From HeartSteps to HeartBeats: Personalized Decision-making

Raaz Dwivedi

HARVARD UNIVERSITY

Massachusetts Institute of Technology

Stanford University, OIT Seminar, Jan 25

Personalized Decision-making

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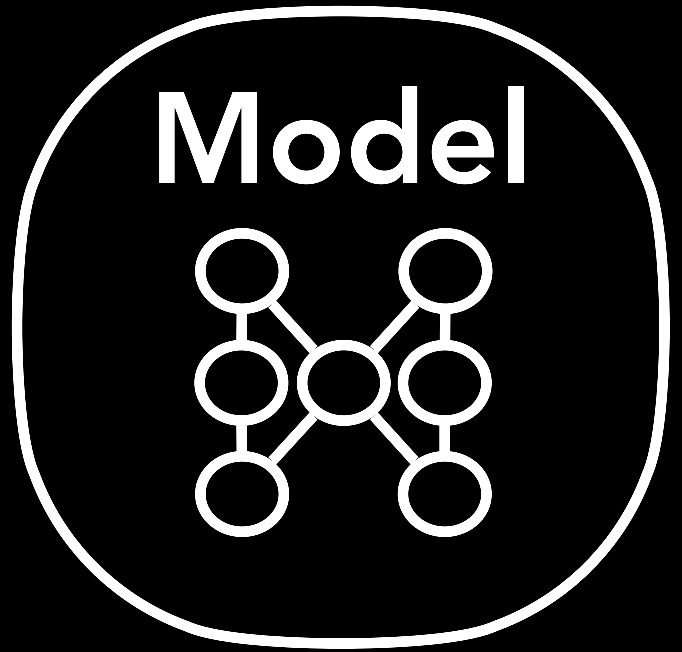
Driven by
extensive data collection,
decreasing cost of computation,
synergy between disciplines

Personalized Decision-making



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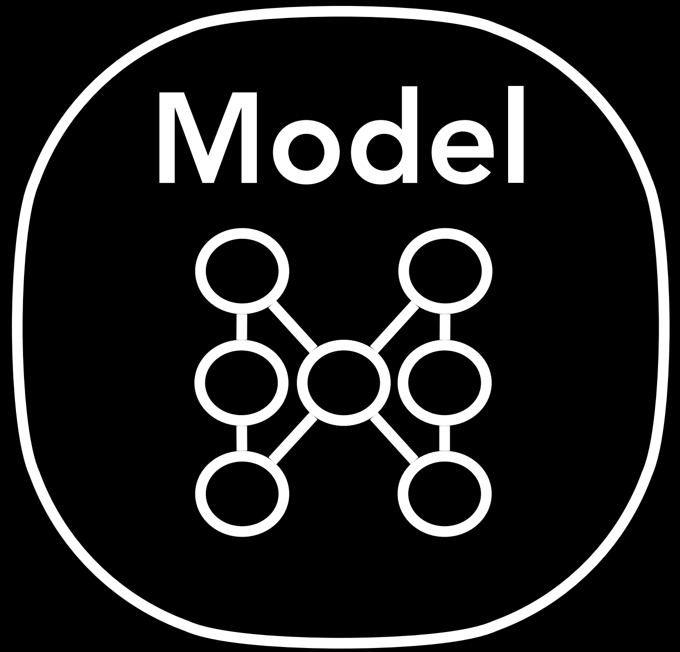

Driven by
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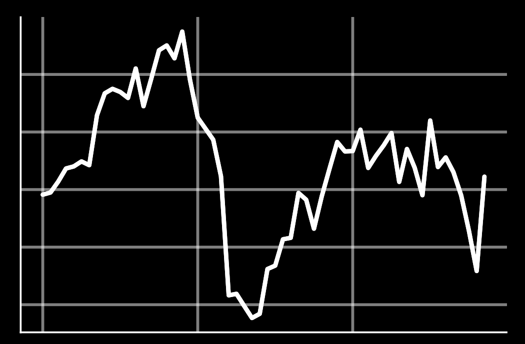
Medical records

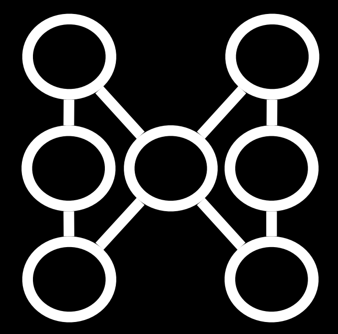
Observational studies

Data



Personalized Decision-making

Model



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Drug trial

Randomized experiments

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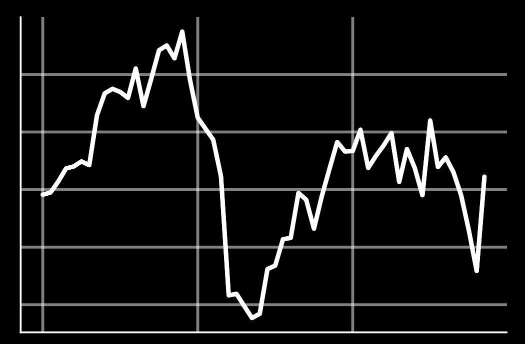
Medical records

Data



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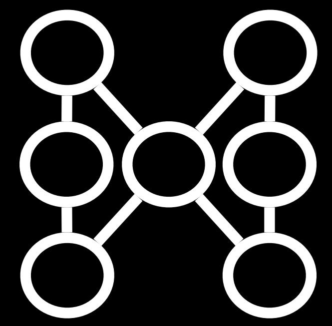

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Sequential experiments

Mobile health

Data

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1. Use **real data** to infer decision's effect

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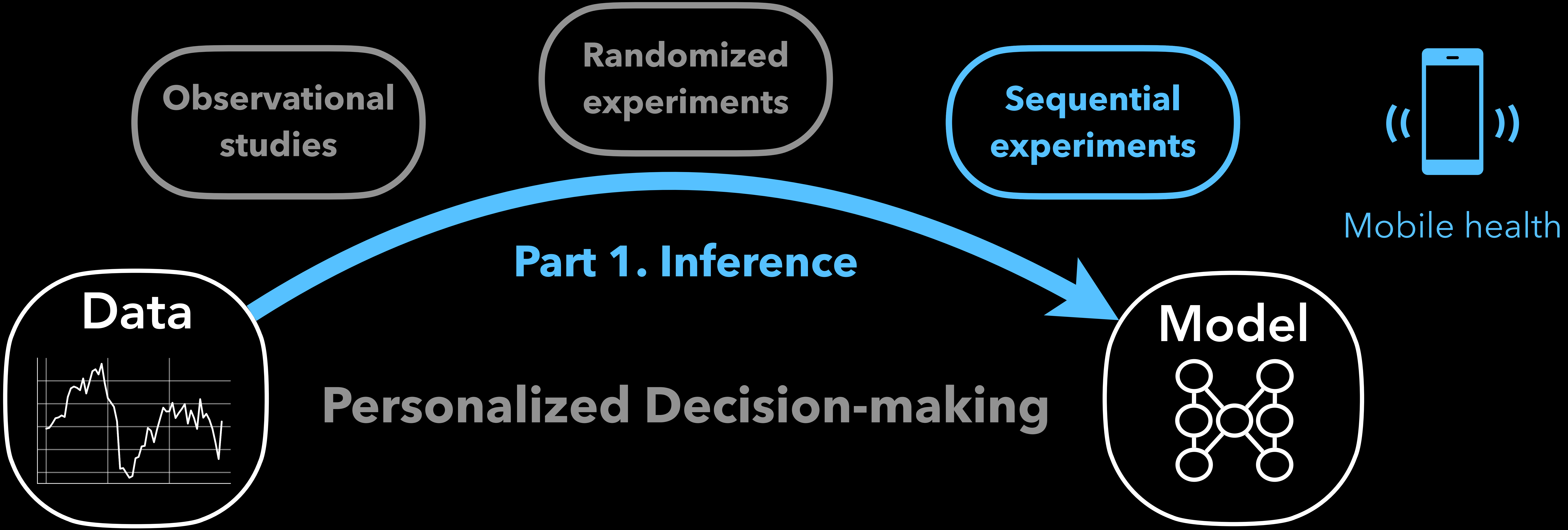

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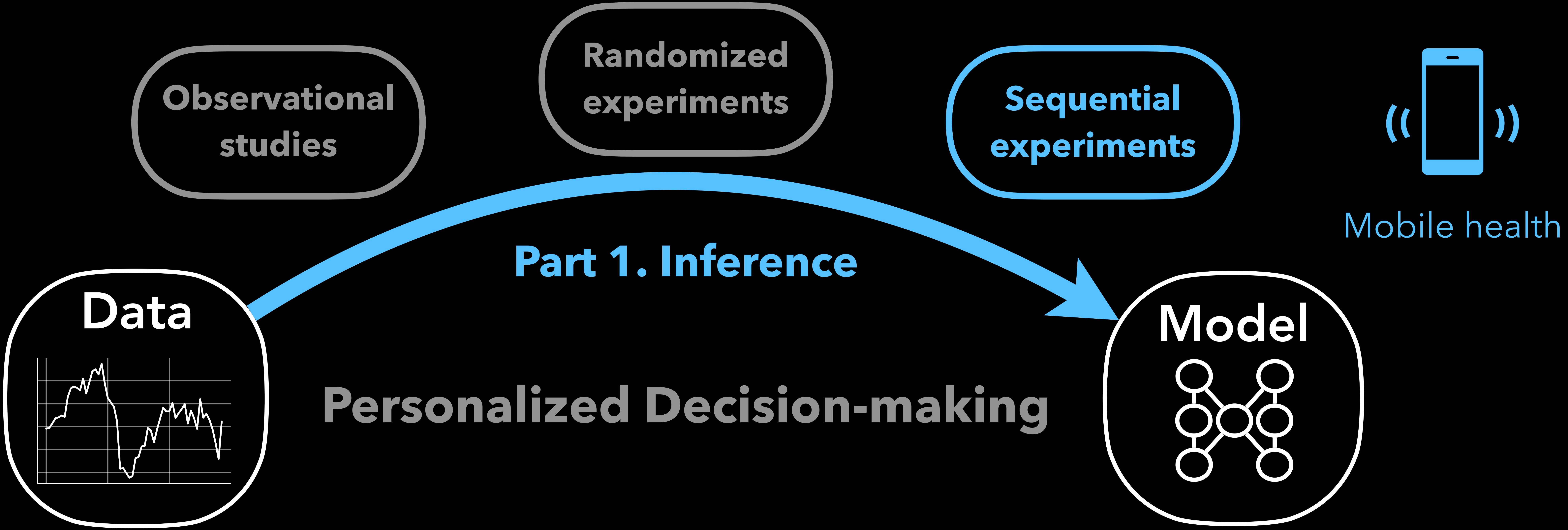

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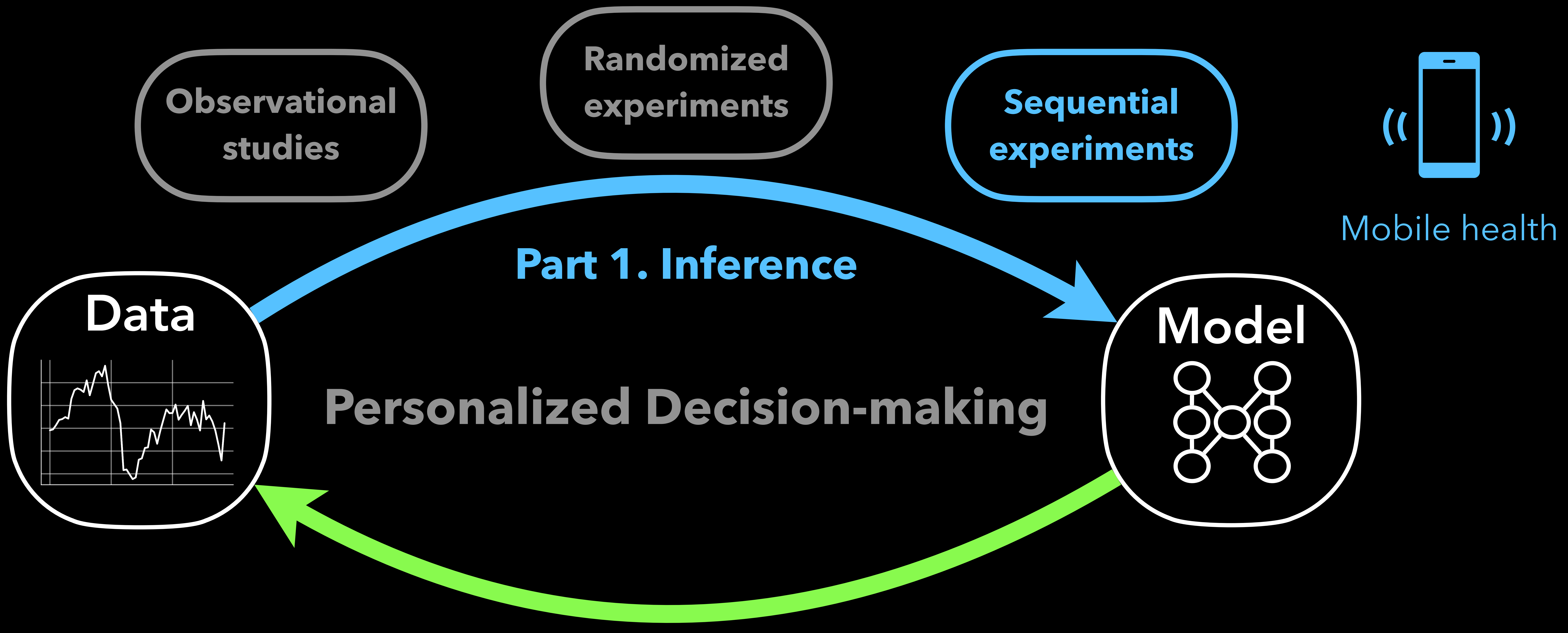

2. Use **simulated data** to predict decision's effect

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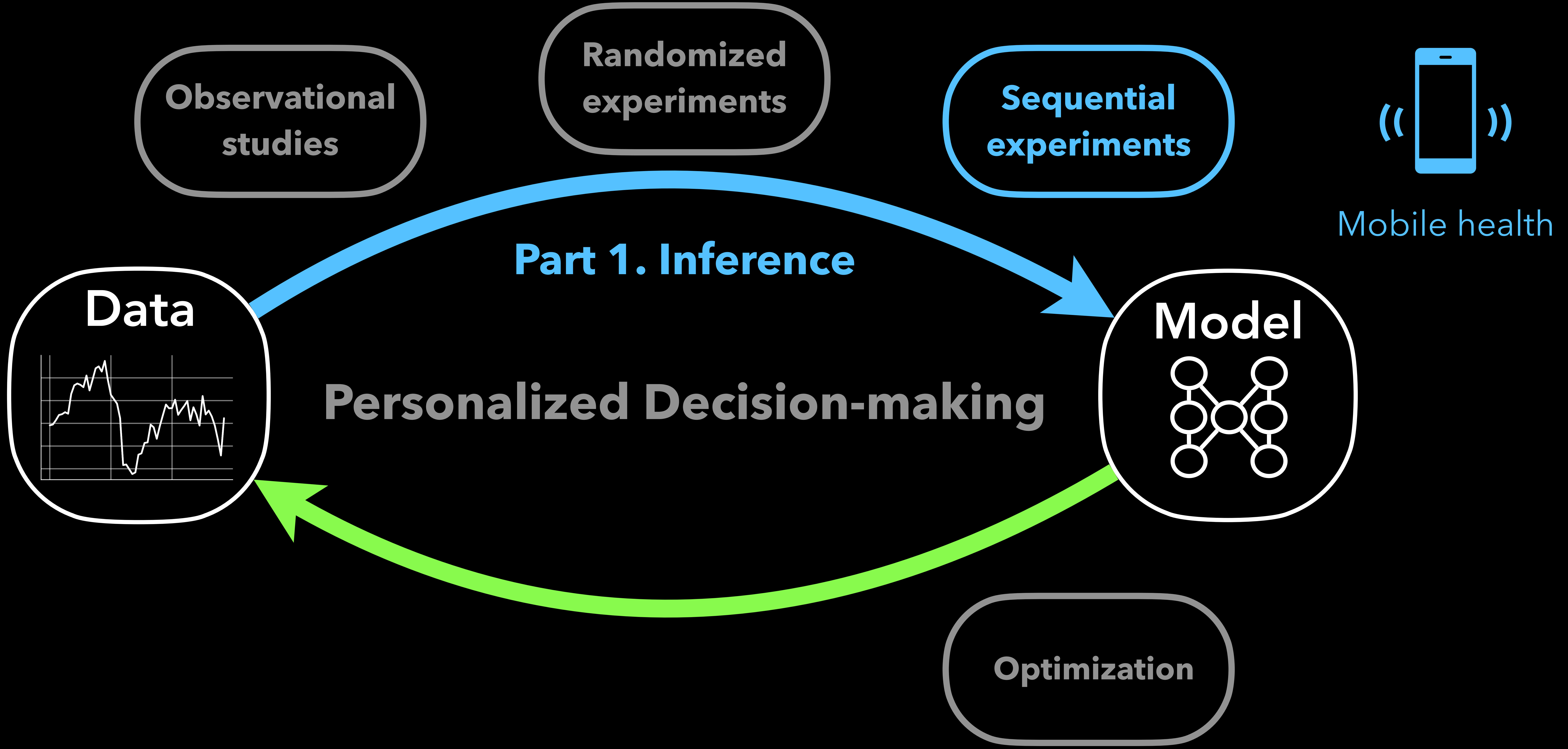

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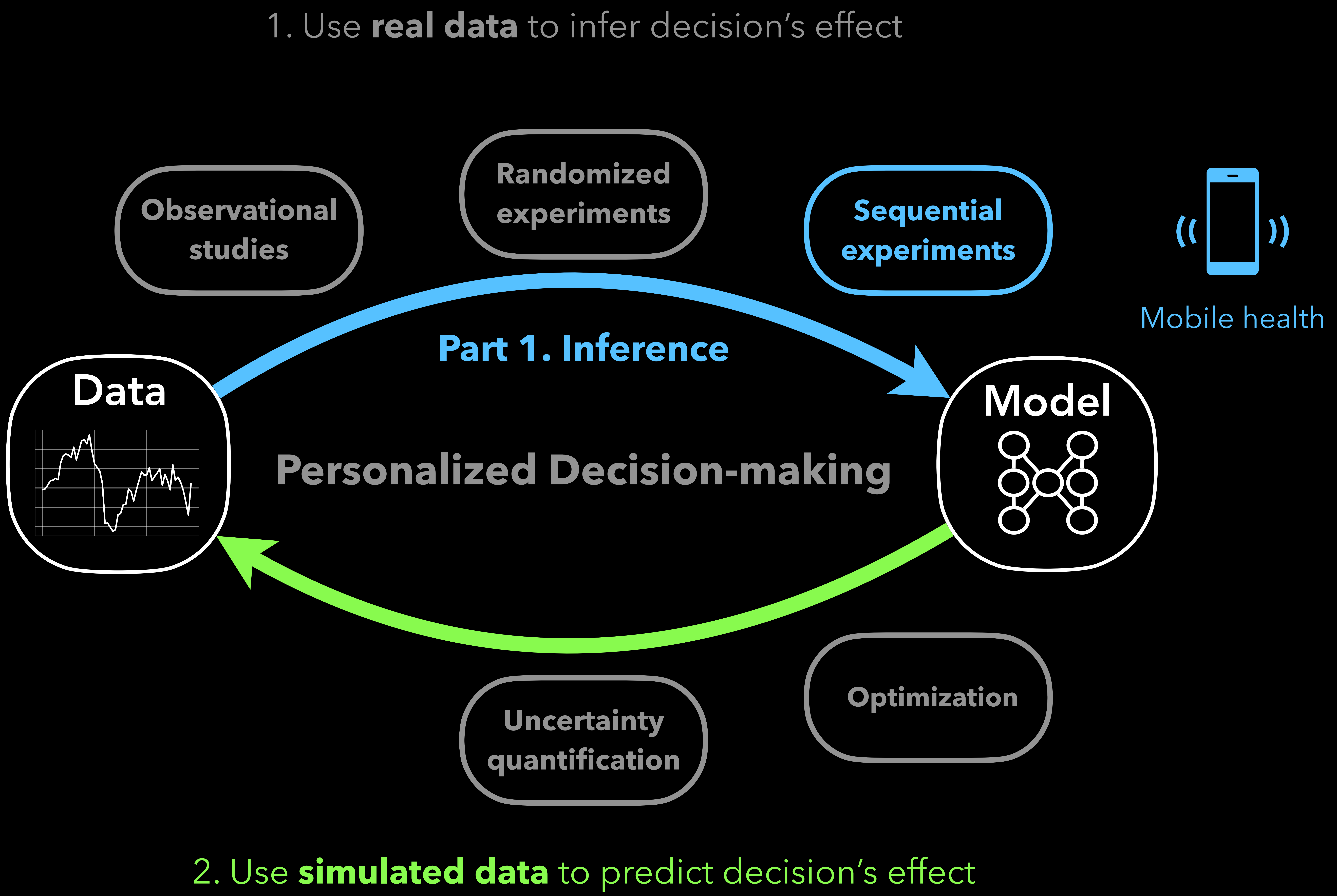
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Observational studies
Randomized experiments
Sequential experiments
Mobile health
Part 1. Inference
Data
Personalized Decision-making
Model
Uncertainty quantification
Optimization
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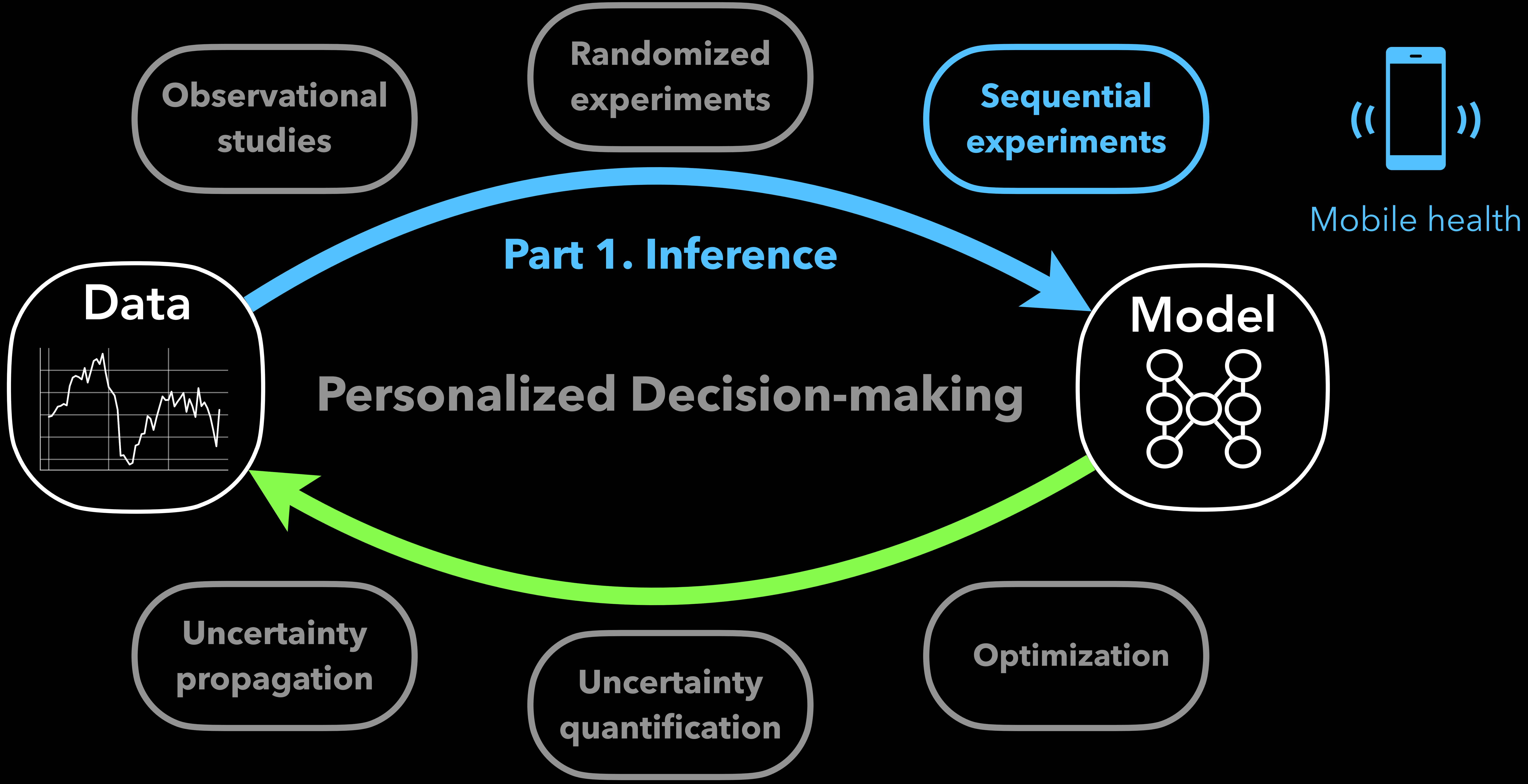

