

# Causal Falsification of Digital Twins

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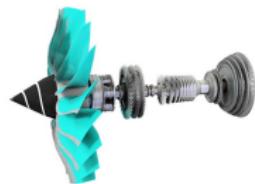
June 25, 2023

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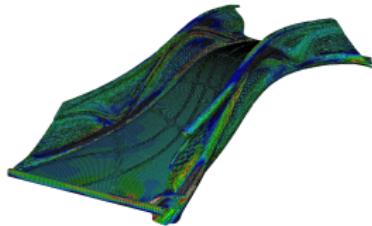
\*Equal contribution

# Motivation

Simulators called **Digital Twins** are increasingly used to guide **safety-critical** decision-making



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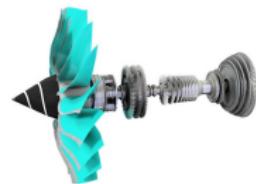
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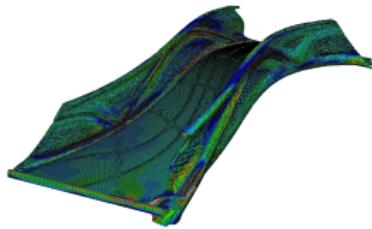
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In these environments, the **accuracy** of a twin is paramount

# High-level goal

Our question: Often **large datasets** taken from the underlying phenomena are available

How can we use this data to **assess the accuracy** of a given twin?

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How can we use this data to **assess the accuracy** of a given twin?

Constraints: Assessment procedure itself must be reliable:

⇒ Prefer **soundness** over completeness

Want a procedure that can **realistically** scale to real twins

⇒ Want to make **minimal assumptions**

# Key insight and challenge

A natural approach is to **compare directly** the output of the twin with observational data

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However, if **causal** conclusions are sought (e.g. for planning), then this is **unsound** for most datasets in practice

## Motivating example

## Toy scenario

Consider modelling effect of **drug** on **weight** for some population

Drug interacts with an **enzyme**  $U \in \{0, 1\}$  present in a subpopulation:

- If  $U = 1$ , drug increases weight
- If  $U = 0$ , drug has no effect

Suppose drug is **only administered** when  $U = 1$

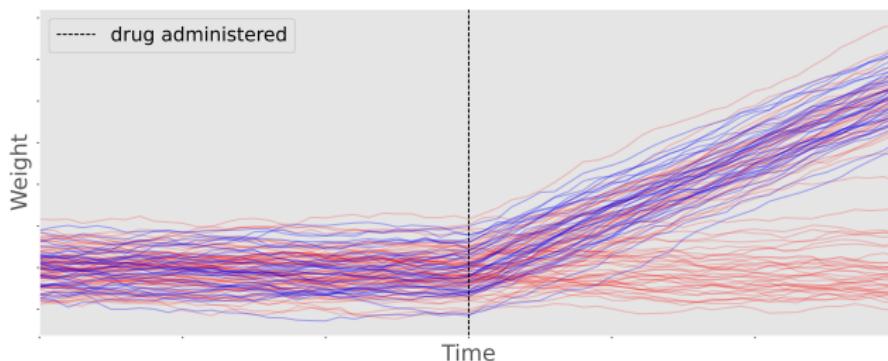
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Blue: outcomes that *were observed* for patients administered drug;  
Red: outcomes that *would be observed* across whole population

## Key point

This phenomenon occurs because the data are **confounded**

Confounding is well-studied in the causal inference literature

However, **implications for simulators** are less appreciated

Key point: in general **wrong** to compare the data with the output of twin under the corresponding actions

# Overview

Motivated by this observation, our paper:

- Formulates twin assessment as a **causal inference** problem
- Argues for an approach based on **falsification** rather than **verification**
- Presents a **statistical methodology** valid under **minimal assumptions**
- Illustrates via a large-scale **case study**

## Aside: Causal Inference

# Overview

Causal inference provides a mathematical framework for reasoning about the causal effects of interventions based on observational data

