital trial”

- *Now funded for trial*
- Plan:
  - Intense discussions with entire clinical team
  - Deepen understanding of clinical workflow
  - Where does the system come in?
  - Push or pull?
  - Unit of Randomization?
    - “Defer” or “Run Model”
- Main outcomes
  - Clinician acceptance, adherence, reaction
  - Safety
  - Kidney function



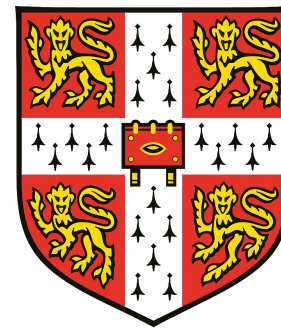
• **Looking forward to your thoughts and comments!**

# We are hiring!

- Joint PhD / Postdoc with Prof. Mihaela van der Schaar at Cambridge University
- Work on methods for causal inference and machine learning in healthcare
- Email [shalit-lab@technion.ac.il](mailto:shalit-lab@technion.ac.il)



**TECHNION**  
Israel Institute  
of Technology



**UNIVERSITY OF  
CAMBRIDGE**

# Thank you!

- Technion:
  - Rom Gutman
  - Shimon Sheiba
  - Omer Noy
- Rambam Health Care Campus & Technion:
  - Dr. Oren Caspi
  - Prof. Doron Aronson
- Clalit Research Institute:
  - Ohad Levinkron
  - Dr. Janni Yuval
  - Galit Shaham
  - Dr. Becca Feldman
  - Prof. Ran Balicer