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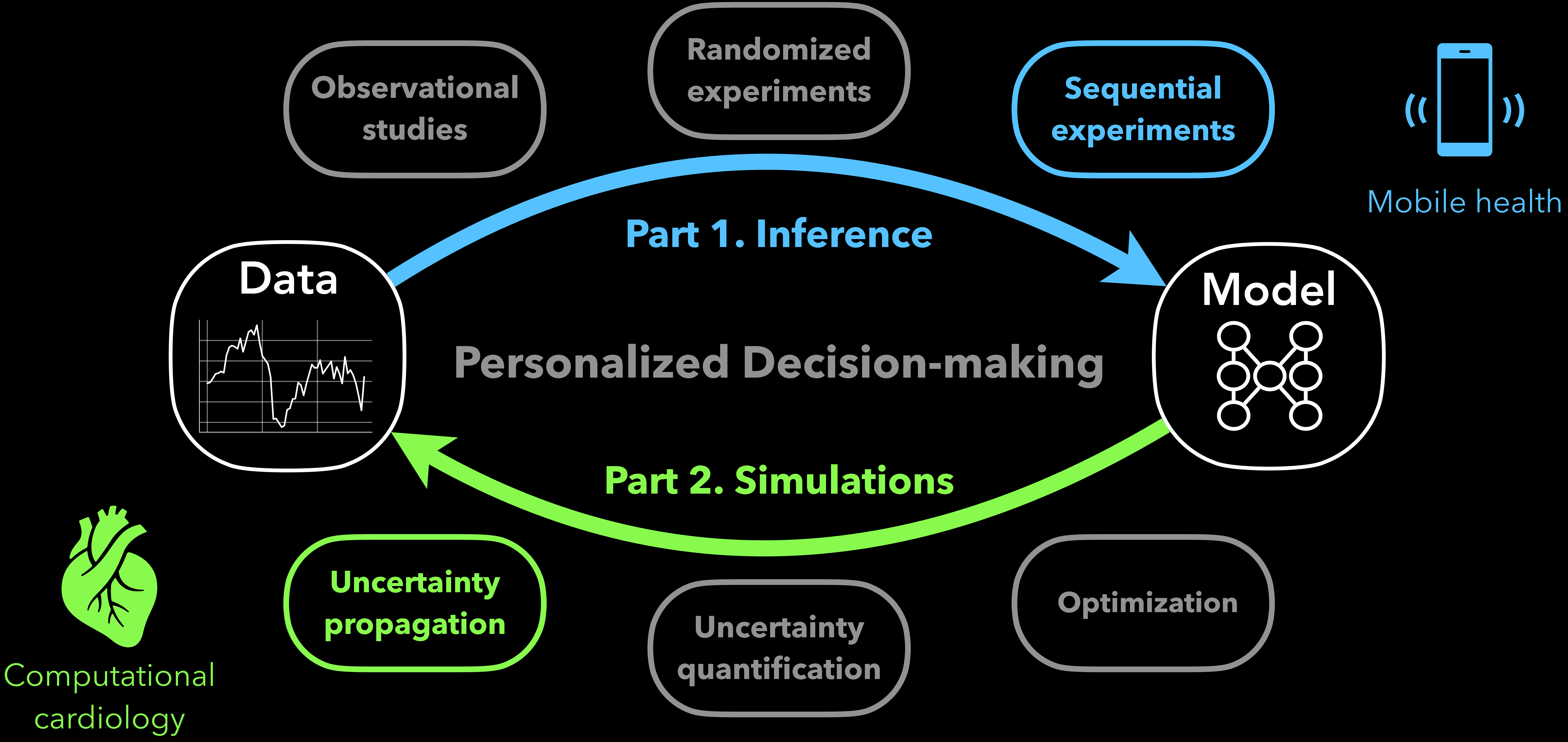

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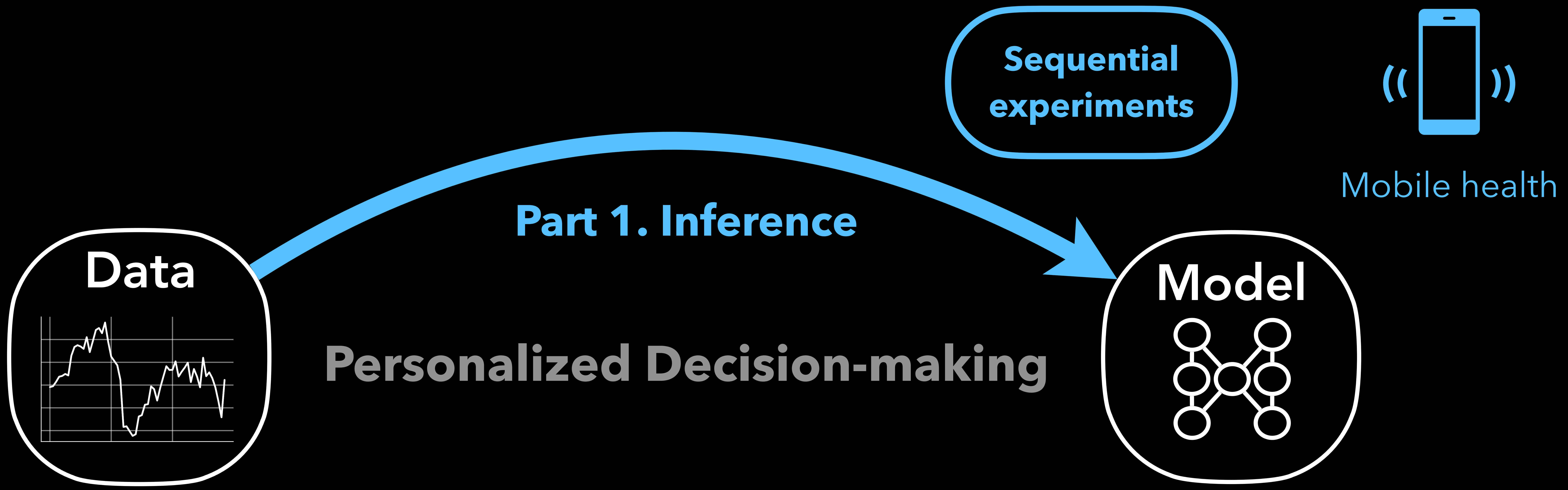
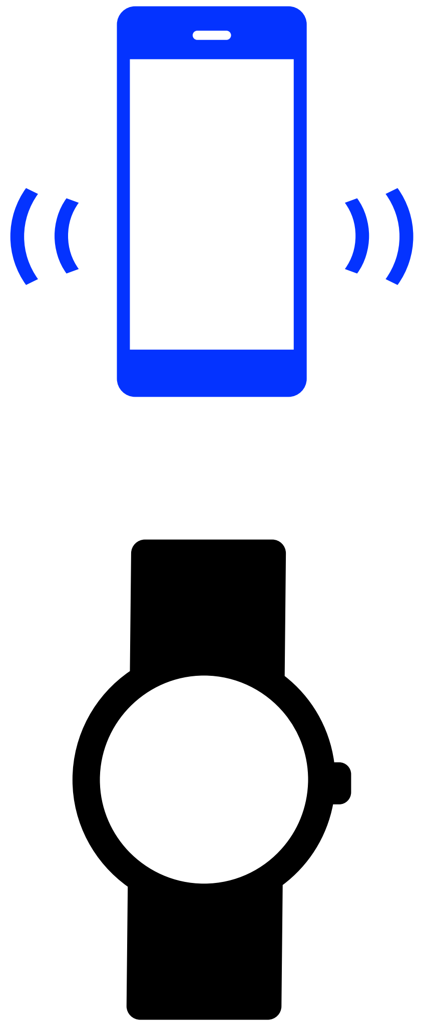
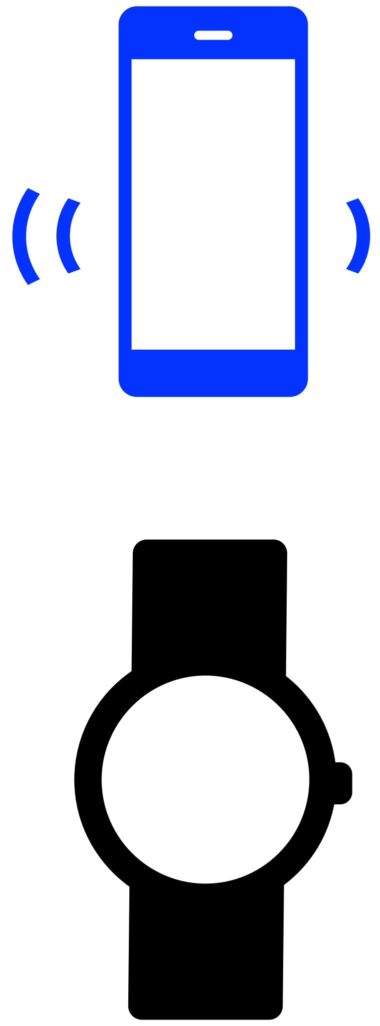


Building AI agents for personalized treatments



Building AI agents for personalized treatments

How to assign personalized digital treatments to help you?



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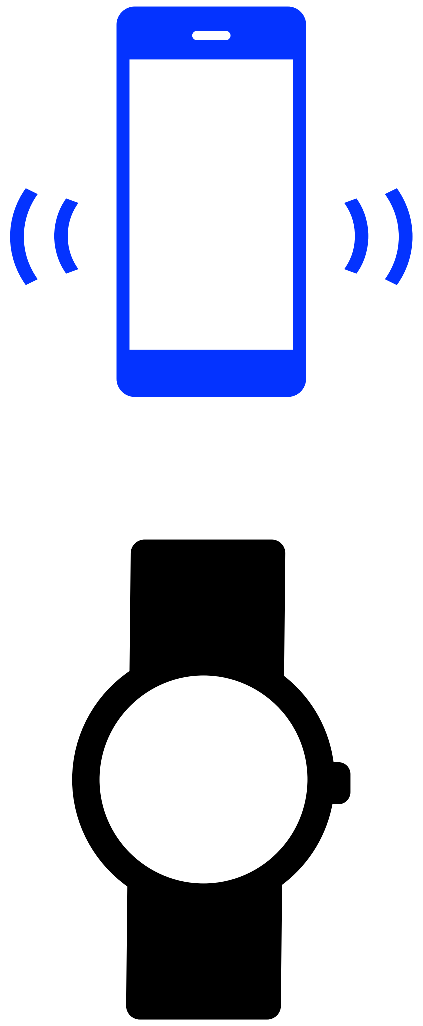

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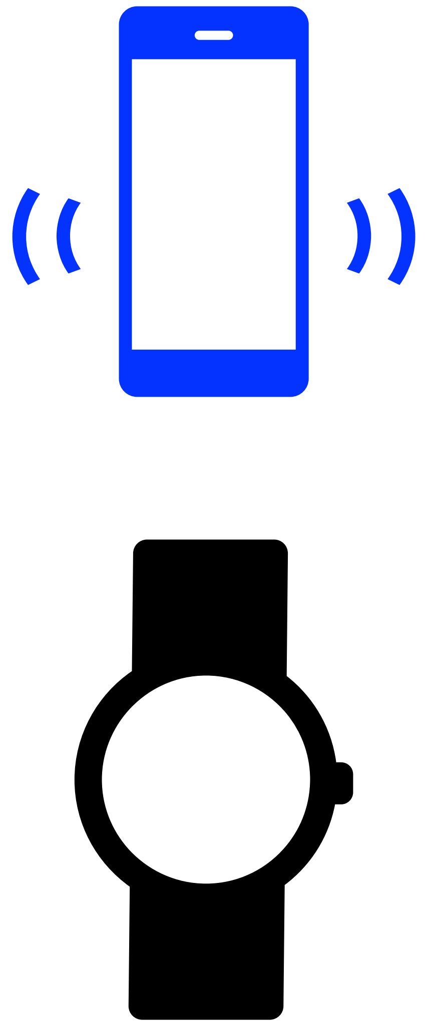

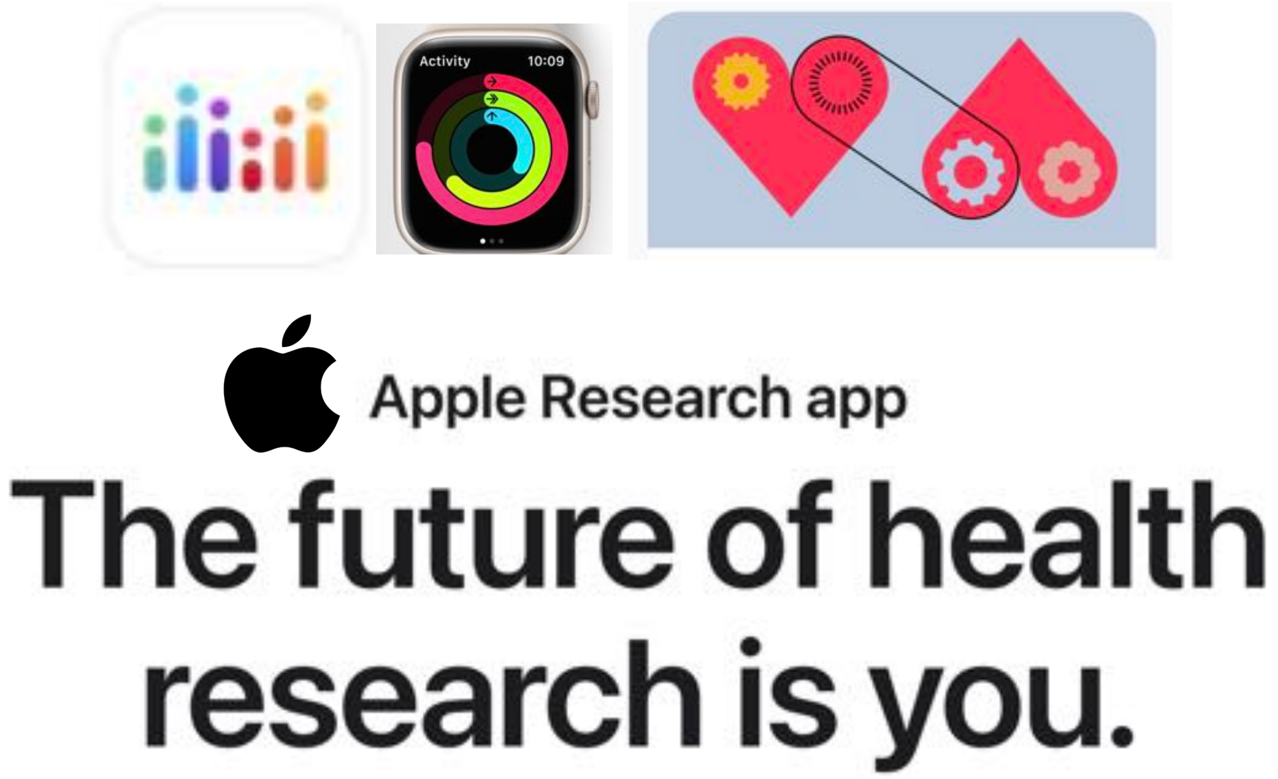

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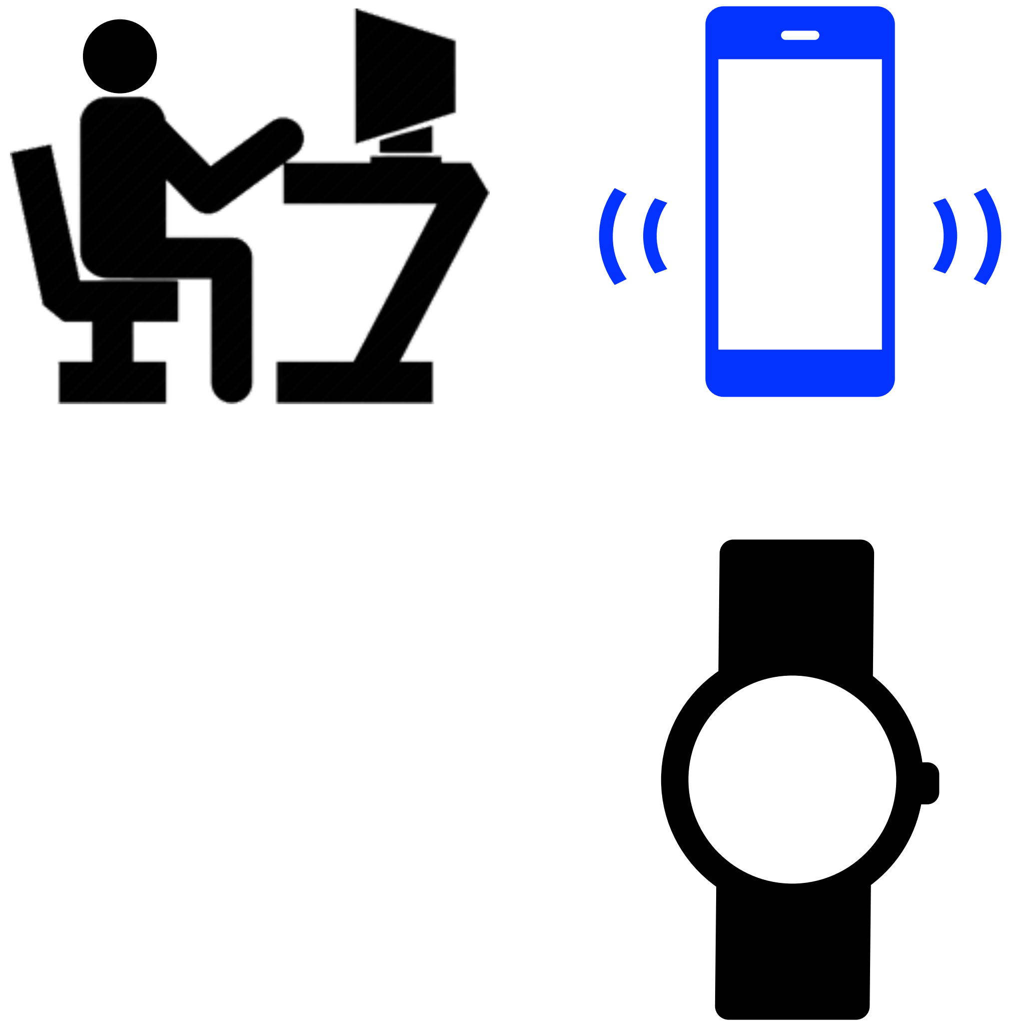
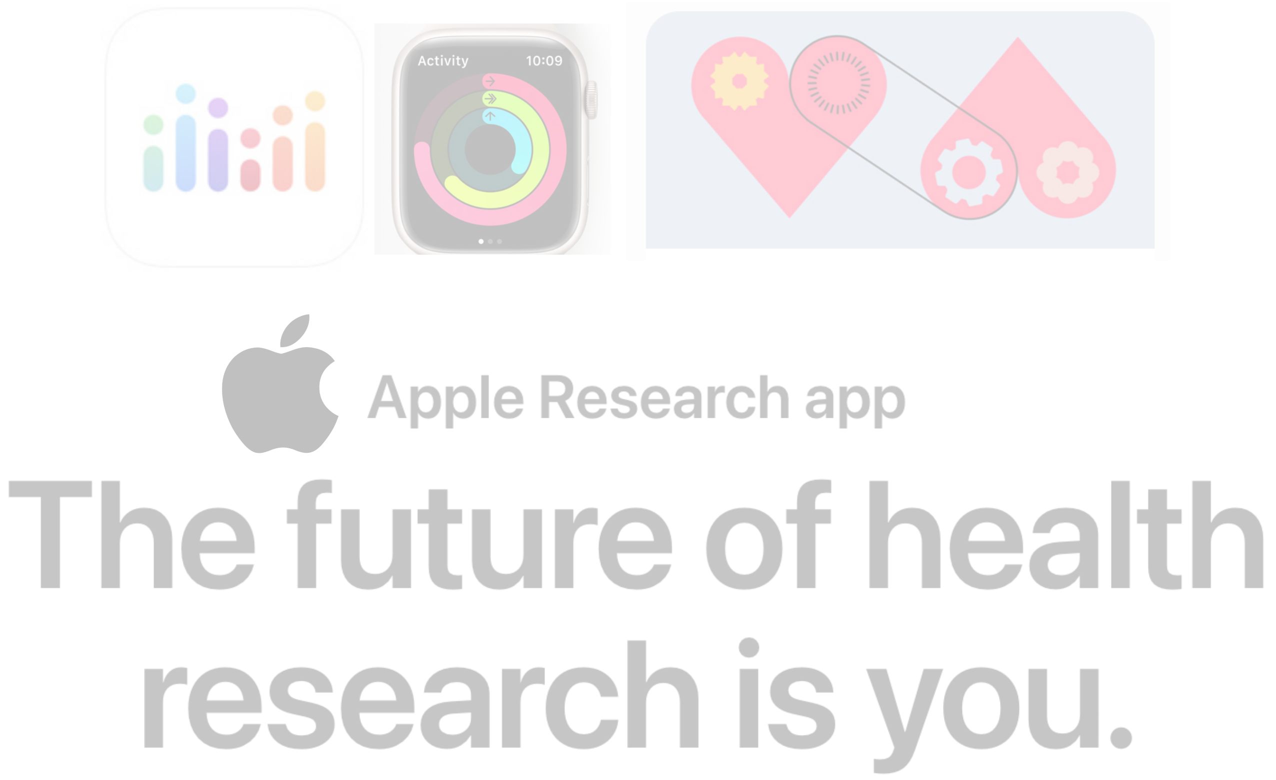


Mobile health study:
Personalized HeartSteps [Liao+ '20]

- **Goal**: Promote physical activity via mobile app
- **Population:** 91 hypertension patients, 90 days