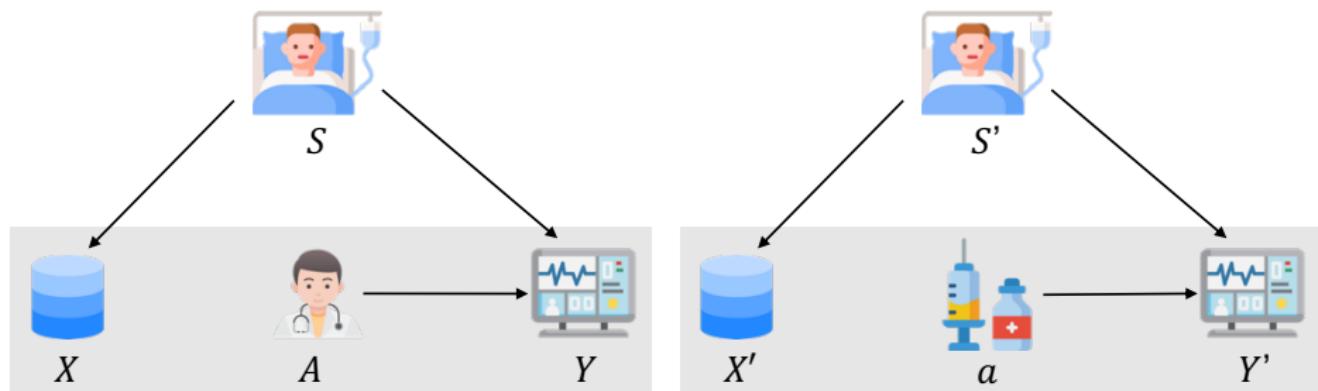
Many questions we care about in practice are of a causal nature

- “What should I do to make things a certain way?” vs. “How do things evolve on their own?”

For this reason, highly suitable for Twins, for which decision-making and acting in the world are primary concerns

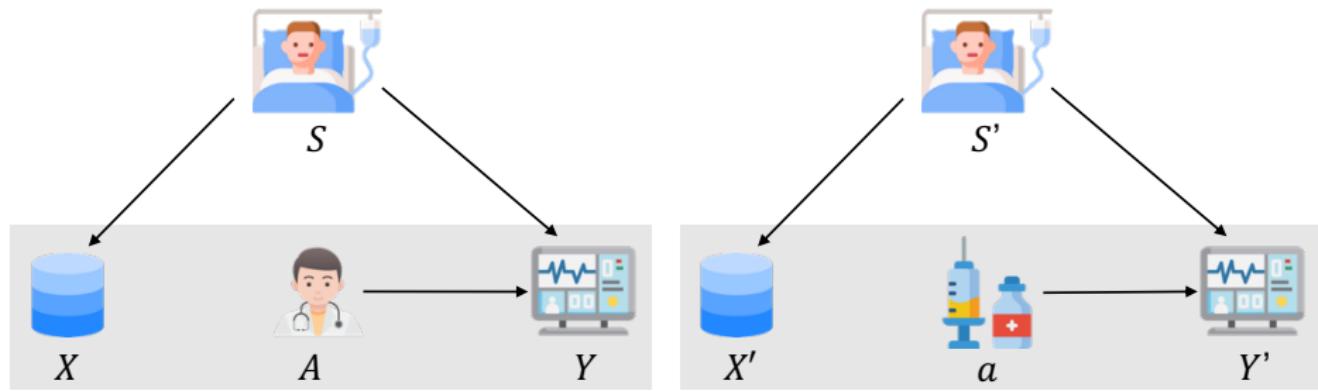
# A Typical Problem

**Straightforward problem:** Given distribution of  $(X, A, Y)$  from the left-hand system, what is distribution of  $(X', Y')$  in the right-hand system?



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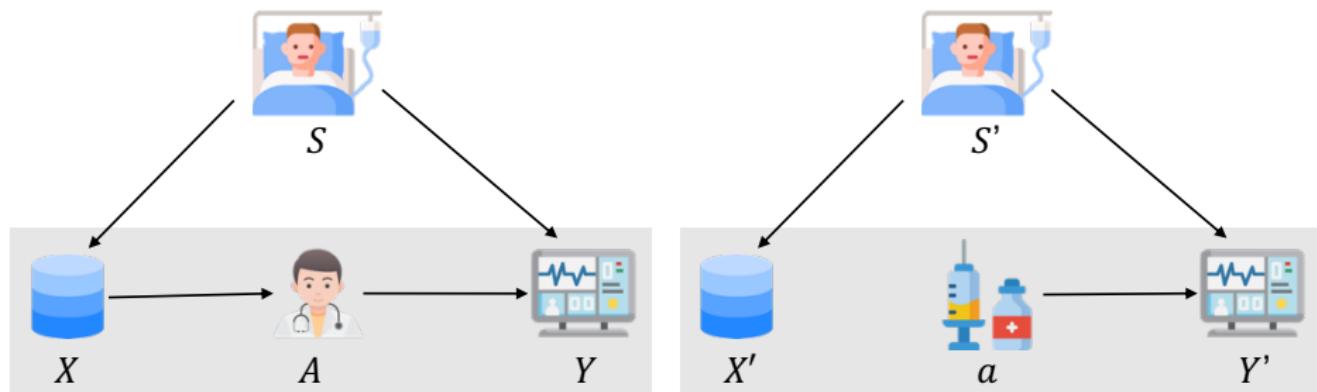
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**Answer:**  $P(X' = x, Y' = y)$  on right is  $P(X = x, Y = y \mid A = a)$  on left

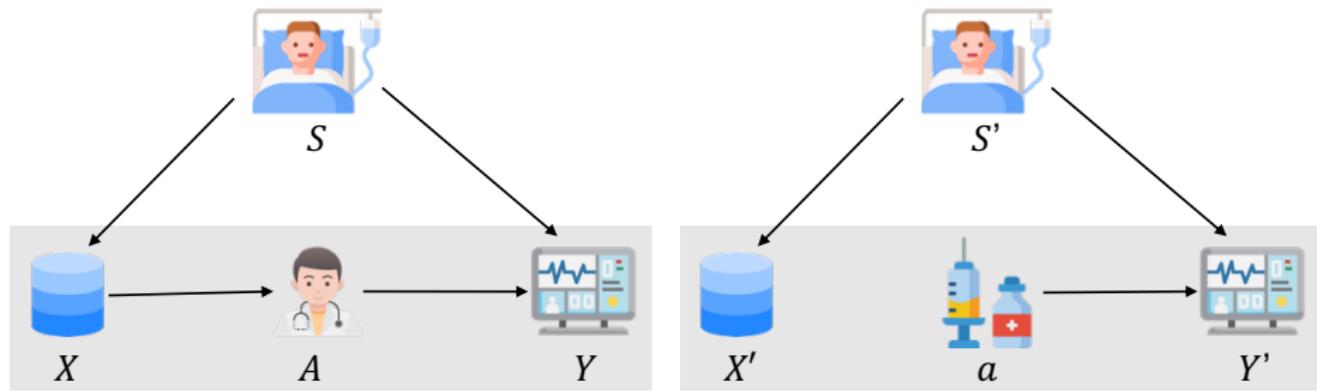
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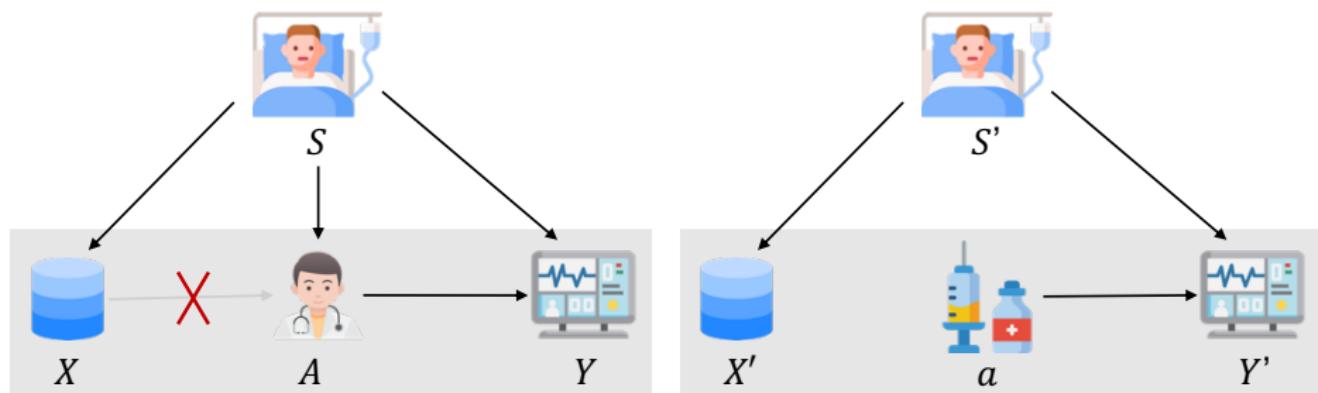


**Answer:**

$P(X' = x, Y' = y)$  on right is  $P(X = x) P(Y = y | X = x, A = a)$  on left  
 $(\neq P(X = x, Y = y | A = a))$

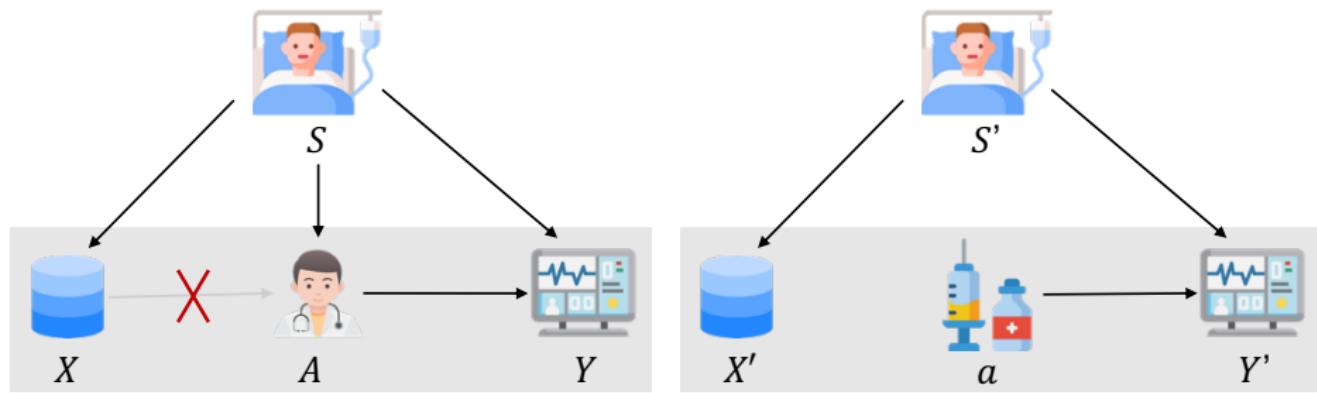
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**Answer:** Don't know! (without further assumptions)