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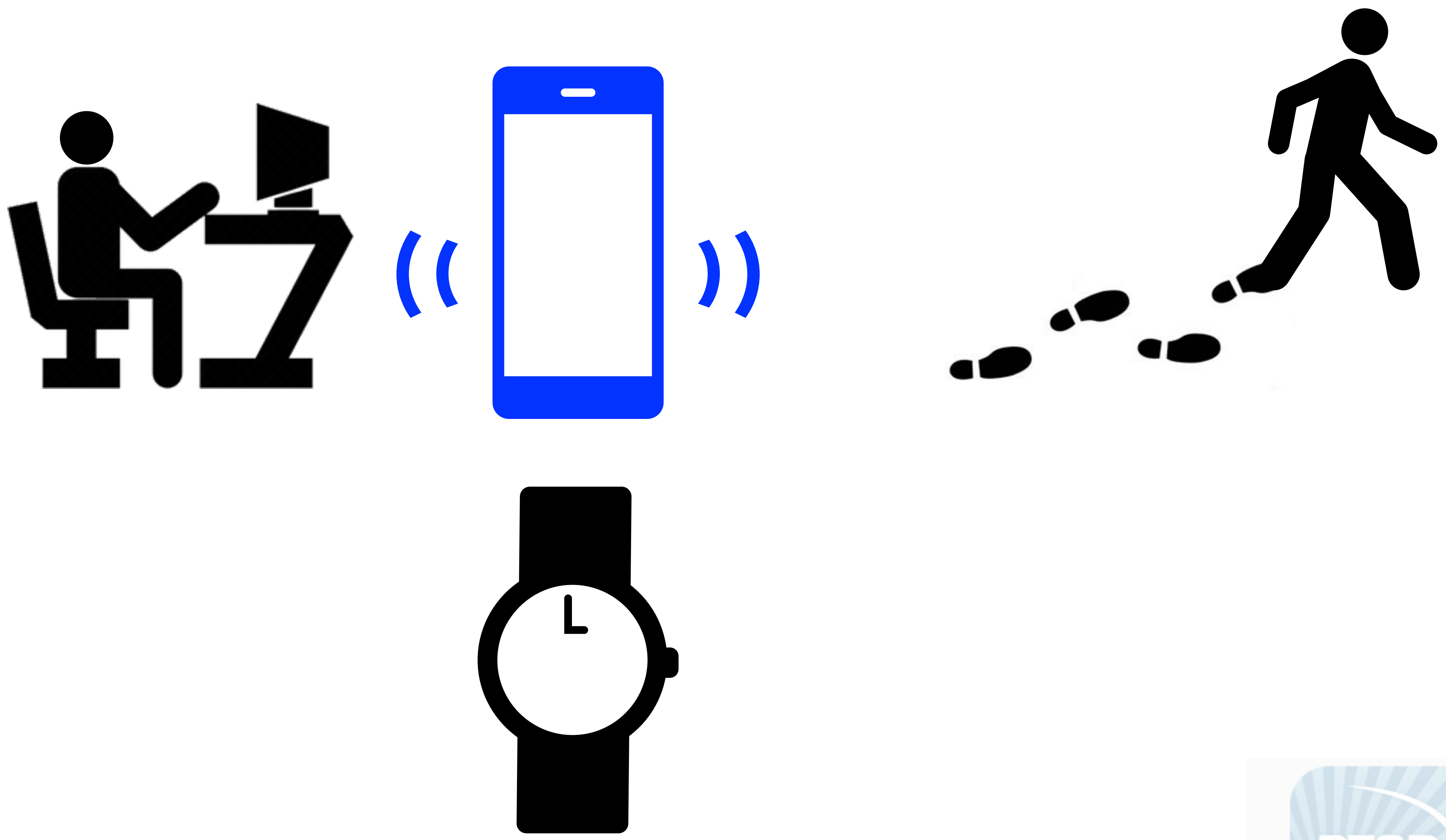

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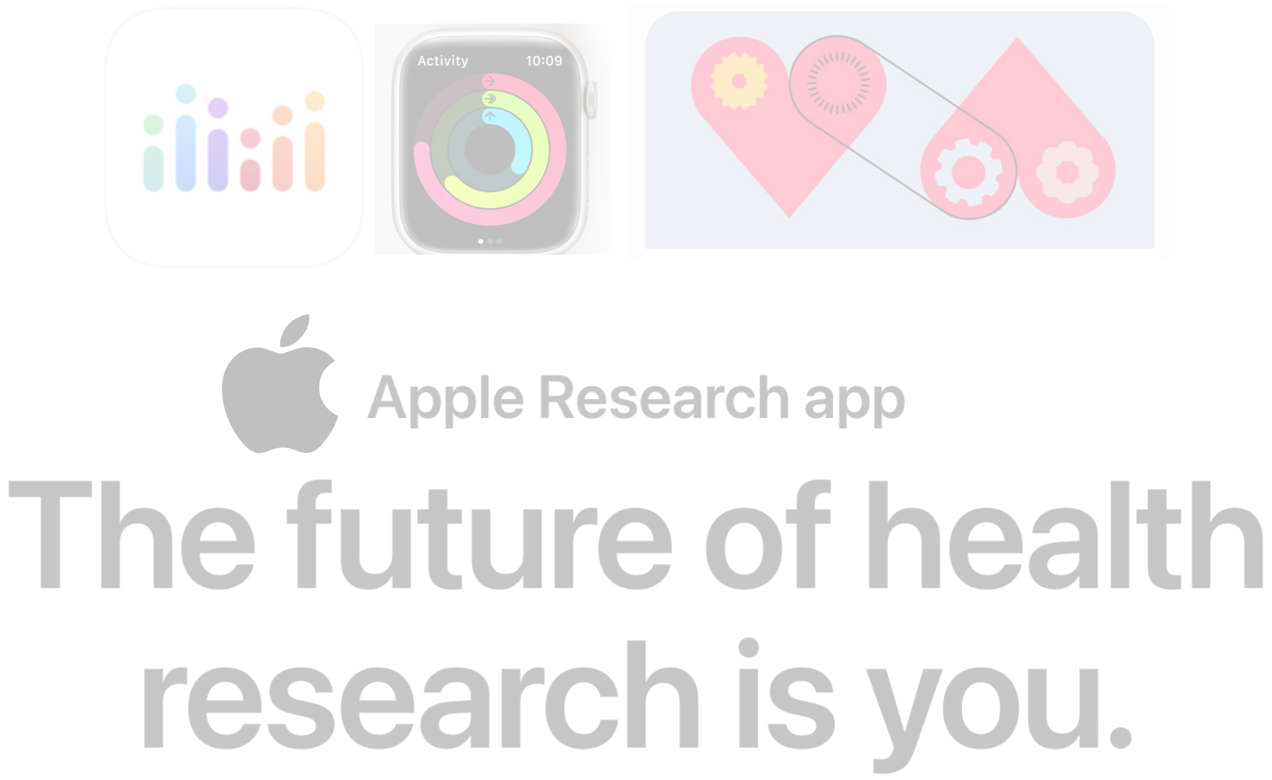

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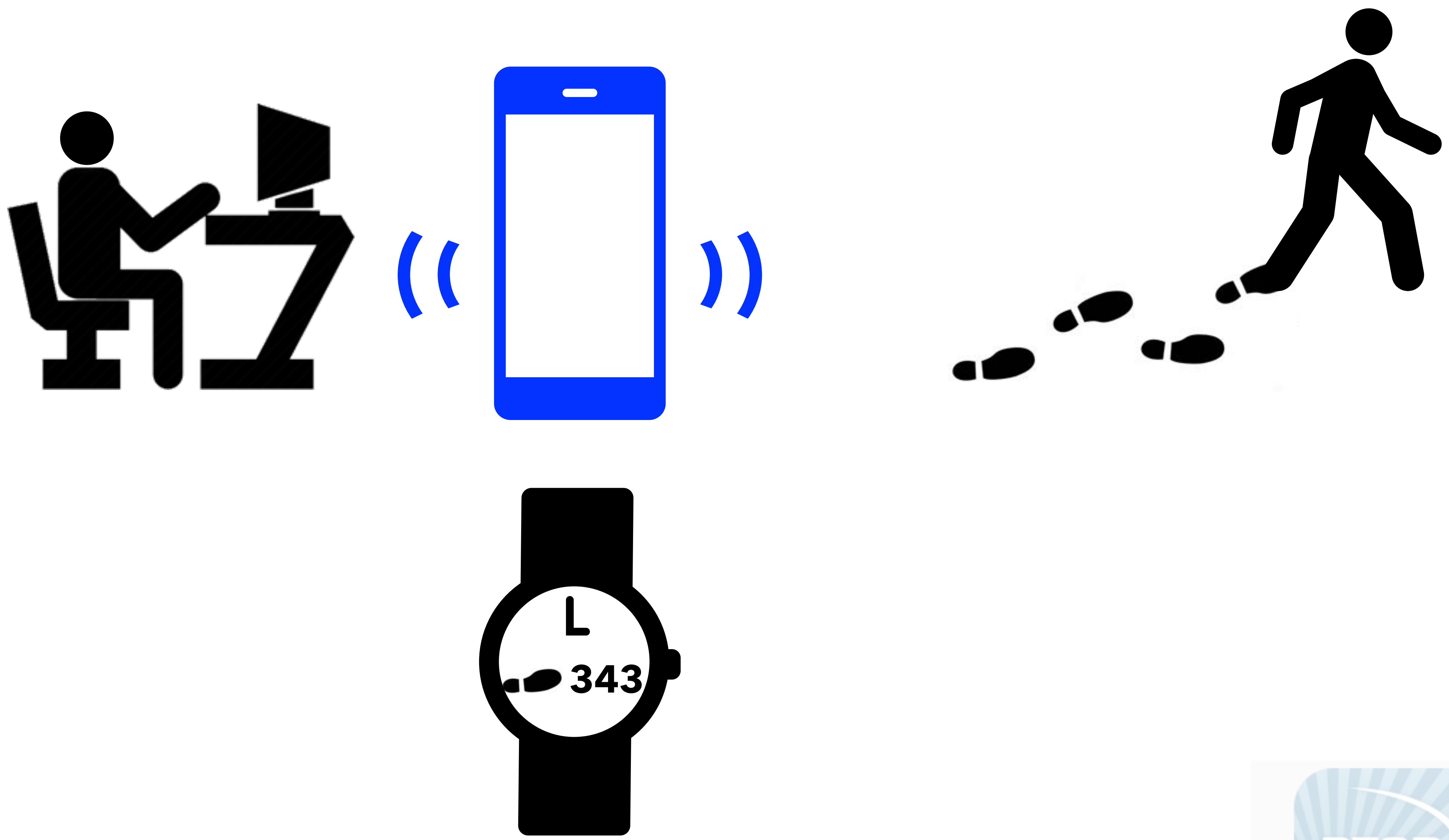

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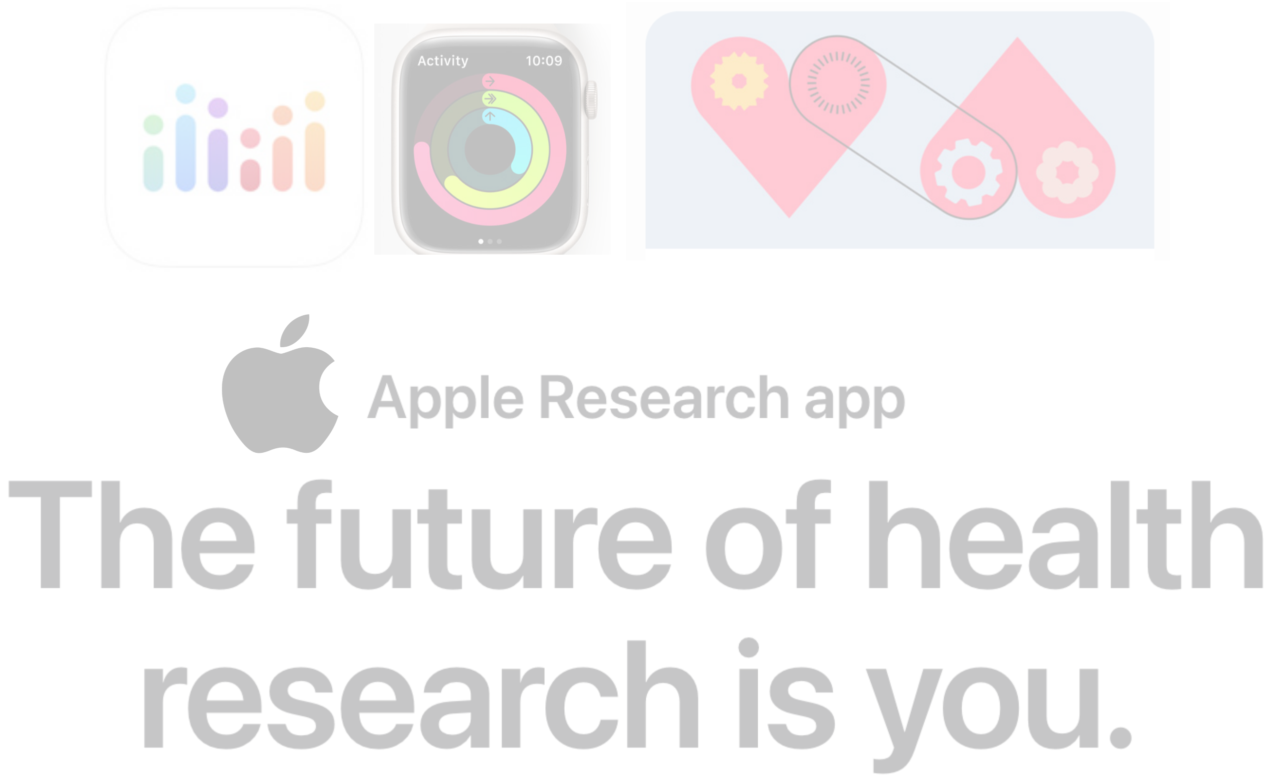
- **Outcome**: 30-min step count after decision time

image credits: va.gov apple.com



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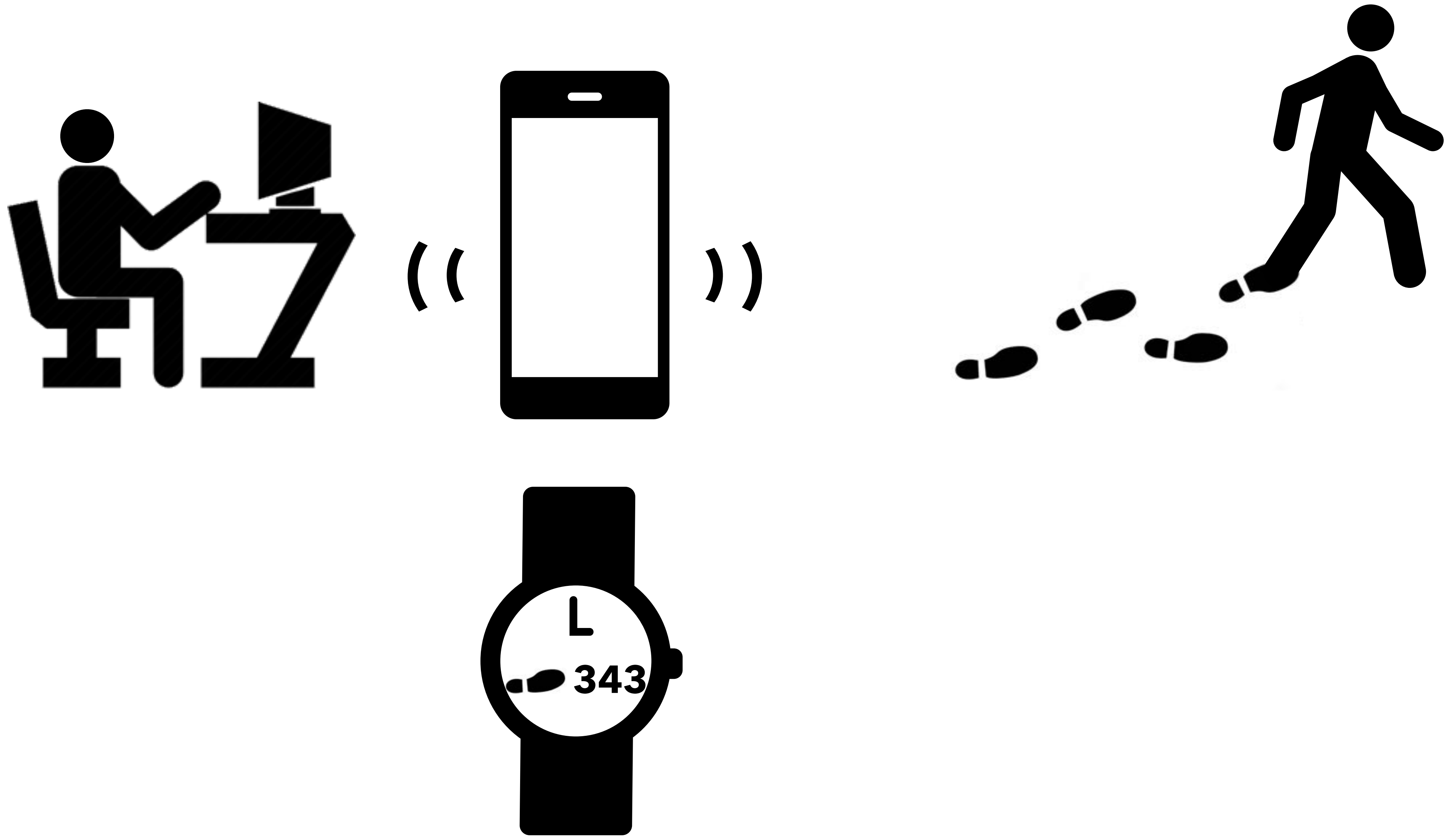

After-study personalized inference questions

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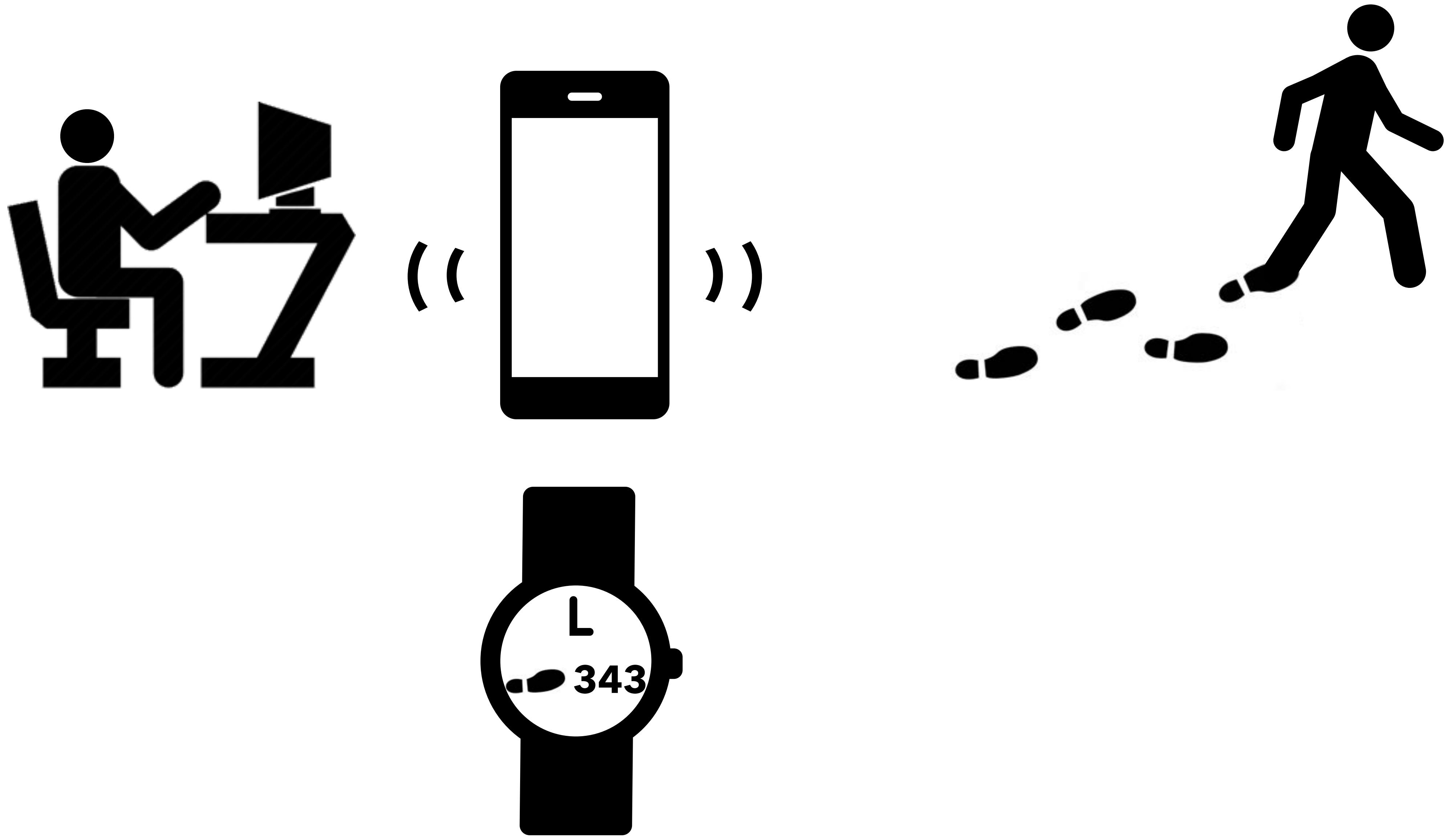

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After-study personalized inference questions

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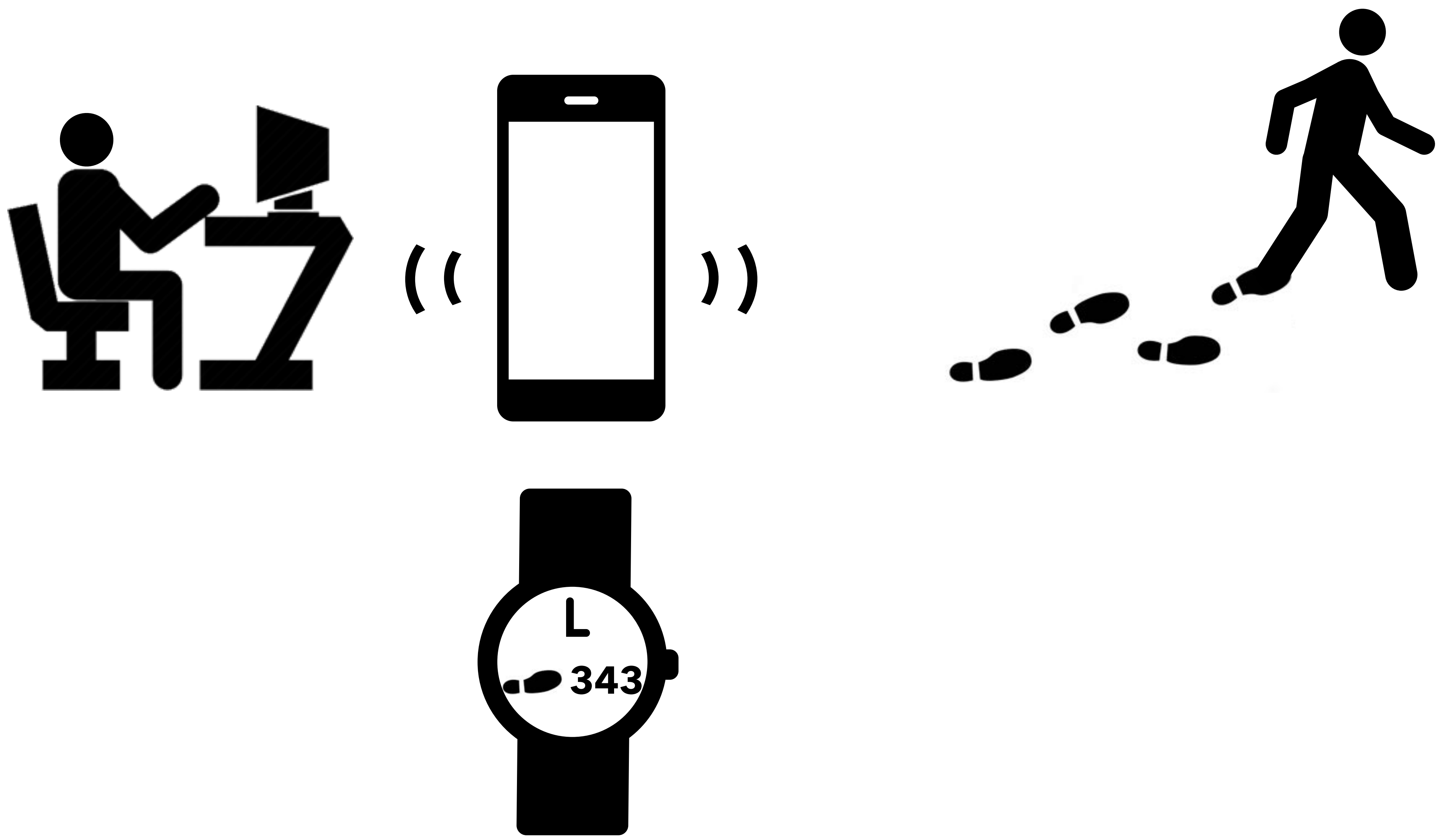

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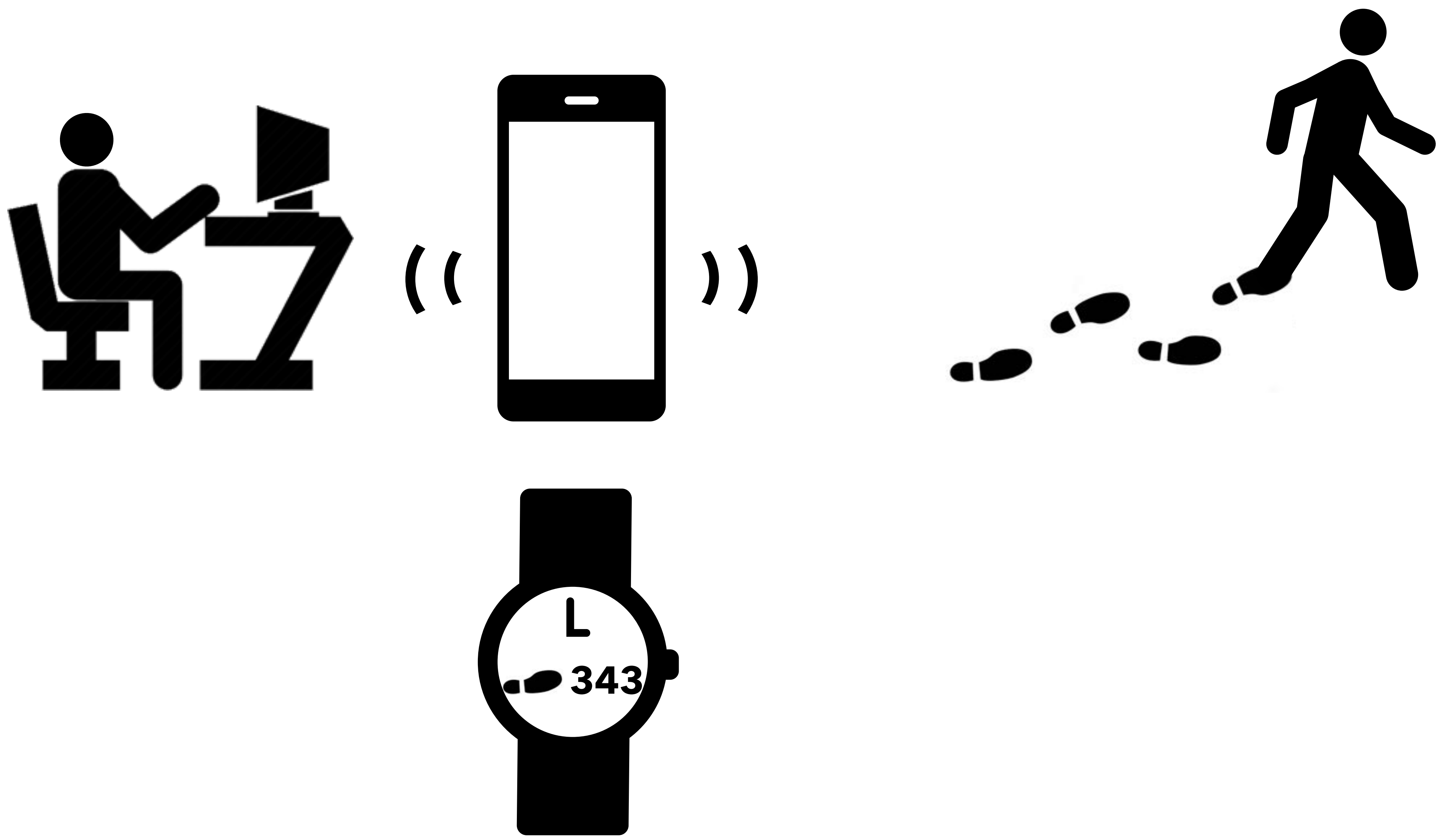

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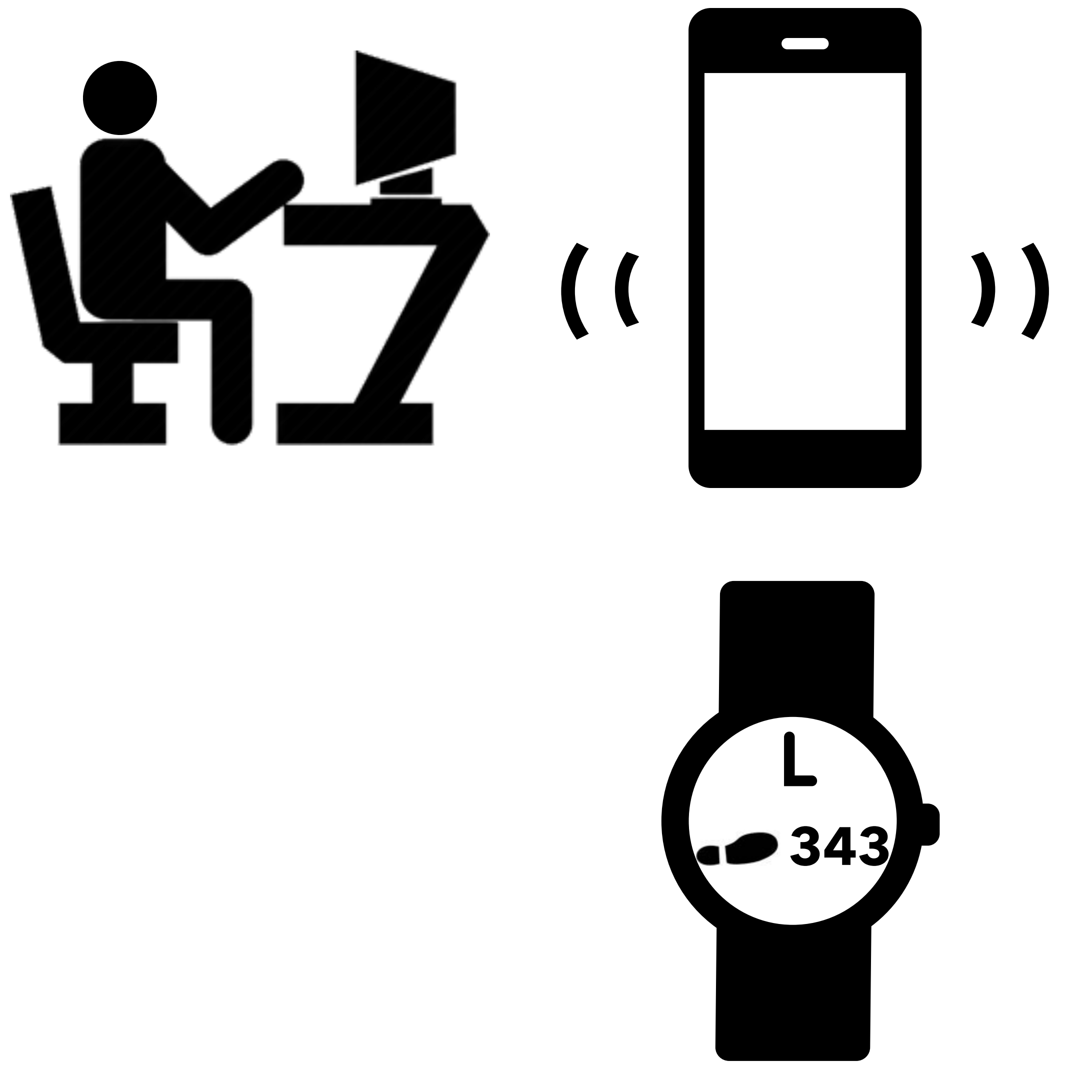

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Challenges: Lack of mechanistic models, adaptively collected data, expensive data collection

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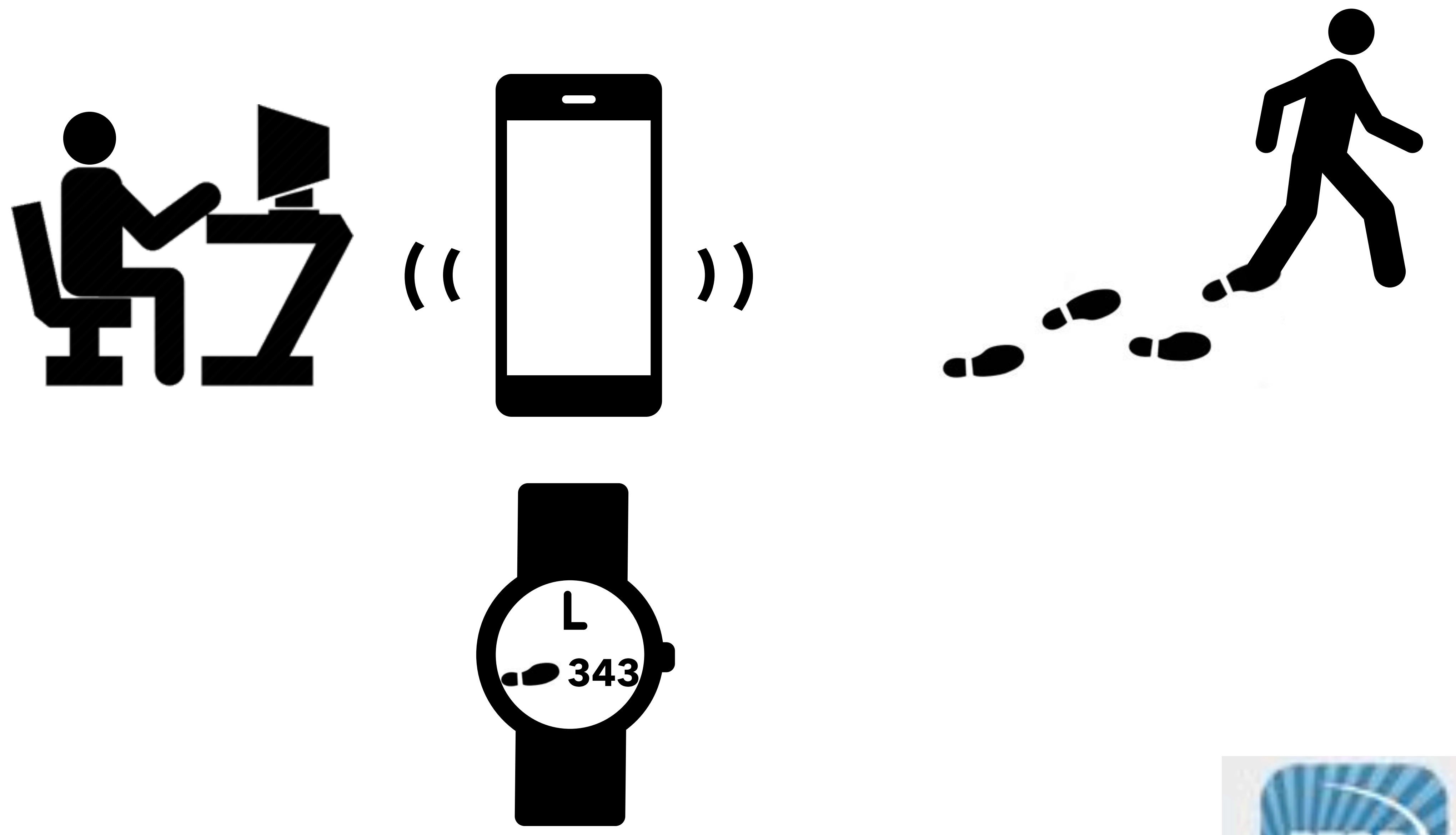

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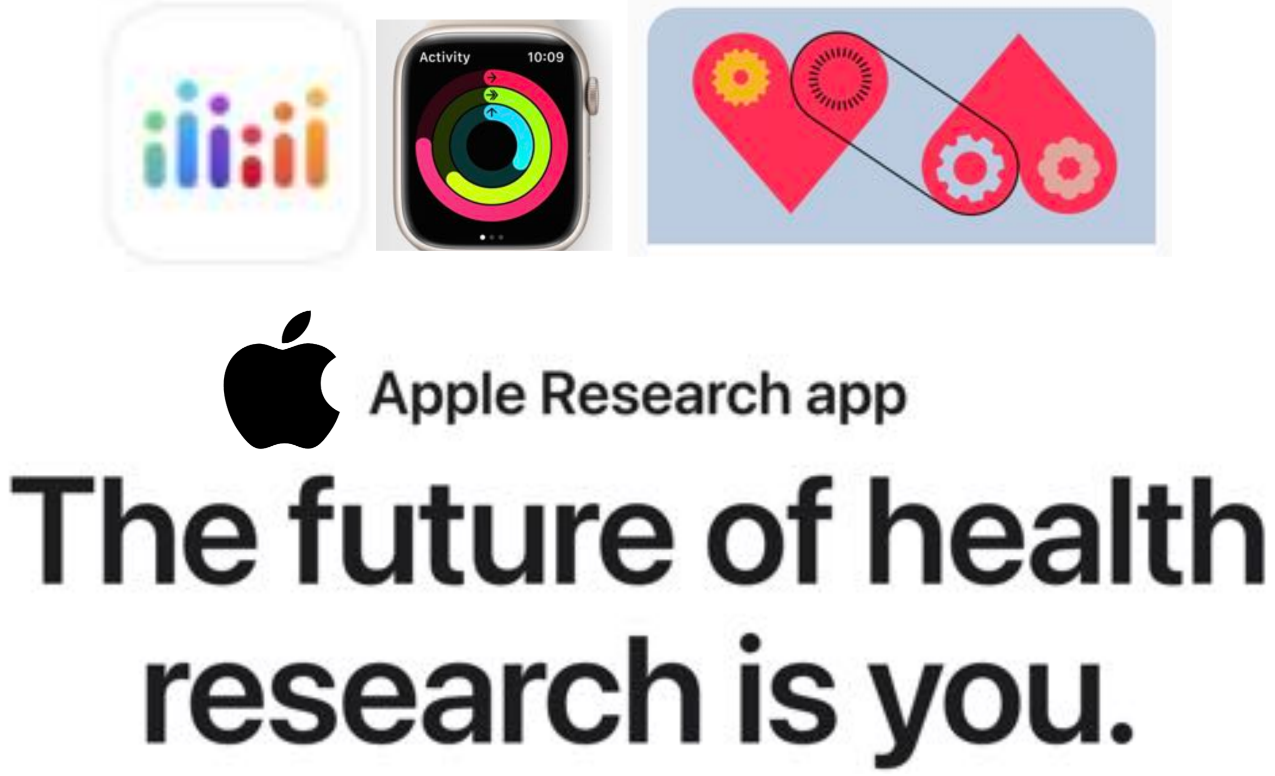
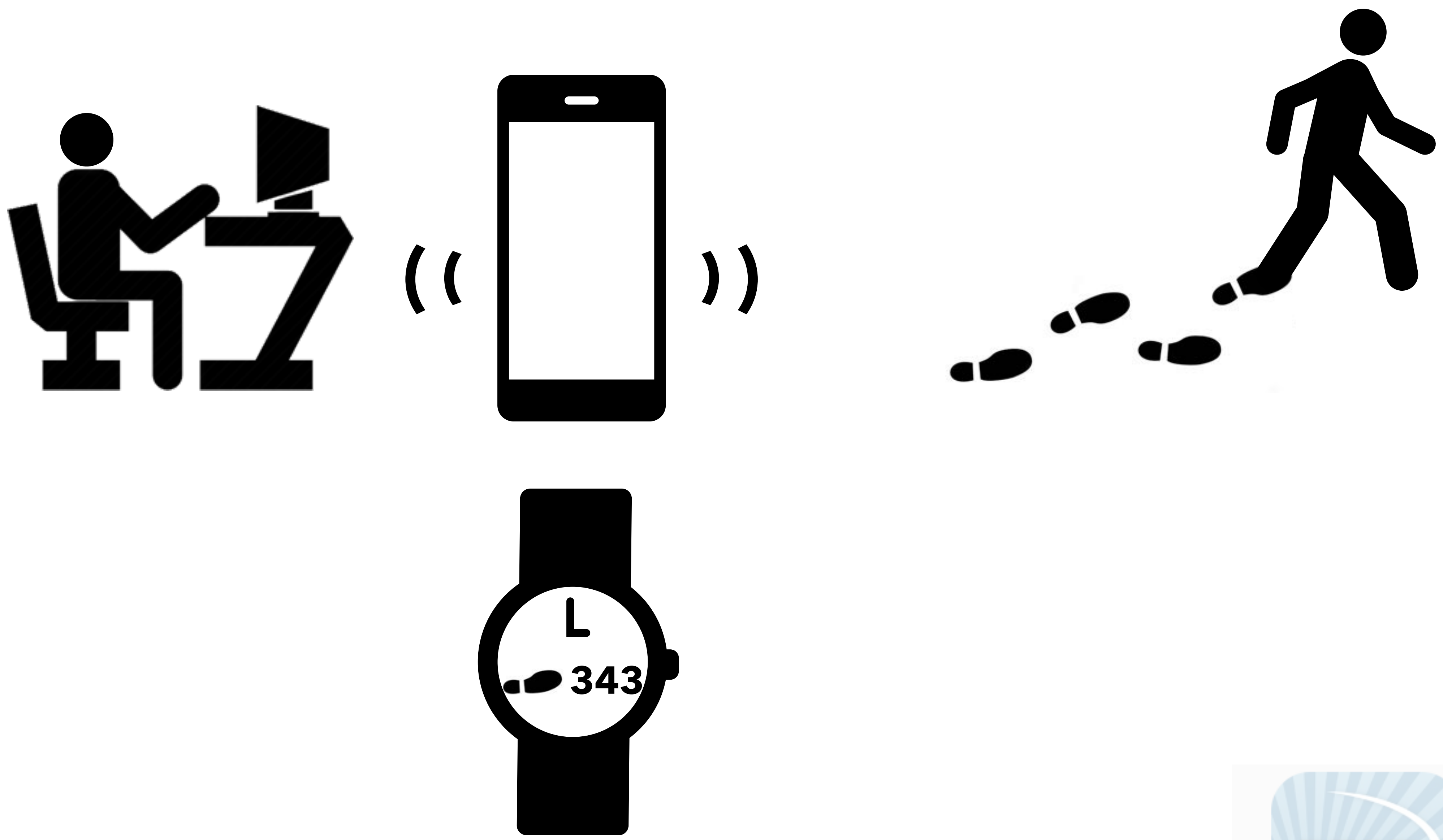


Part 1 overview: Sample-efficient personalized inference in sequential experiments

How to assign personalized digital treatments to help you?