# Unmeasured confounding

In last case, the data contains **unmeasured confounding** (cf. second case)

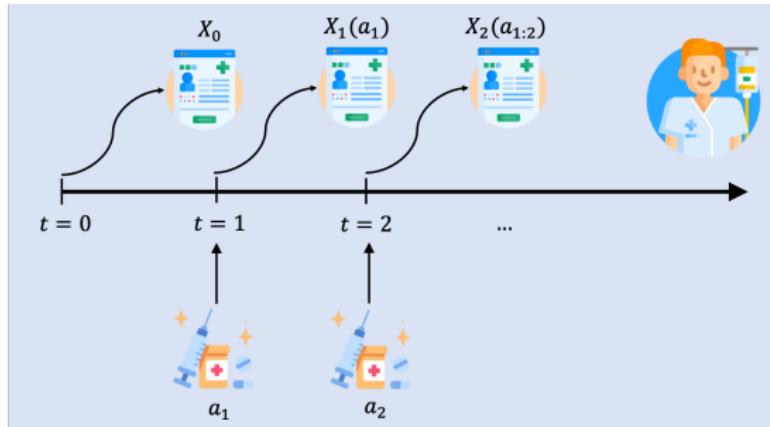
Unmeasured confounding is usually assumed away, but it is in fact **extremely common** (e.g.  $U$  as enzyme from earlier)

For no unmeasured confounding, **every factor** that affects both  $A$  and  $Y$  must be included explicitly in the data

- Often **tenuous**, especially for safety-critical applications

# Our Problem Setup

# Real World Process

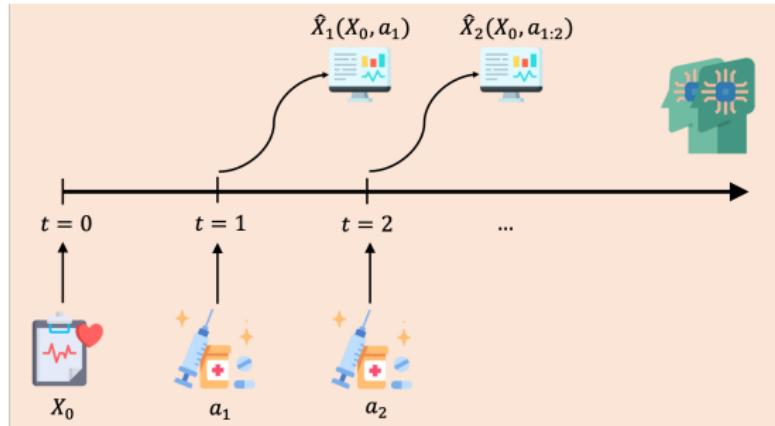


Model **real-world process** via potential outcomes:

$X_0, X_1(a_1), X_2(a_{1:2}), \dots, X_T(a_{1:T})$  for each sequence  $a_{1:T}$  of **actions**.

Idea:  $X_t(a_{1:t})$  represents what **would** be observed after actions  $a_{1:t}$

# Digital Twin Process

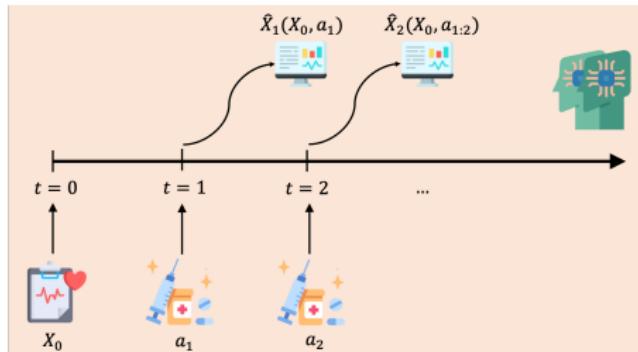
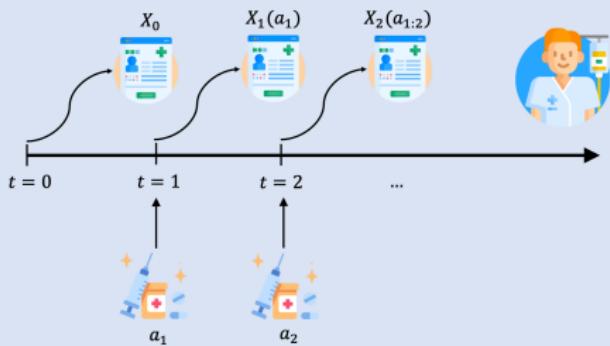


Model **twin** similarly as

$\hat{X}_1(x_0, a_1), \dots, \hat{X}_T(x_0, a_{1:T})$  where additionally  $x_0$  is an **initialisation**

Idea:  $\hat{X}_t(x_0, a_{1:t})$  represents output of twin after inputs  $x_0$  and  $a_{1:t}$

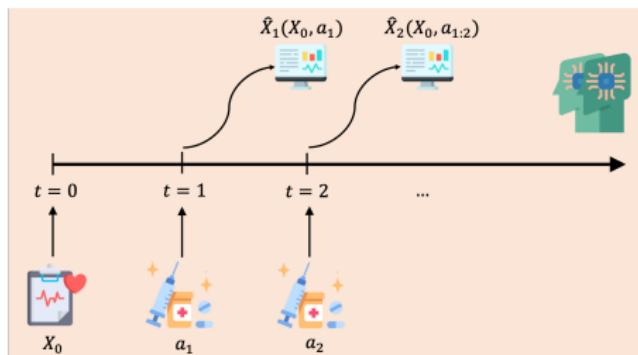
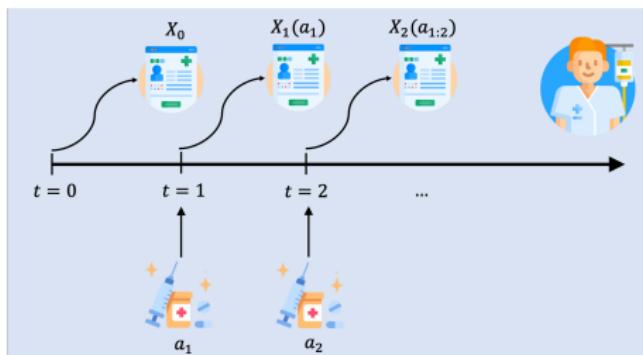
# Interventional Correctness



## Interventional correctness

Would like the distribution of each  $\hat{X}_{1:t}(x_0, a_{1:t})$  to be equal to the conditional distribution of  $X_{1:t}(a_{1:t})$  given  $X_0 = x_0$

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⇒ Can recover real-world distribution via Monte Carlo (e.g. for **planning**)

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Overall model is intentionally very weak, which seems appropriate for the assessment problem

- Do not assume  $X_t(a_{1:t}) \perp\!\!\!\perp A_t \mid X_{0:t-1}(A_{1:t-1}), A_{1:t-1}$  (sequential randomisation assumption, i.e. no unmeasured confounding)

## Verification and falsification

# Verification approaches

Standard assessment approaches have the following logical structure:

## Verification assessment

- ① Choose a **hypothesis**  $\mathcal{H}$  such that, if  $\mathcal{H}$  is true, then the twin is correct
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## Theorem

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⇒ Does not exist  $\mathcal{H}$  with this property whose truth can be determined from the **data alone**

# Our alternative: falsification

We consider the following **alternative structure**:

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Advantage: **can** choose  $\mathcal{H}$  with this property whose falsity can be determined from data

However: lack of falsification does not imply the twin is correct

## Hypotheses from causal bounds

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$$Y_{\text{lo}} := \mathbb{I}(A_{1:t} = a_{1:t}) Y(A_{1:t}) + \mathbb{I}(A_{1:t} \neq a_{1:t}) y_{\text{lo}}$$

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## Theorem (Causal bounds)

If  $\mathbb{P}(y_{\text{lo}} \leq Y(a_{1:t}) \leq y_{\text{up}} \mid X_{0:t}(a_{1:t}) \in B_{0:t}) = 1$ , then

$$\begin{aligned} \mathbb{E}[Y_{\text{lo}} \mid X_{0:N}(A_{1:N}) \in B_{0:N}] &\leq \mathbb{E}[Y(a_{1:t}) \mid X_{0:t}(a_{1:t}) \in B_{0:t}] \\ &\leq \mathbb{E}[Y_{\text{up}} \mid X_{0:N}(A_{1:N}) \in B_{0:N}]. \end{aligned}$$

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Key point: left and right-hand sides are **identifiable** (in fact, **unbiasedly**) from observational data

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Corresponds to Manski [1990] (cf. Zhang and Bareinboim [2019])

# Optimality of bounds

- Without further assumptions, these bounds **cannot be improved upon** for general  $Y(a_{1:t})$  (or if  $Y(a_{1:t}) = f(X_t(a_{1:t}))$ )