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This talk

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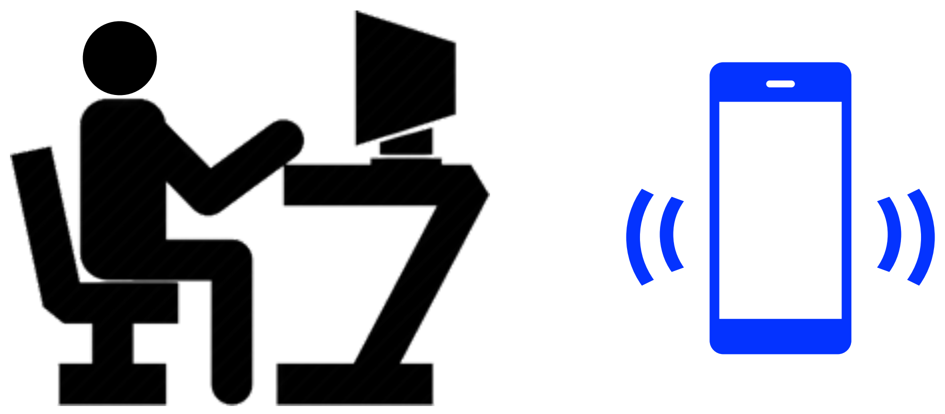

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For user $i \in [N]$ at time $t \in [T]$

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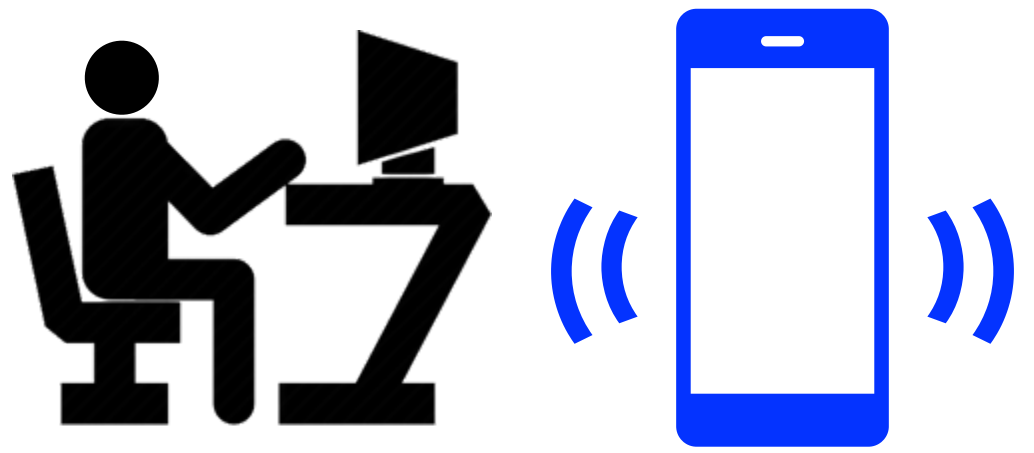

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e.g., ε-greedy, Thompson sampling, softmax, multiplicative weights, pooled variants,…

Sequentially adaptive policy that **can pool** observed data **across users** to speed up learning

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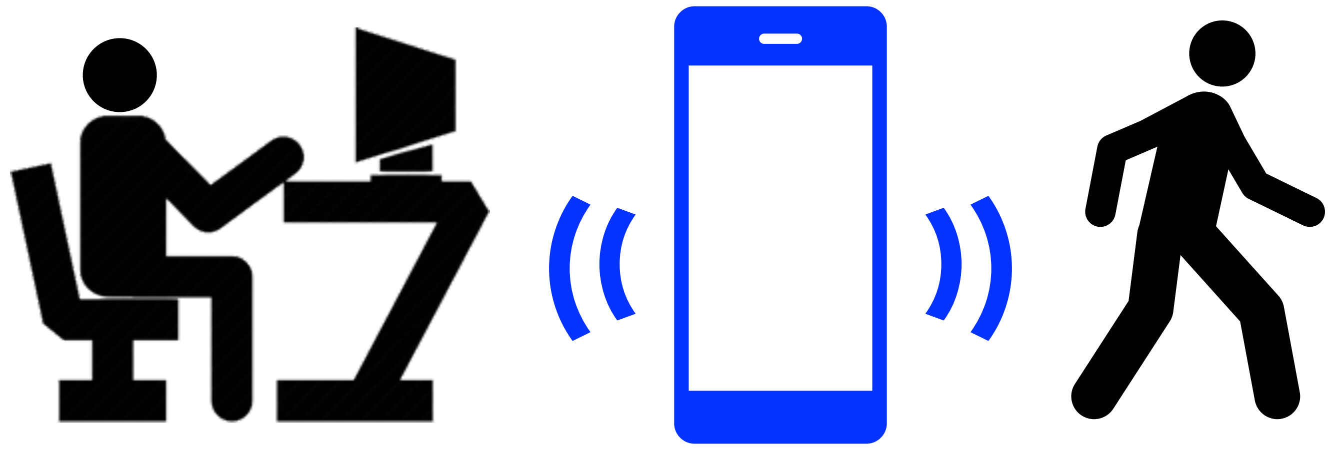

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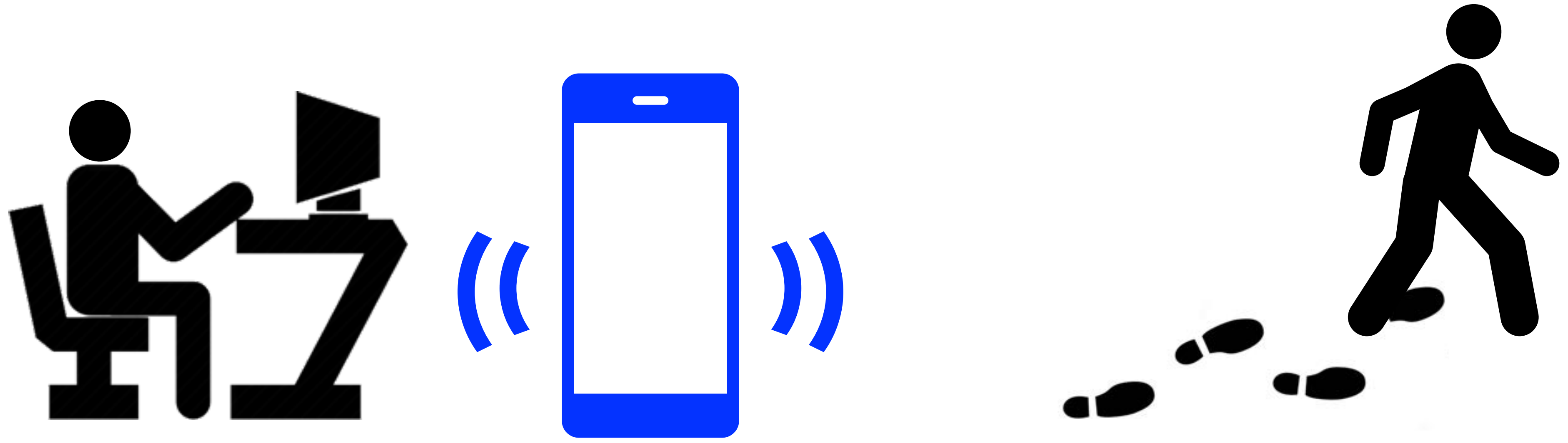

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[Neyman-Rubin framework + SUTVA]

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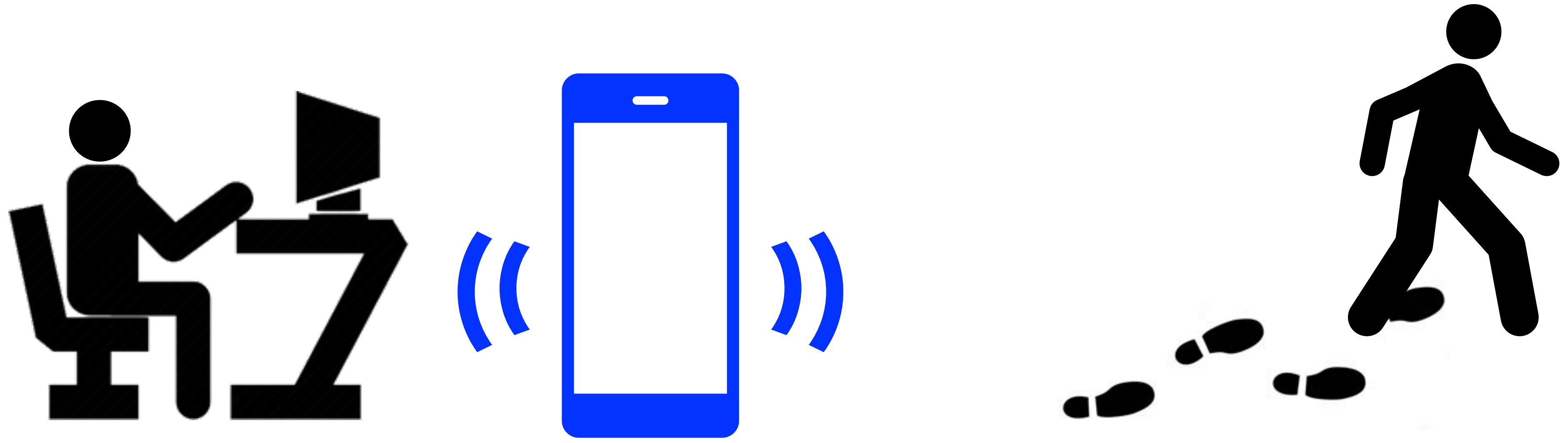

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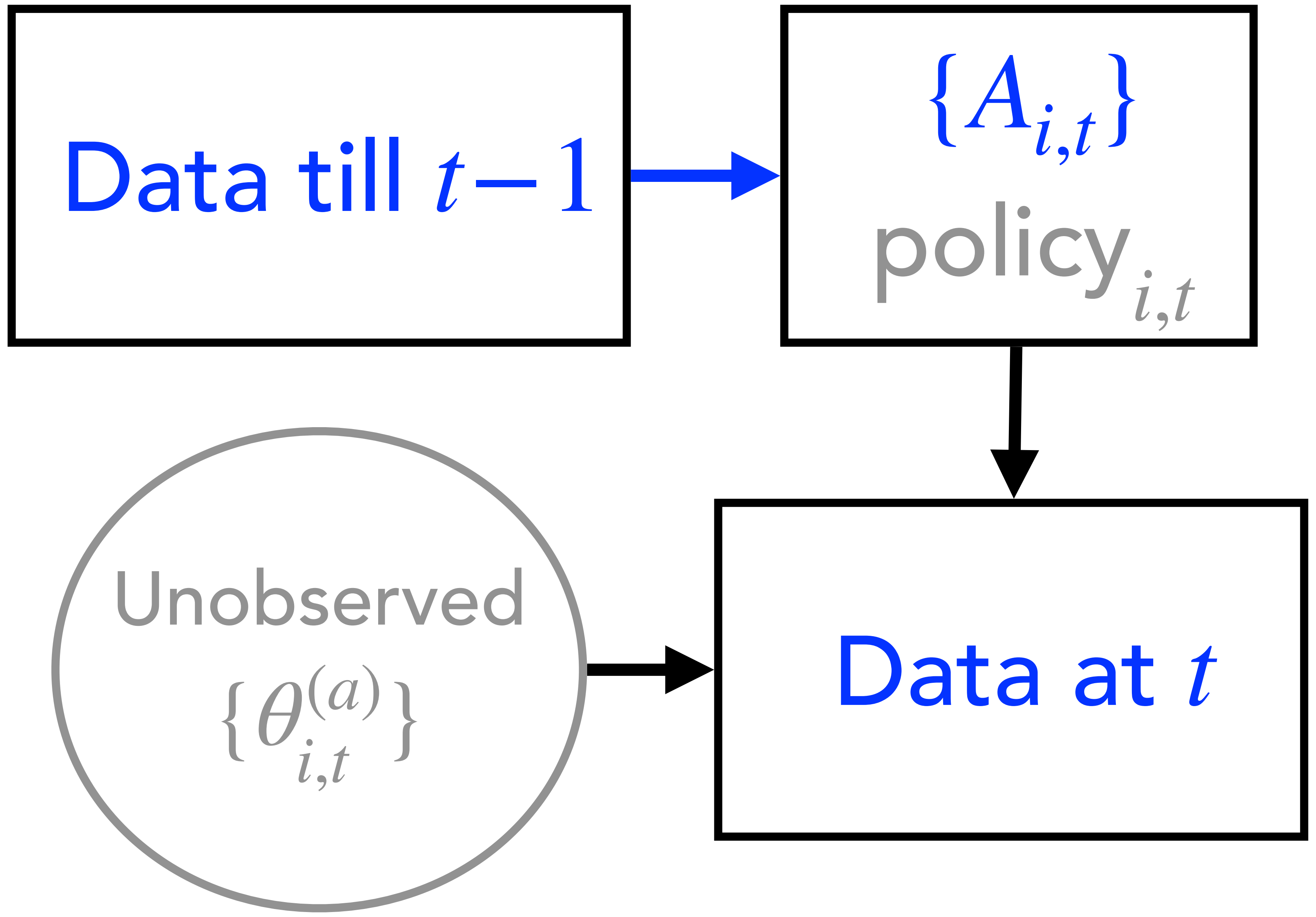

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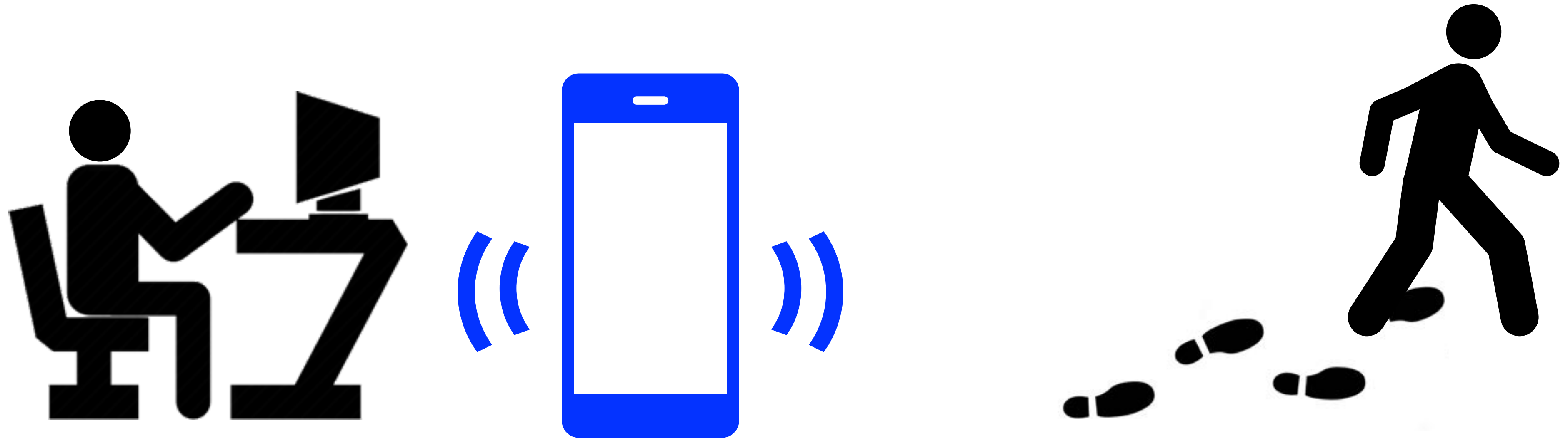

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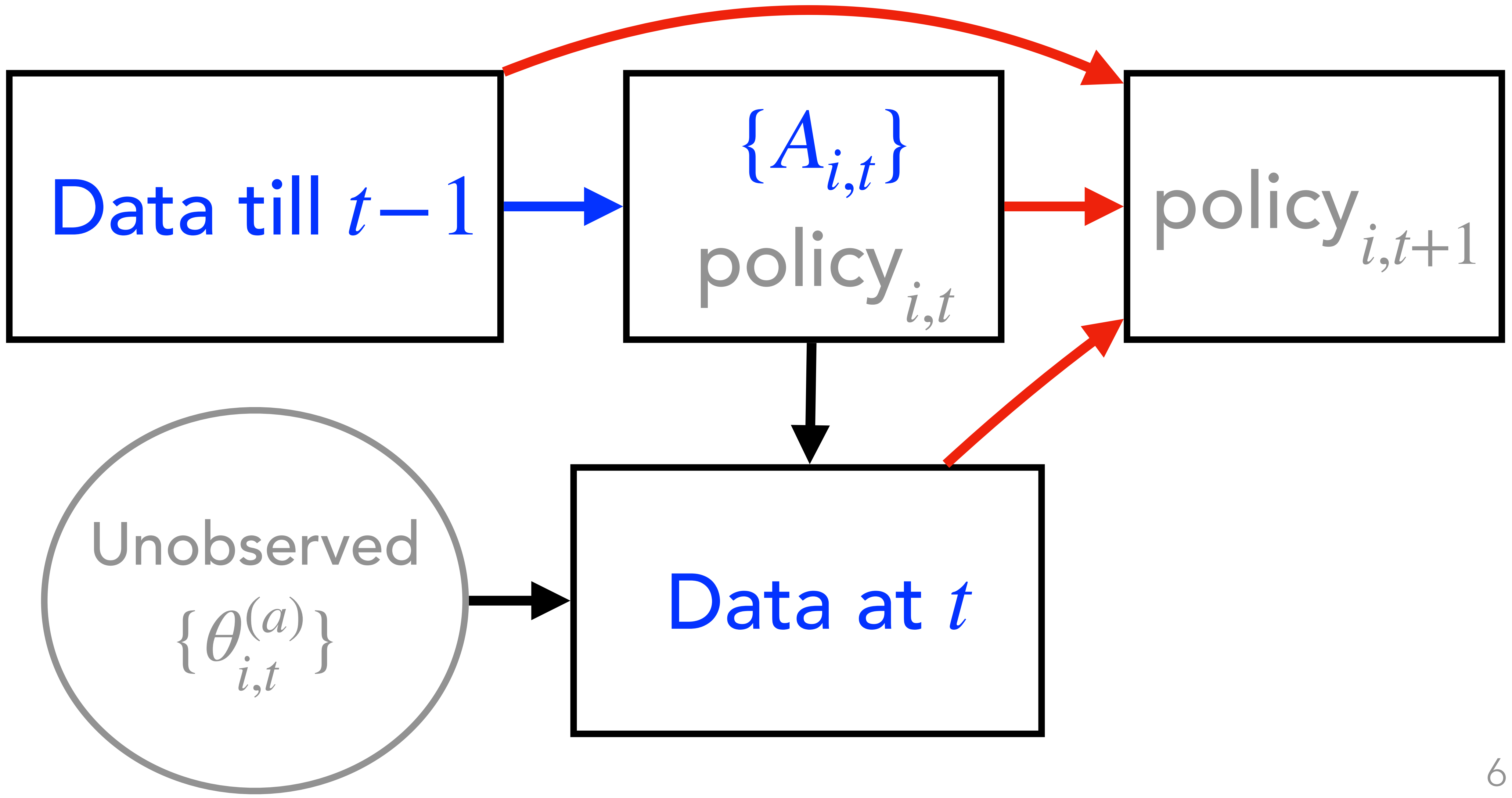

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Estimate counterfactual means $\{\theta_{i,t}^{(a)}\}$ for $a \in \{0,1\}$, **all** N users & T times

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- Enable generic after-study analyses and assist next study design
- E.g., how effective was the notification for user i at time t ($\theta_{i,t}^{(1)} - \theta_{i,t}^{(0)}$)?

Estimate counterfactual means $\{\theta_{i,t}^{(a)}\}$ for $a \in \{0,1\}$, all N users & T times

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Challenges:

➡ More unknowns than (noisy) observations

An impossible task without structural assumptions…

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Hope:

★ N iid users

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- ★ If users are not all too different & multiple observations can help find similarities

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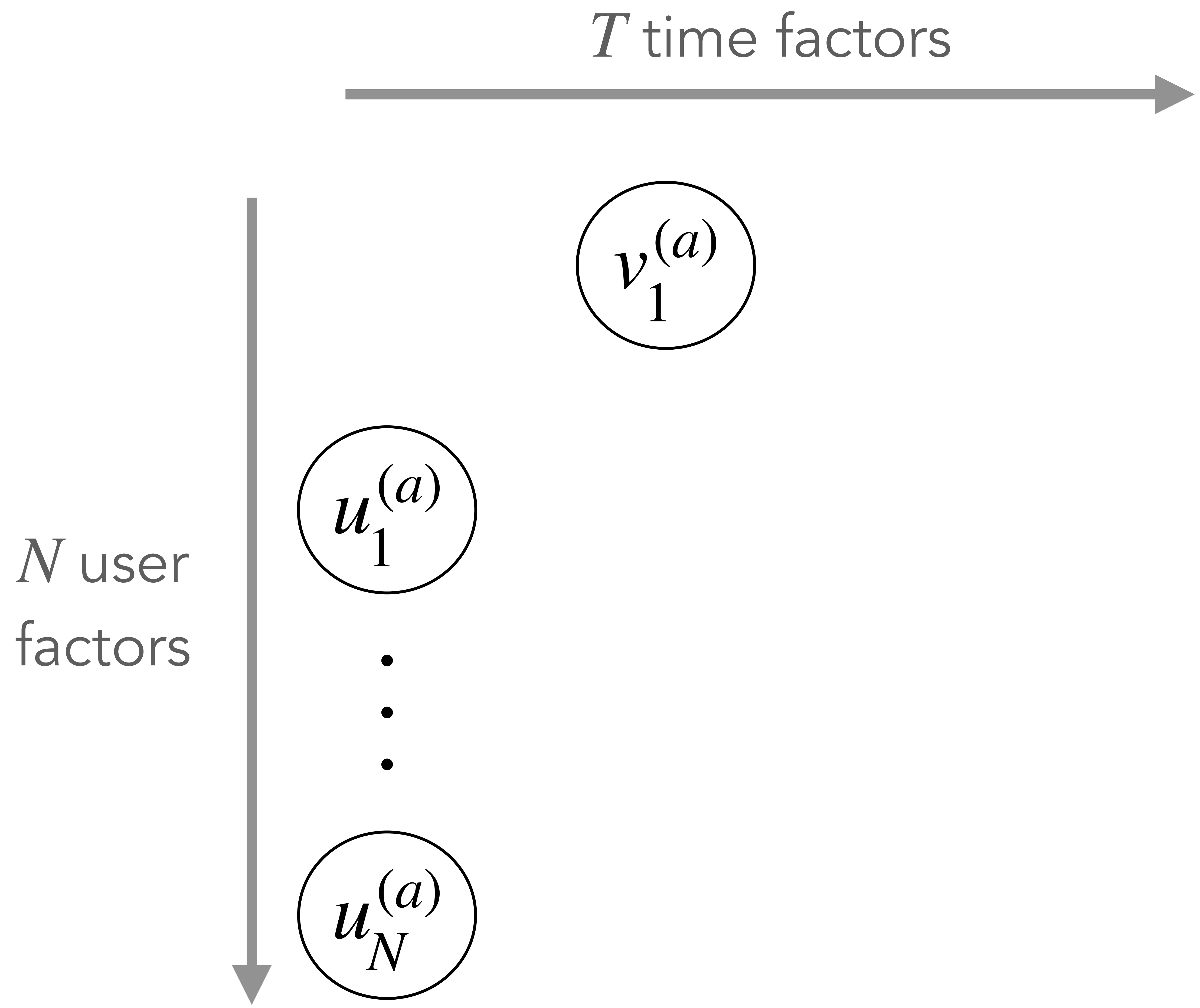

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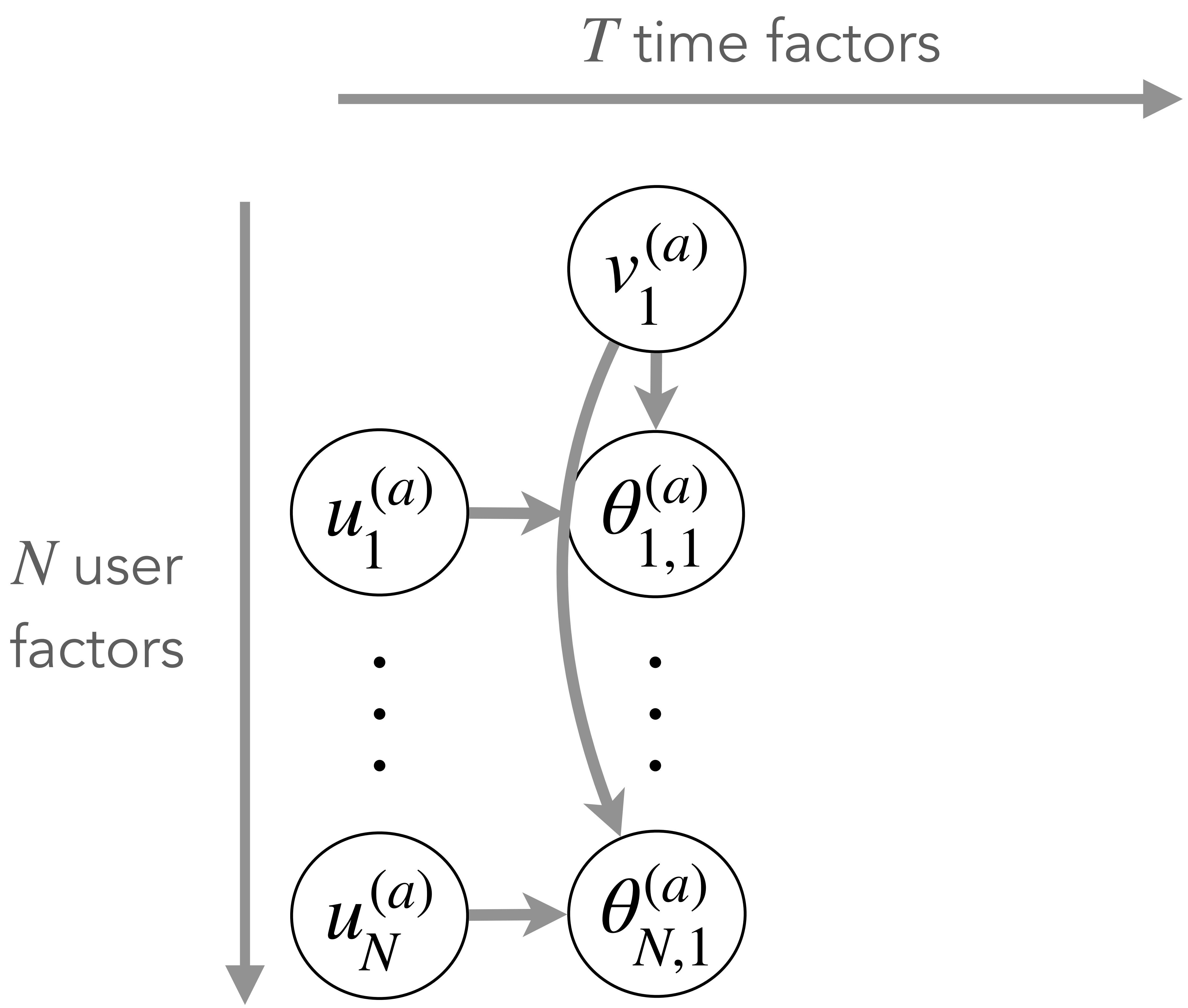