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- Also, cannot bound  $\mathbb{E}[Y(a_{1:t}) \mid X_{0:t}(a_{1:t})]$  nontrivially if  $X_{1:t}(a_{1:t})$  is **continuous**

## Derived hypotheses

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Let  $Q_{\text{lo}}$  and  $Q_{\text{up}}$  be causal bounds from earlier

⇒ If the twin is interventionally correct, then  $\mathcal{H}_{\text{lo}}$  and  $\mathcal{H}_{\text{up}}$  hold, where

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Interpretation: (e.g.) if  $\mathcal{H}_{\text{lo}}$  is false, then when  $(X_0, \hat{X}_{1:t}(X_0, a_{1:t})) \in B_{0:t}$ , the outputs  $f(X_0, \hat{X}_{1:t}(X_0, a_{1:t}))$  are on average **too small**

# Statistical methodology

## High-level overview

Consider testing a given  $\mathcal{H}_{\text{lo}} : Q_{\text{lo}} \leq \hat{Q}$

Recall: we have an **observational dataset** of i.i.d. copies of

$$X_0, A_1, X_1(A_1), A_2, X_2(A_{1:2}), \dots, A_T, X_T(A_{1:T}).$$

For given  $a_{1:t}$ , **generate** dataset of i.i.d. copies of

$$X_0, \hat{X}_1(X_0, a_1), \dots, \hat{X}_t(X_0, a_{1:t})$$

# High-level overview

Consider testing a given  $\mathcal{H}_{\text{lo}} : Q_{\text{lo}} \leq \hat{Q}$

Recall: we have an **observational dataset** of i.i.d. copies of

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For given  $a_{1:t}$ , **generate** dataset of i.i.d. copies of

$$X_0, \hat{X}_1(X_0, a_1), \dots, \hat{X}_t(X_0, a_{1:t})$$

Use e.g. Hoeffding's inequality to obtain one-sided conf. intervals  $R_{\text{lo}}^\alpha$ ,  $\hat{R}^\alpha$ ,

$$\mathbb{P}(Q_{\text{lo}} \geq R_{\text{lo}}^\alpha) \geq 1 - \frac{\alpha}{2} \quad \mathbb{P}(\hat{Q} \leq \hat{R}^\alpha) \geq 1 - \frac{\alpha}{2}$$

and **reject**  $\mathcal{H}_{\text{lo}}$  if  $\hat{R}^\alpha < R_{\text{lo}}^\alpha$ , or return a **p-value**

## Other aspects

Control for **multiple testing** via e.g. Holm-Bonferroni or Benjamini-Yekutieli

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Can choose parameters  $(t, f, a_{1:t}, B_{0:t})$  for each  $\mathcal{H}_{\text{lo}}$  and  $\mathcal{H}_{\text{up}}$  in a data-dependent way, provided we use **sample splitting**

- Useful e.g. for  $y_{\text{lo}}$  and  $y_{\text{up}}$

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- Useful e.g. for  $y_{\text{lo}}$  and  $y_{\text{up}}$

No additional assumptions required by construction

## Case study: Pulse Physiology Engine

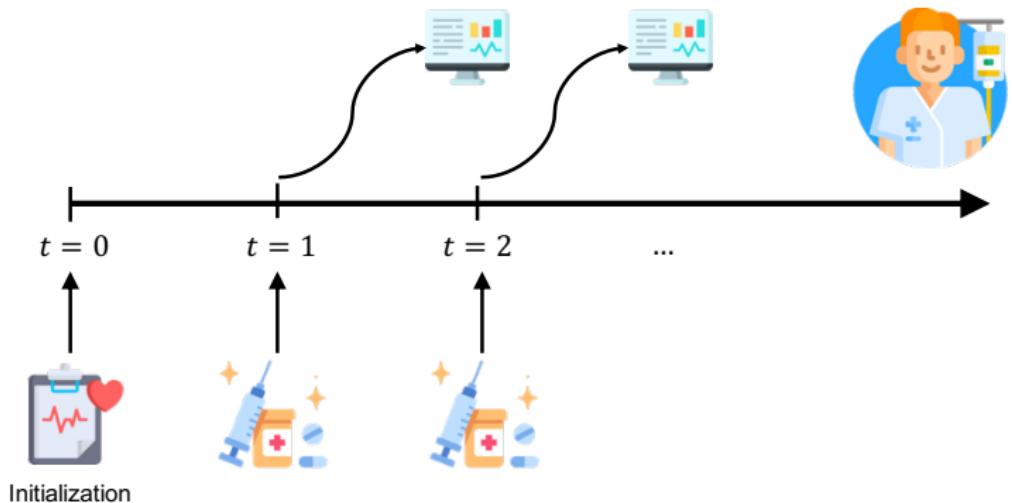
# Pulse Physiology Engine

We apply our methodology to **Pulse Physiology Engine**, an open source computational model designed for human physiology simulation

Validate using the **MIMIC-III** dataset, generated from 40,000+ ICU patients at Beth Israel Hospital



# Pulse Physiology Engine

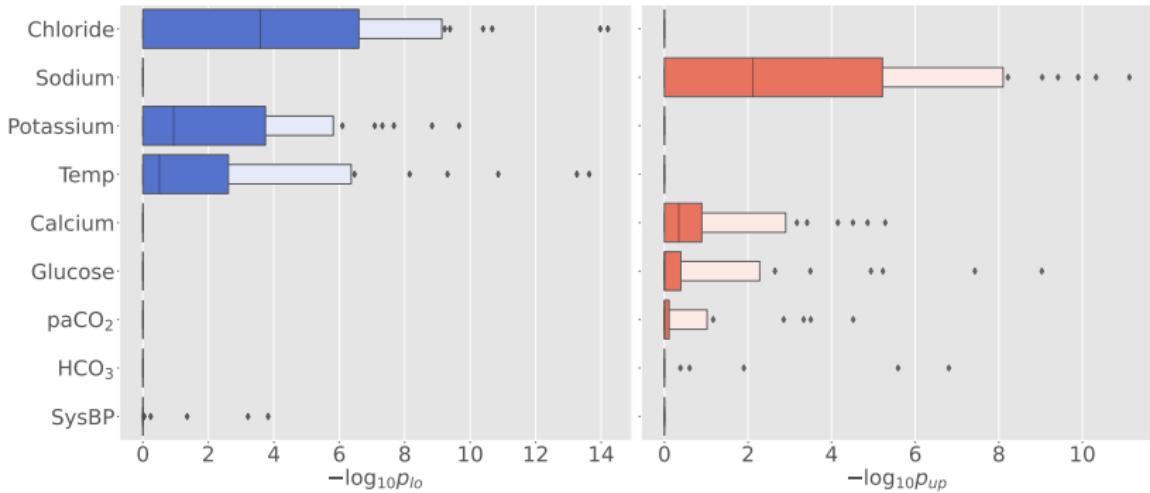


# Results

Physiological quantity	# Rejections	# Hypotheses
Chloride Blood Concentration (Chloride)	24	94
Sodium Blood Concentration (Sodium)	21	94
Potassium Blood Concentration (Potassium)	13	94
Skin Temperature (Temp)	10	86
Calcium Blood Concentration (Calcium)	5	88
Glucose Blood Concentration (Glucose)	5	96
Arterial CO <sub>2</sub> Pressure (paCO <sub>2</sub> )	3	70
Bicarbonate Blood Concentration (HCO <sub>3</sub> )	2	90
Systolic Arterial Pressure (SysBP)	2	154
Arterial O <sub>2</sub> Pressure (paO <sub>2</sub> )	0	78
Arterial pH (Arterial_pH)	0	80
Diastolic Arterial Pressure (DiaBP)	0	72
Mean Arterial Pressure (MeanBP)	0	92
Respiration Rate (RR)	0	172
Heart Rate (HR)	0	162

Table: Overall rejections (FWER = 0.05)

# Additional granularity

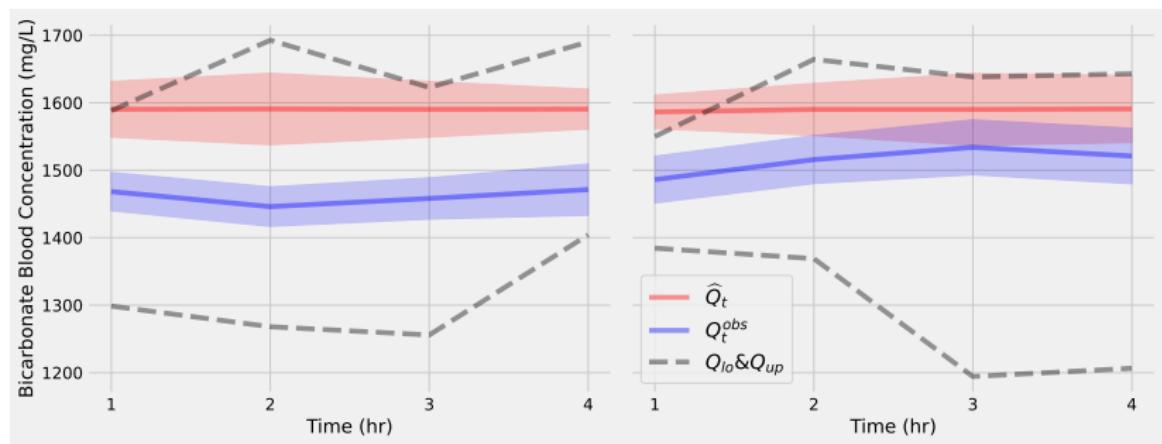


p-values for physiological quantities some rejections (notice consistent over/underestimation)