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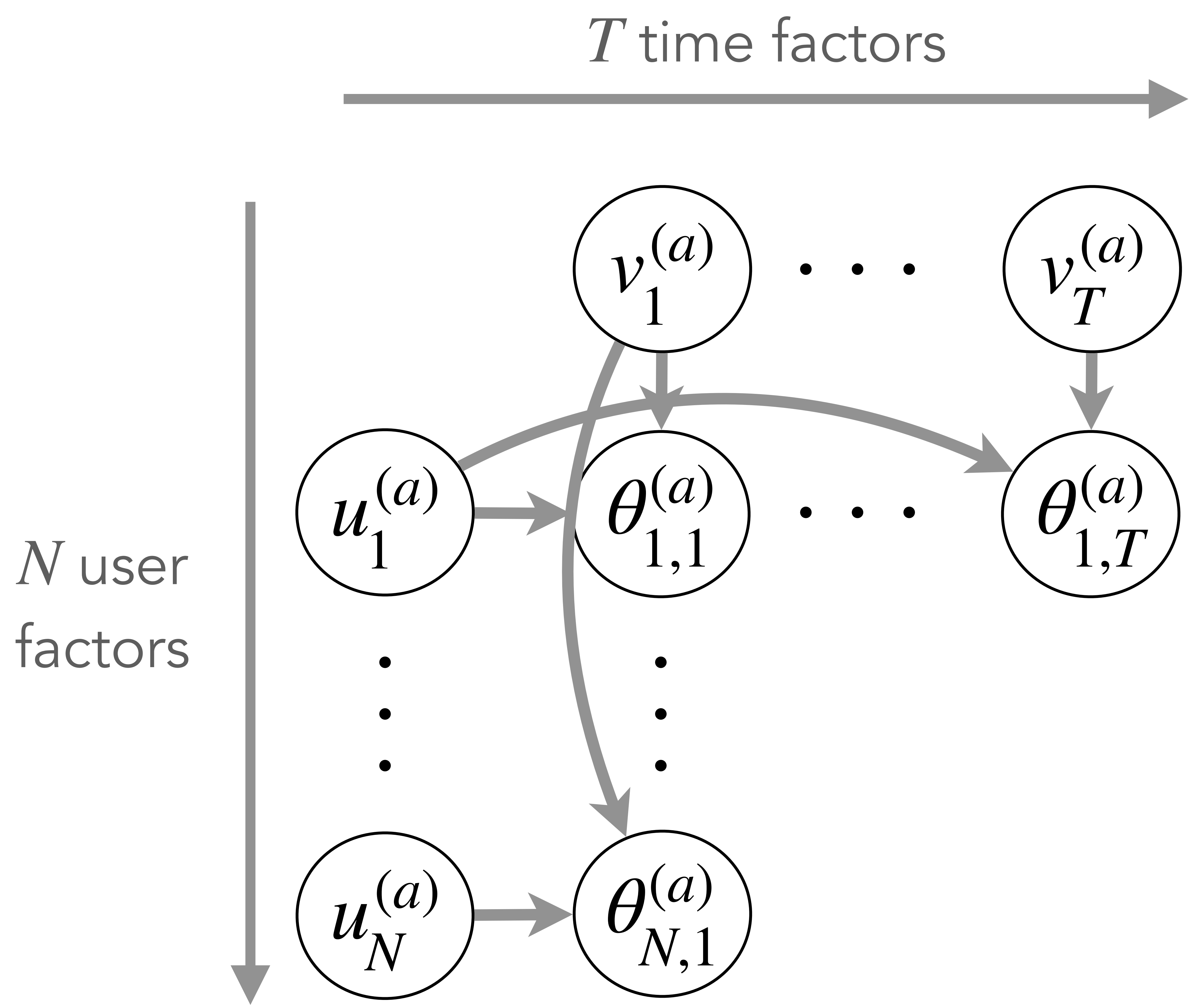

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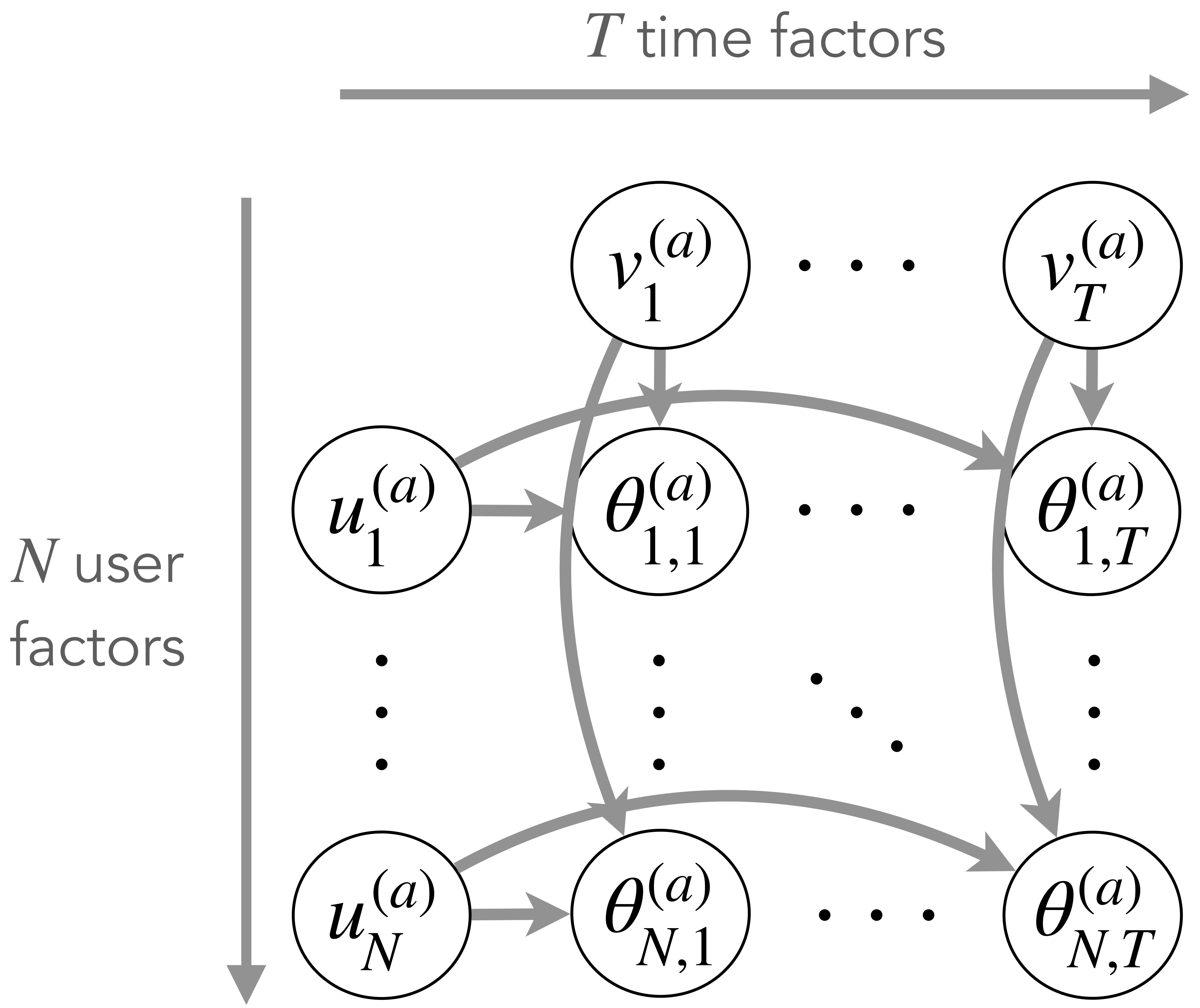

User nearest neighbors estimator for $\theta_{i,t}^{(a)}$

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$$\rho_{i,j}^{(a)} = \frac{\sum_{t'=1}^{T} (Y_{i,t'} - Y_{j,t'})^2 \cdot \mathbf{1}(A_{i,t'} = A_{j,t'} = a)}{\sum_{t'=1}^{T} \mathbf{1}(A_{i,t'} = A_{j,t'} = a)}$$

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$$\mathbf{1}(\rho_{i,j}^{(a)} \leq \eta, A_{j,t} = a)$$

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2. Average outcome across user neighbors treated with a at time t

$$\widehat{\theta}_{i,t,\text{user-NN}}^{(a)} = \frac{\sum_{j=1}^{N} Y_{j,t} \cdot \mathbf{1}(\rho_{i,j}^{(a)} \le \eta, A_{j,t} = a)}{\sum_{j=1}^{N} \mathbf{1}(\rho_{i,j}^{(a)} \le \eta, A_{j,t} = a)}$$

Main result: A non-asymptotic guarantee for each (i, t, a)

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User factor distribution

$\downarrow$

(Uniform on finite set of size M)

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User-NN guarantees: Advantages

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Challenges tackled: First guarantee for user-time-level counterfactuals

✓ More unknowns than observations

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$$\Downarrow$$

$$|\,??\, - \theta^{(a)}_{i,t}\,| = \tilde{O}\left(\frac{1}{\sqrt{T}} + \frac{1}{\sqrt{N}}\right)$$

Can we improve the <u>slow rate</u> in T?

Challenges tackled: First guarantee for user-time-level counterfactuals

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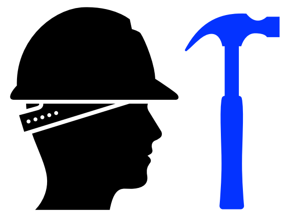

Martingale concentration, **new sandwich argument** for nn

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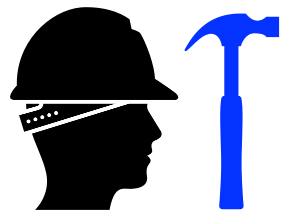

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$$\approx \max\{|\hat{u}_i - u_i|, |v_t - \hat{v}_t|\}$$

- **Convert + to ×:** $|u_i v_t - ??| = O(|u_i - \hat{u}_i| \times |v_t - \hat{v}_t|)$

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$$Y_{j,t} + Y_{i,t'} - Y_{j,t'}$$

$$\rho_{i,j}^{(a)} \leq \eta, \quad \rho_{t,t'}^{(a)} \leq \eta'$$

This is our improved nearest neighbors estimator!

$$\cancel{u_i v_t} - ?? = (u_i - \hat{u}_i) \times (v_t - \hat{v}_t)$$

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$$\widehat{\theta}^{(a)}_{i,t,\mathrm{DR\text{–}NN}} = \frac{\sum_{j,t'} (Y_{j,t} + Y_{i,t'} - Y_{j,t'}) \mathbf{1}_{i,t,j,t'}}{\sum_{j,t'} \mathbf{1}_{i,t,j,t'}}$$

$$\mathbf{1}_{i,t,j,t'} = \mathbf{1}(\rho^{(a)}_{i,j} \leq \eta,\ \ \rho^{(a)}_{t,t'} \leq \eta', A_{j,t} = A_{i,t'} = A_{j,t'} = a)$$

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DR-NN error $\approx$ **user-NN error** $\times$ **time-NN error**

$\lesssim$ **min{user-NN error, time-NN error}**

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Doubly robust to heterogeneity in user factors & time factors

Double robustness, double machine learning…

[… Cassel+ '77, Robinson '88, Särndal+ '89, Robins+ '94, '95, '08, '09, Newey+ '94, '18, Bickel+ '98, van der Laan+ '03, Lunceford+ '04, Davidian+ '05, Li+ '11, Jiang+ '15, Chernozhukov+ '18, Hirshberg+ '18, Diaz '19, Arkhangelsky+ '21, Dorn+ '21 …]

Simulation results

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Uniform latent factors on $[-0.5,0.5]^4$, Gaussian noise, pooled ε-greedy policy ($\varepsilon = 0.5$)

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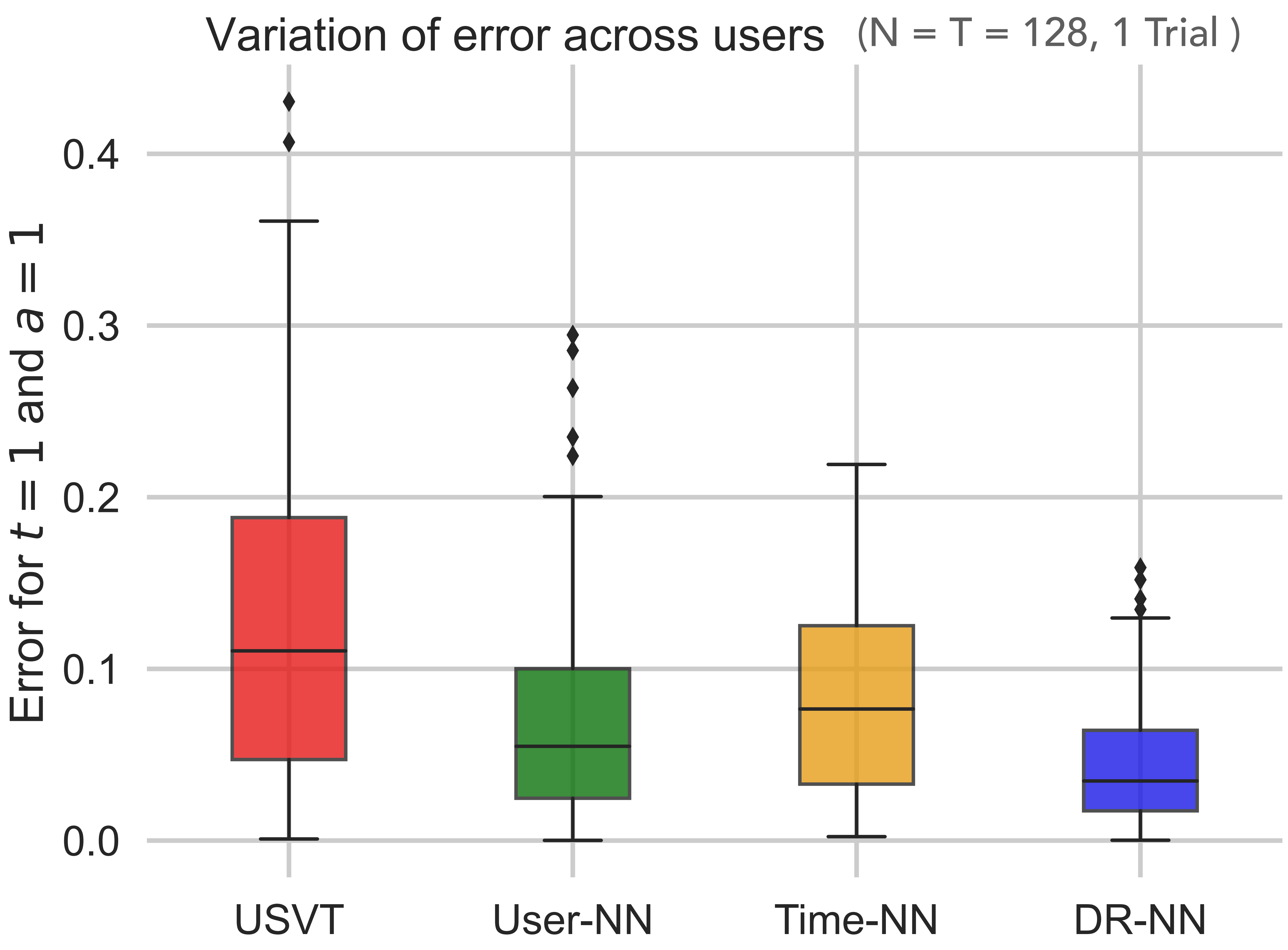

A baseline algorithm from [Chatterjee 2014]

Simulation results

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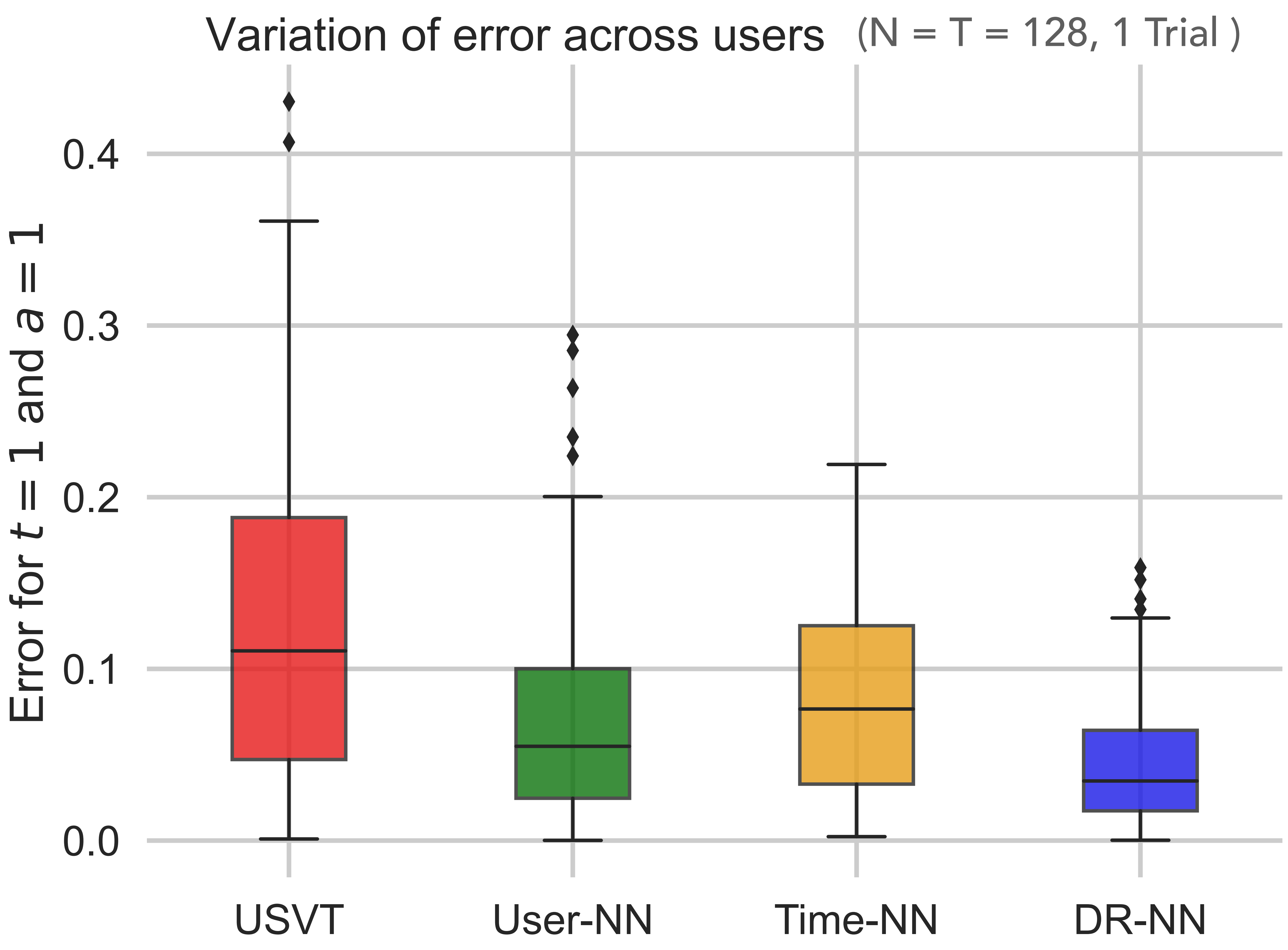

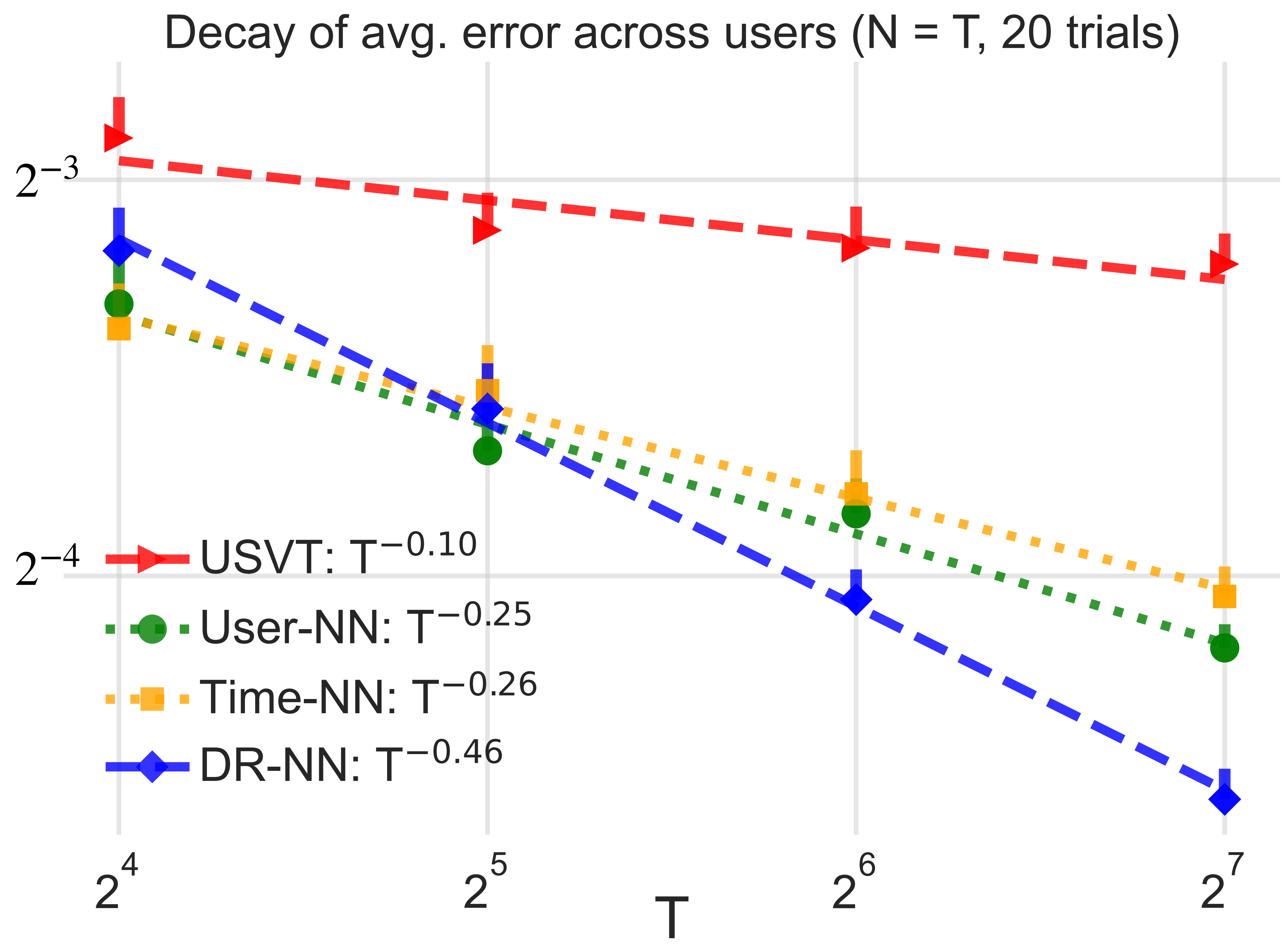

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Personalized HeartSteps results

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Treatments assigned with Thompson sampling independently for 91 users for 90 days, 5 times a day

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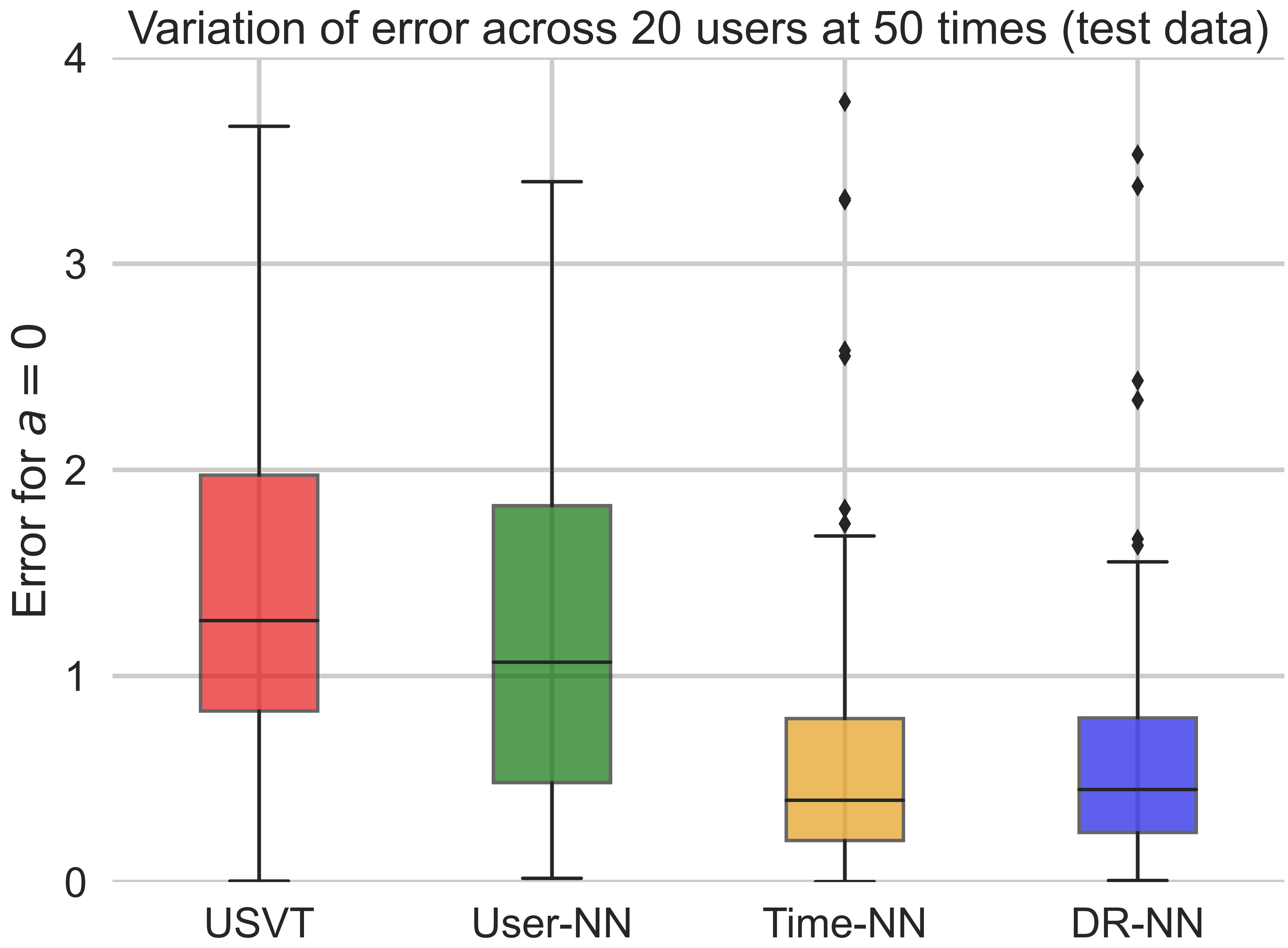

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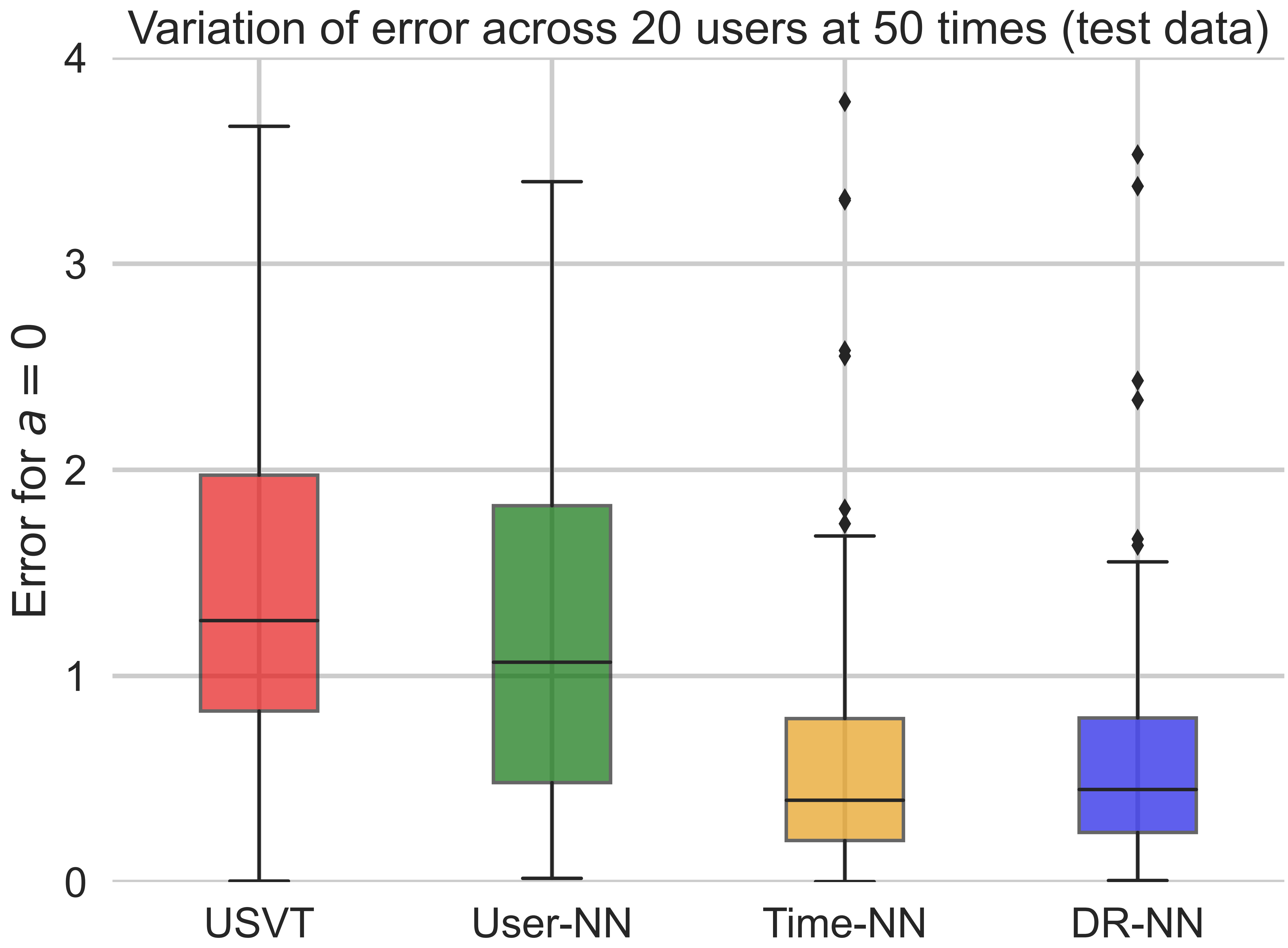

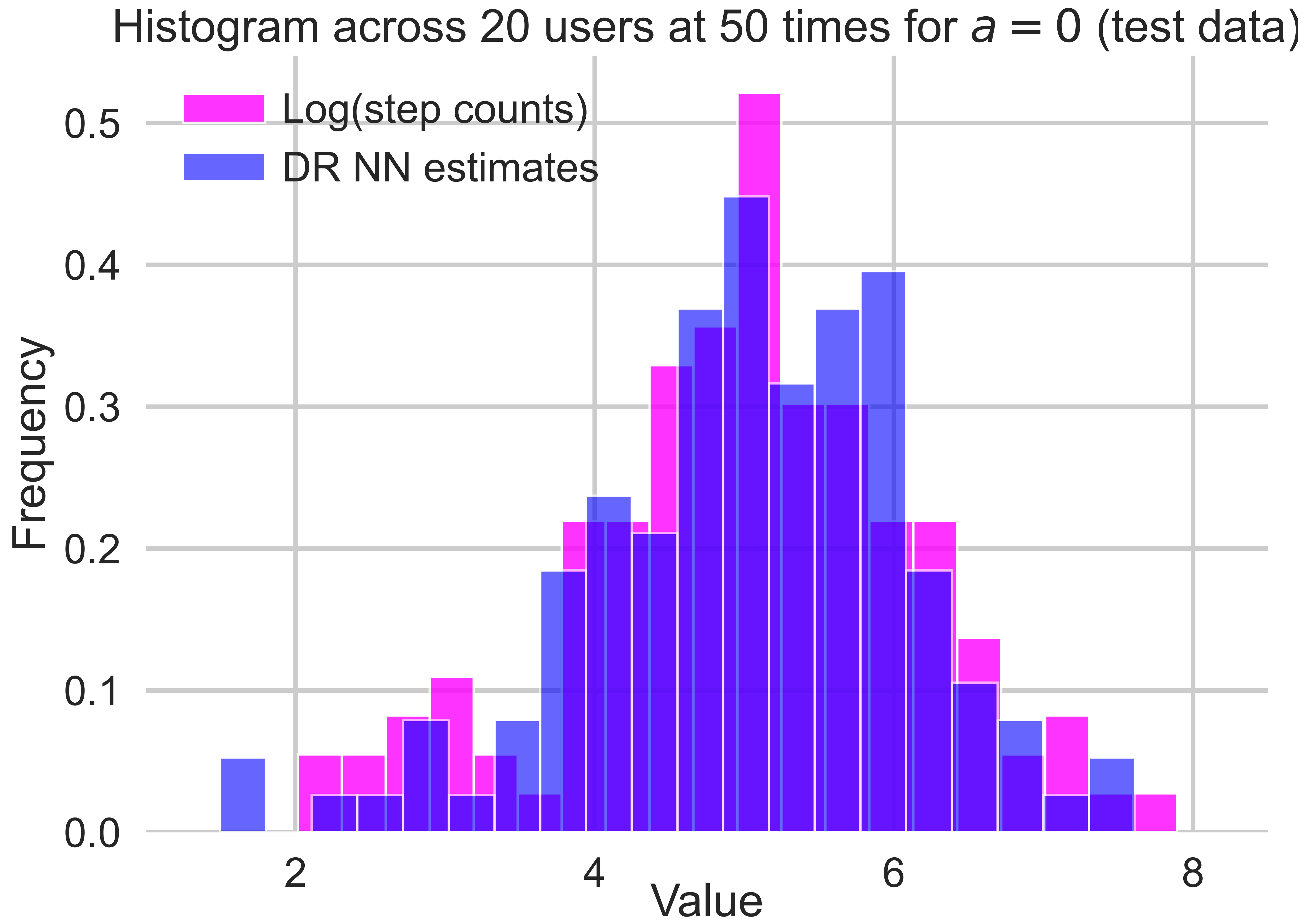

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Part 1 summary:
Sample-efficient inference with non-parametric factor models

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Sample-efficient inference with non-parametric factor models

✓ Inference in sequential experiments: User-NN with $\tilde{O}(T^{-1/4})$ error

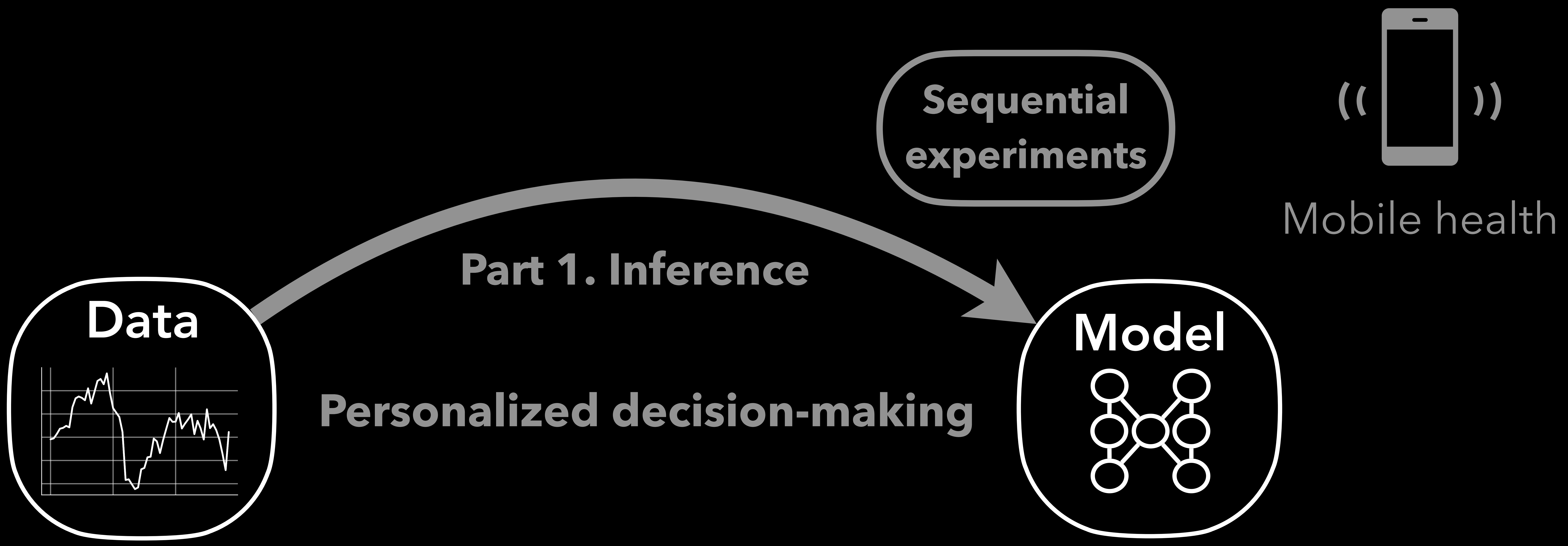
✓ Efficient estimators: Doubly robust-NN with $\tilde{O}(T^{-1/2})$ error

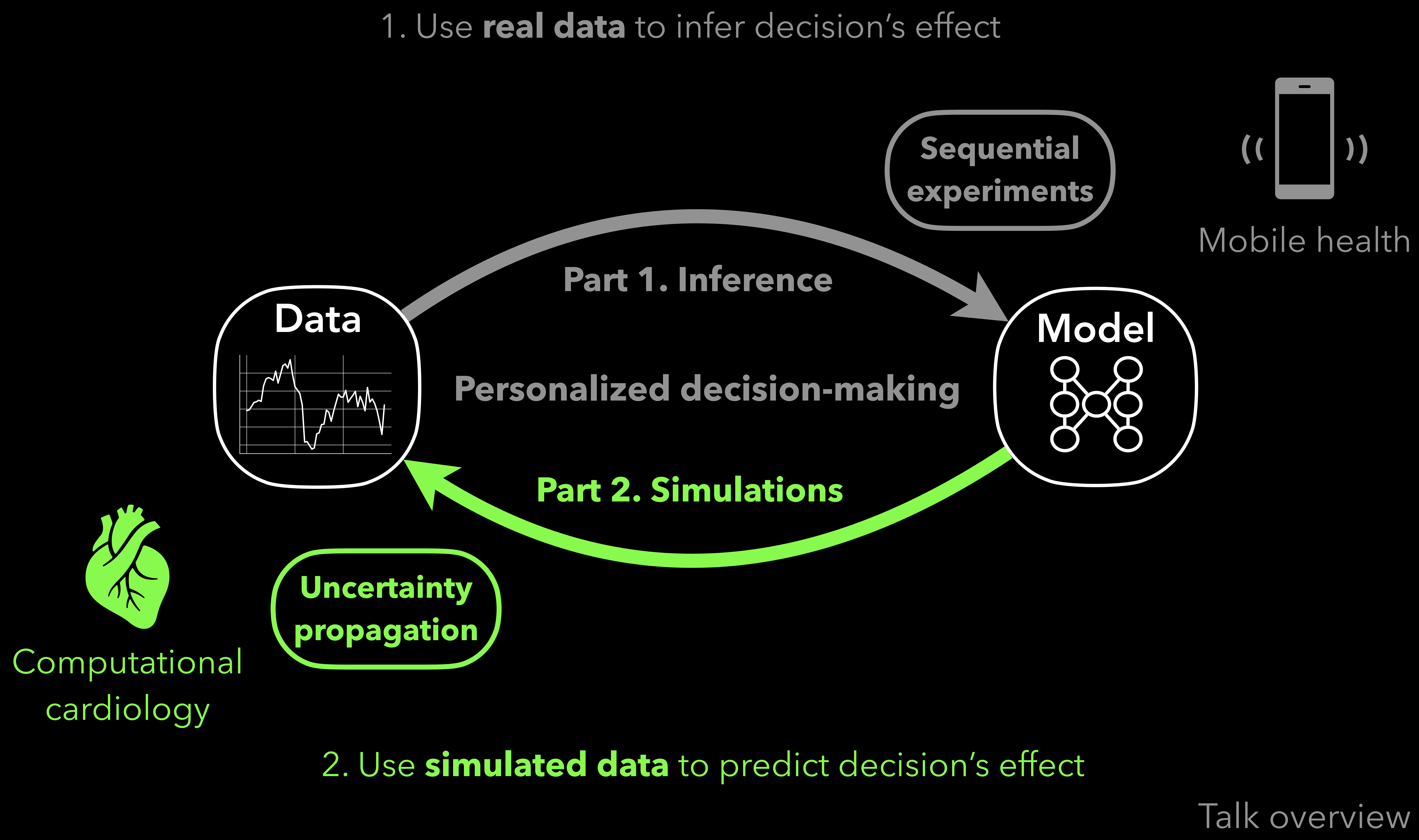
DR-NN error $\approx$ **user-NN error** $\times$ **time-NN error**

$\lesssim$ **min{user-NN error, time-NN error}**

✦ Future: Settings with contexts and covariates

1. Use **real data** to infer decision's effect



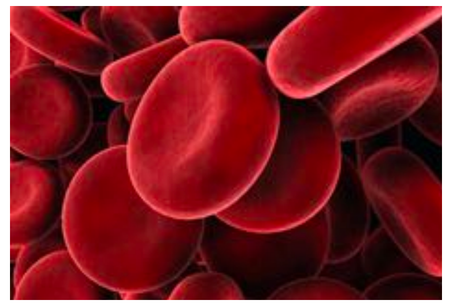

1. Use **real data** to infer decision's effect
Sequential experiments
Mobile health
Part 1. Inference
Data
Model
Personalized decision-making
Part 2. Simulations
Uncertainty propagation
Computational cardiology
2. Use **simulated data** to predict decision's effect
Talk overview

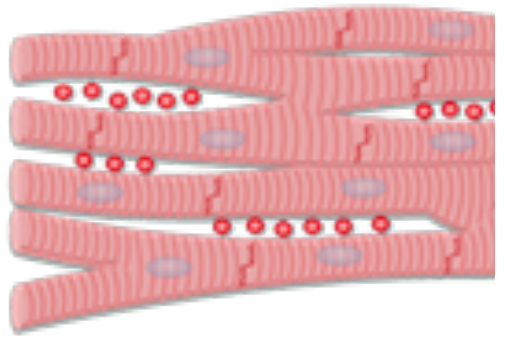
Complex multi-scale simulation systems

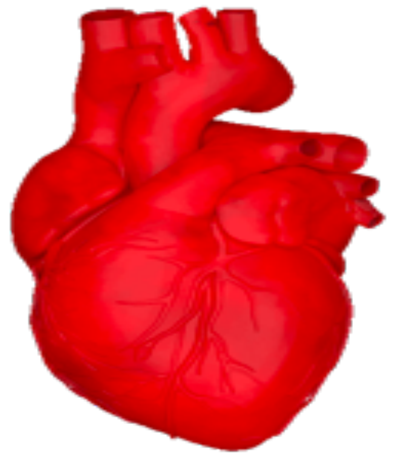
Complex multi-scale simulation systems

Basic unit | Sub-component | System

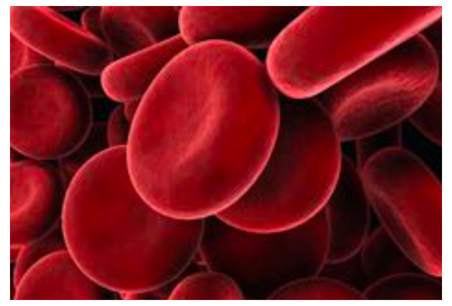
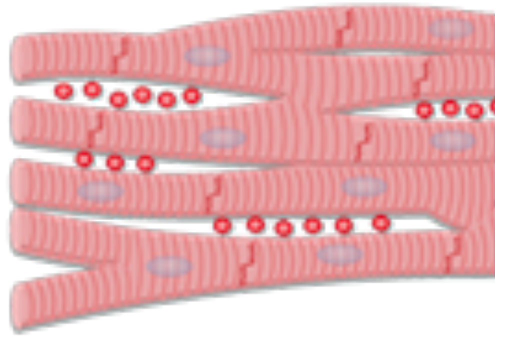
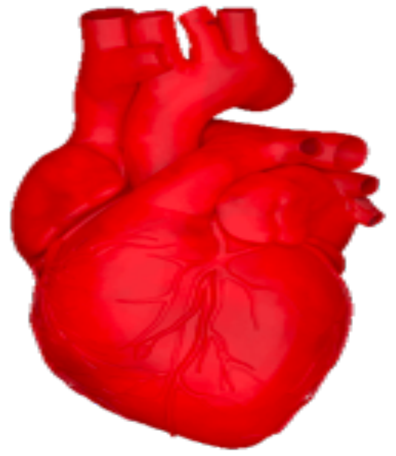
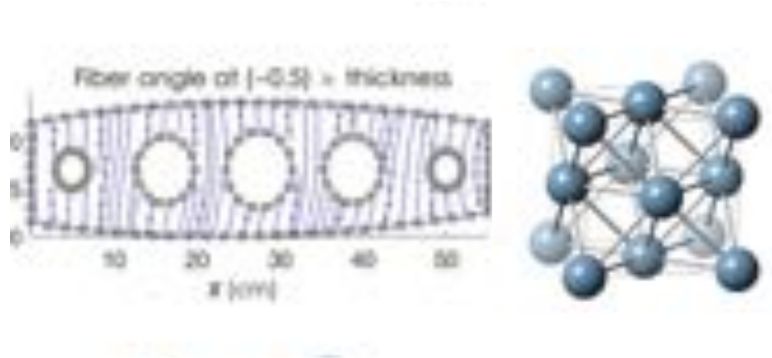
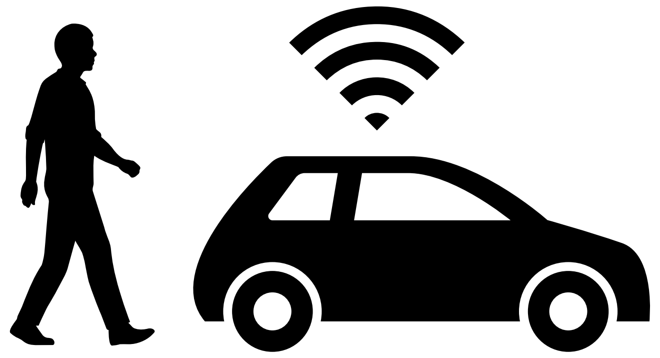
Cardiology