# Pitfalls of naive twin assessment

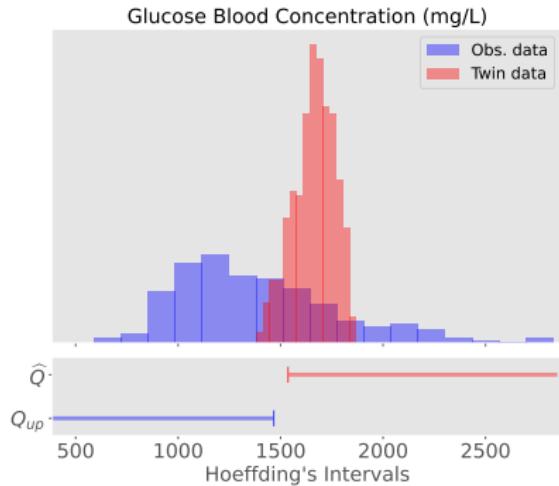
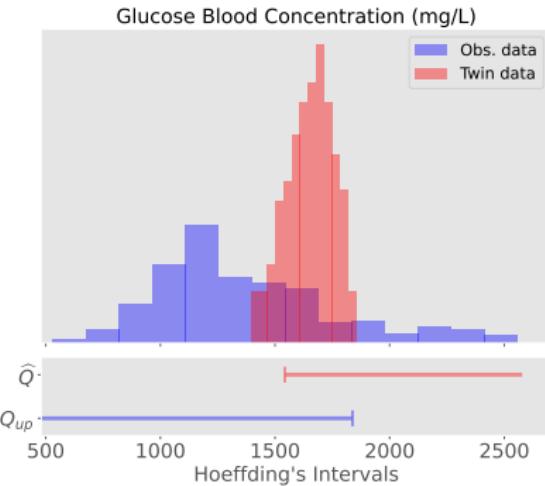
For two separate choices of  $(a_{1:t}, B_{1:t})$ , compare

$$\begin{aligned}\hat{Q}_t &:= \mathbb{E}[\hat{Y}(a_{1:t}) \mid \hat{X}_{0:t}(a_{1:t}) \in B_{0:t}], \\ Q_t^{\text{obs}} &:= \mathbb{E}[Y(A_{1:t}) \mid X_{0:t}(A_{1:t}) \in B_{0:t}, A_{1:t} = a_{1:t}].\end{aligned}$$



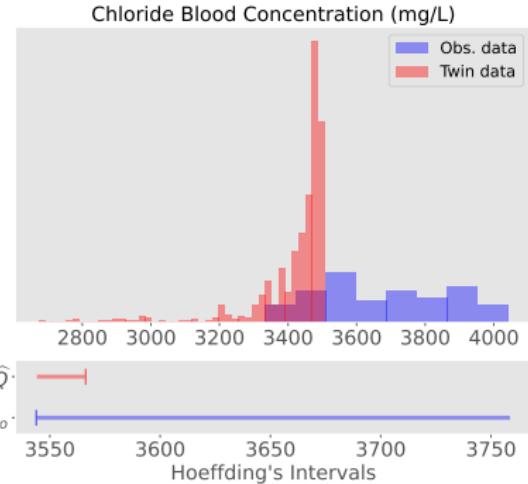
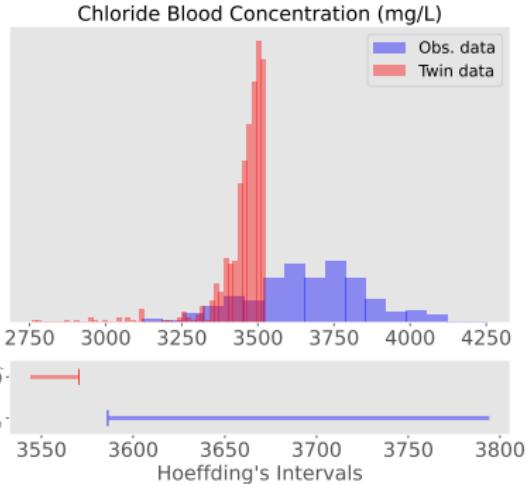
Left case looks worse, but in fact only right case leads to some rejection

## Pitfalls of naive twin assessment (2)



Despite apparent similarity, right hypothesis is rejected but left one is not

# Pitfalls of naive twin assessment (3)



Despite apparent similarity, right hypothesis is rejected but left one is not

# Thank you!



Joint work with Rob Cornish, Arnaud Doucet, and Chris Holmes

# References I

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Junzhe Zhang and Elias Bareinboim. Near-optimal reinforcement learning in dynamic treatment regimes. *Advances in Neural Information Processing Systems*, 32, 2019.