Complex multi-scale simulation systems

	Basic unit	Sub-component	System
Cardiology			
Aerospace			
Self-driving			

Complex multi-scale simulation systems

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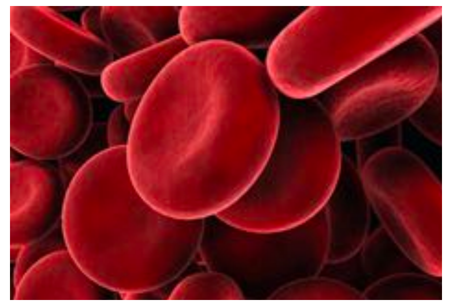
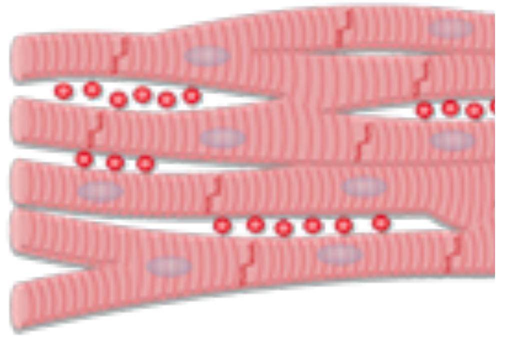
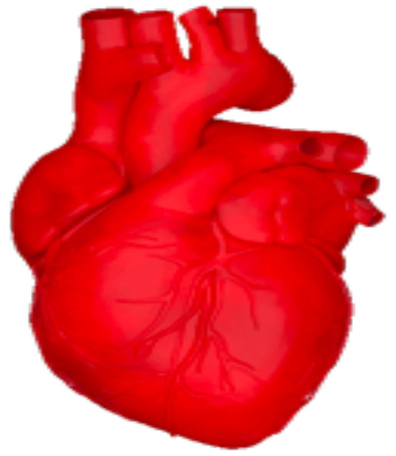
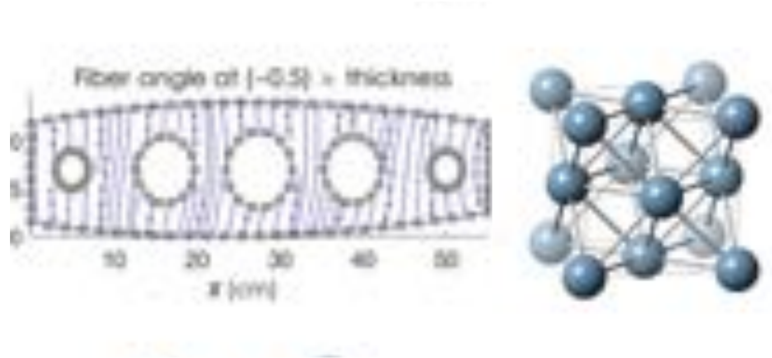
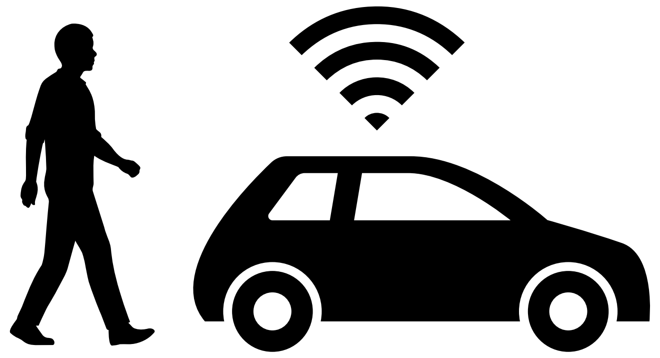
Cardiology			
Aerospace			
Self-driving			

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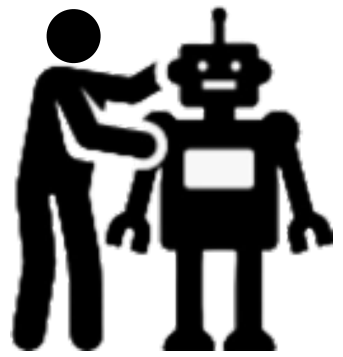
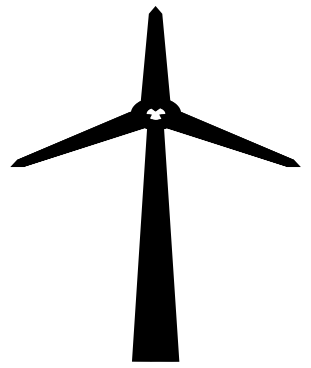


Human-robot interaction

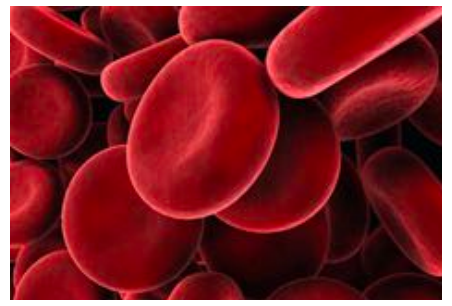
Built environment (city planning)

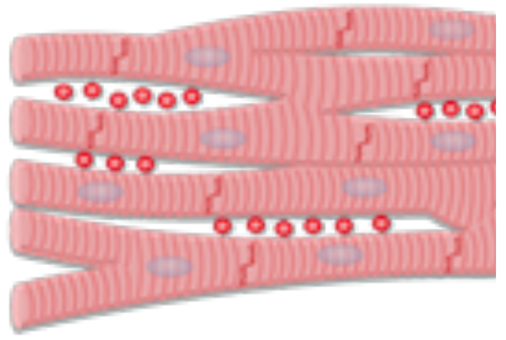


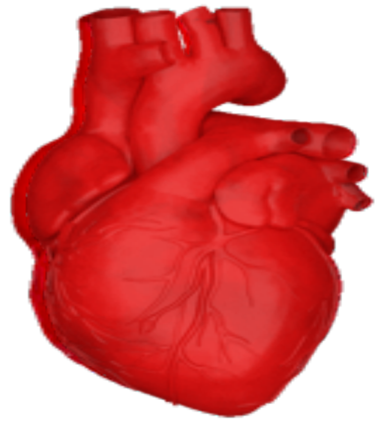
Plant design

Computational cardiology: Personalized HeartBeats

Cell model — Tissue — Heart model

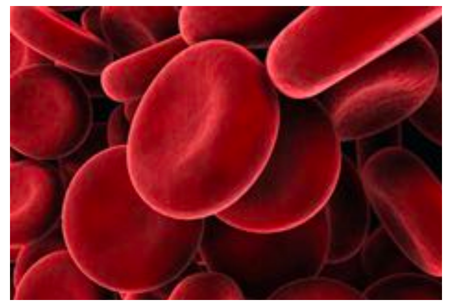




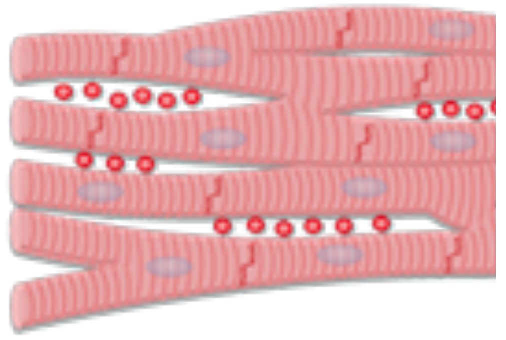


Computational cardiology: Personalized HeartBeats

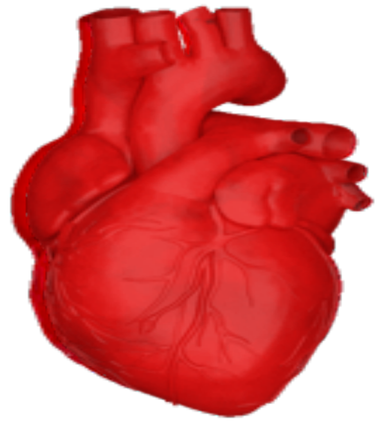
Cell model



Tissue

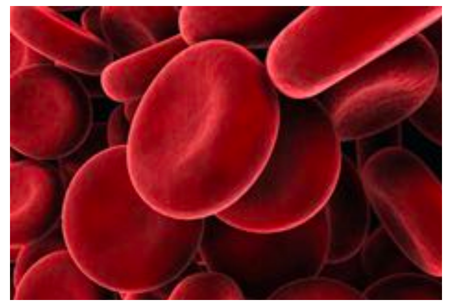


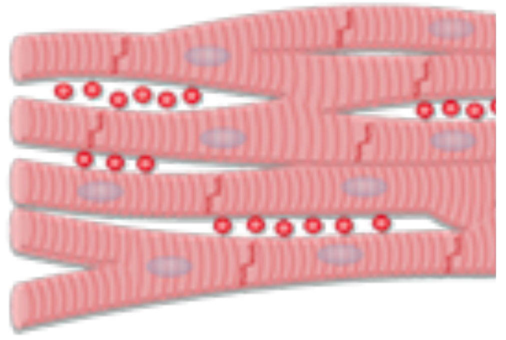
Heart model

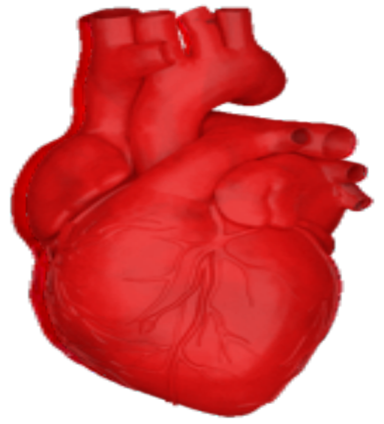


Computational cardiology: Personalized HeartBeats

Cell model | Tissue | Heart model







gif credits
alperdurmaz

- Dysregulation of calcium signaling in heart cells can cause lethal arrhythmias

[Augustin+ '16, Colman '19, Riabiz+ '21, Niederer+ '21]

Computational cardiology: Personalized HeartBeats

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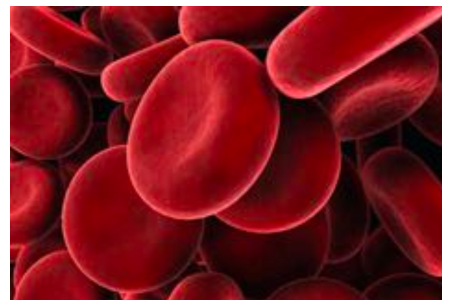
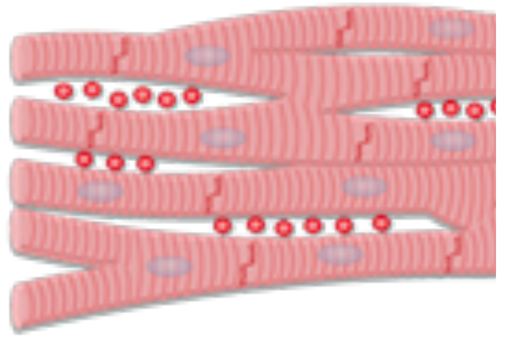
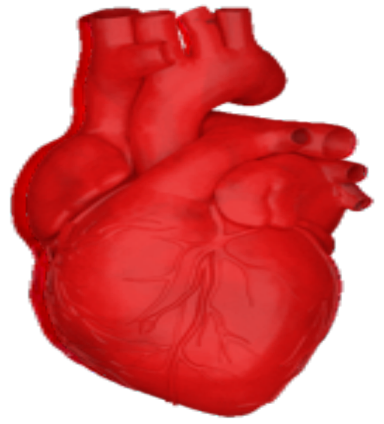

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- Task: **Simulate** multi-scale **digital twin** models of heart for **personalized predictions** of dysregulation's effect on a patient's heartbeat

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Computational cardiology: Personalized HeartBeats

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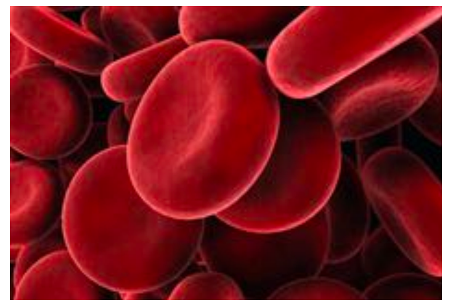

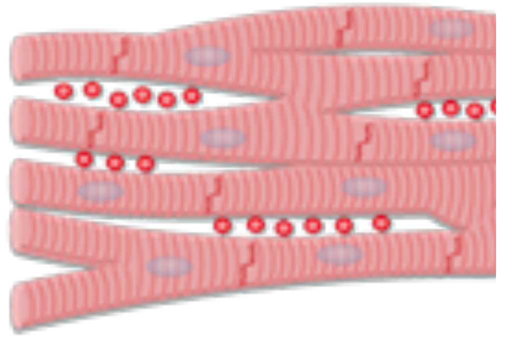

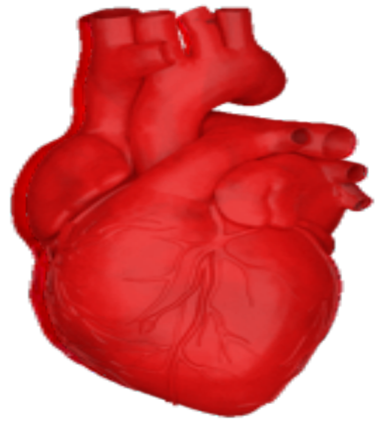

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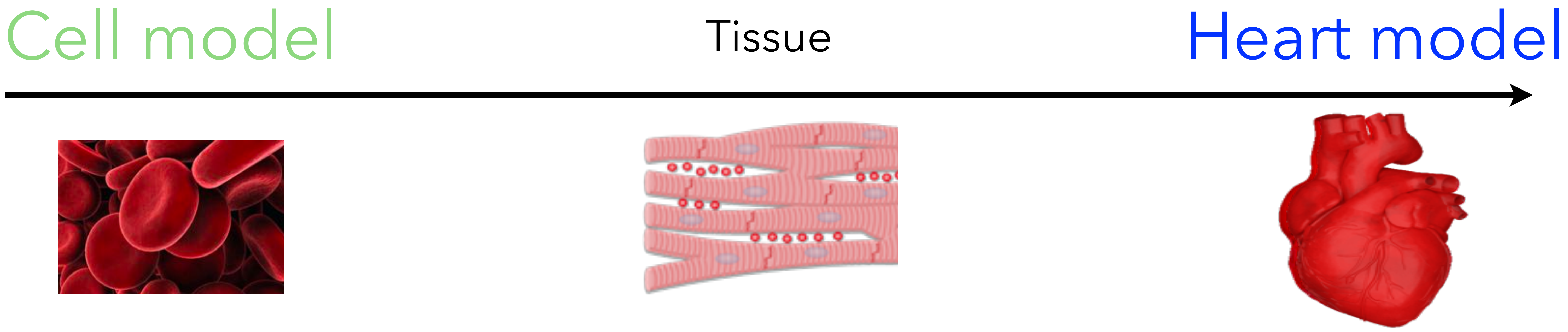
Computational cardiology: Personalized HeartBeats



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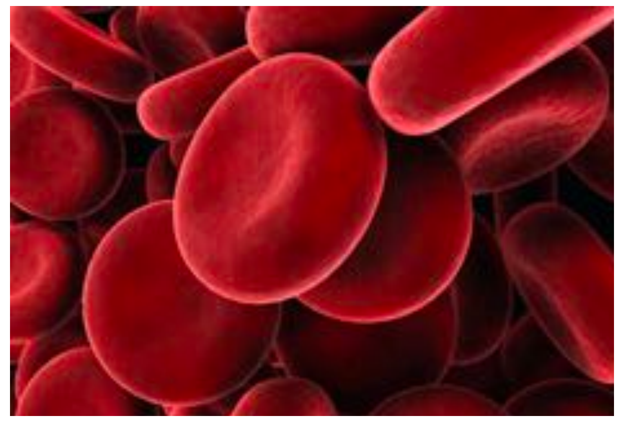
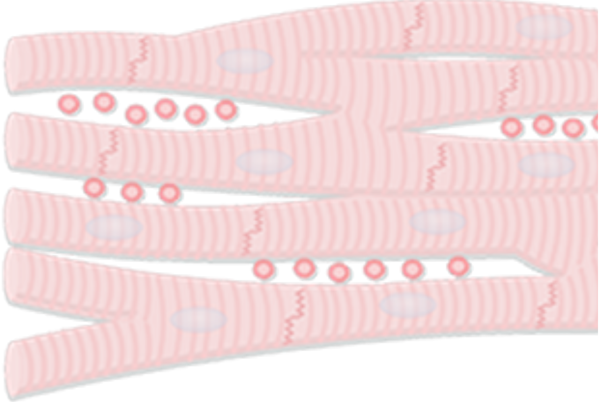
2. **Propagate** cell-model **uncertainty** to whole-heart model via simulations and Monte Carlo **integration**

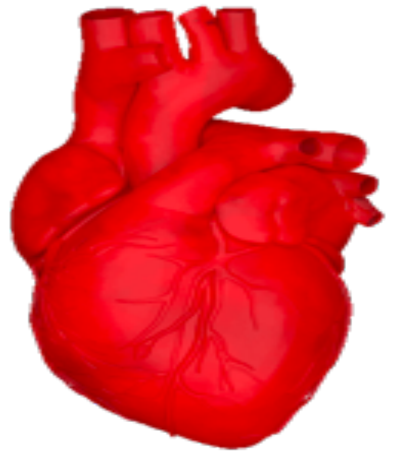
[Augustin+ '16, Colman '19, Riabiz+ '21, Niederer+ '21]

Impact of calcium signaling dysregulation on heartbeat—Two-stage inferential pipeline

Cell model X



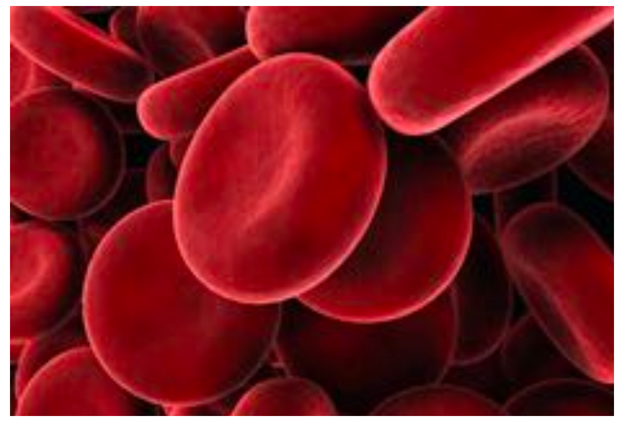


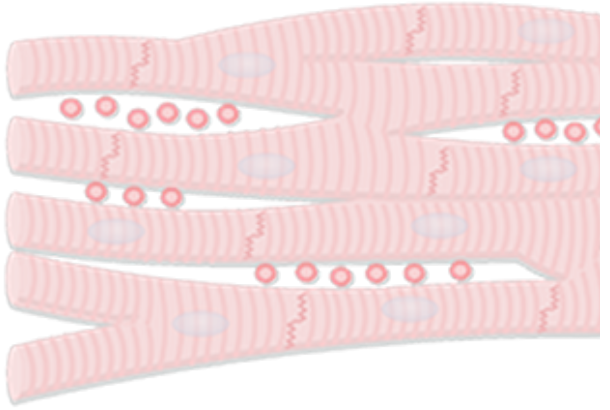


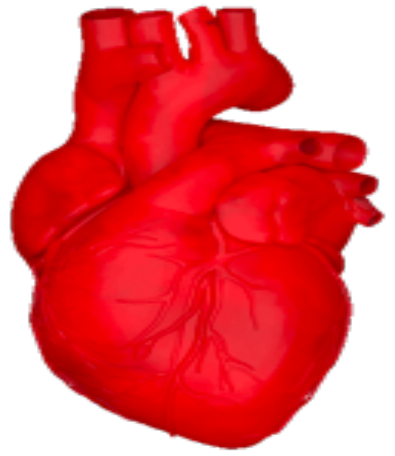
Heart model f

Impact of calcium signaling dysregulation on heartbeat—Two-stage inferential pipeline

Cell model X







Heart model f

1. Random sampling via MCMC

$$X_1, \ldots, X_T \sim \mathbb{P}^\star$$

(posterior in $\mathbb{R}^{38}$)

Impact of calcium signaling dysregulation on heartbeat—Two-stage inferential pipeline

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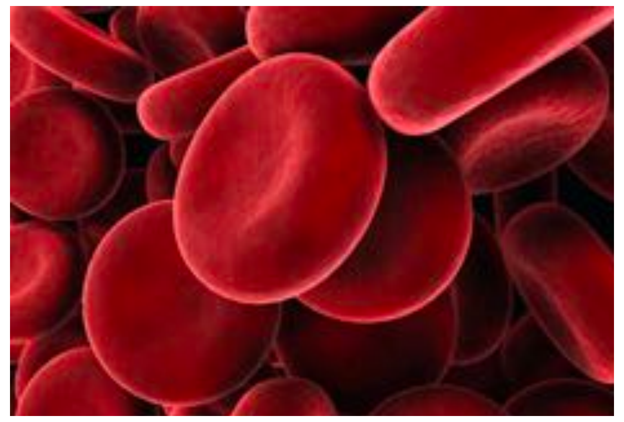

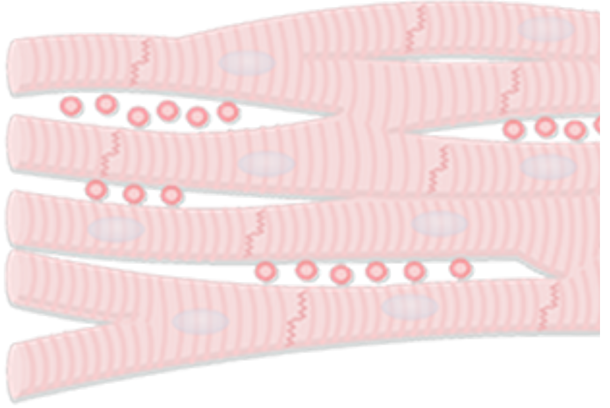

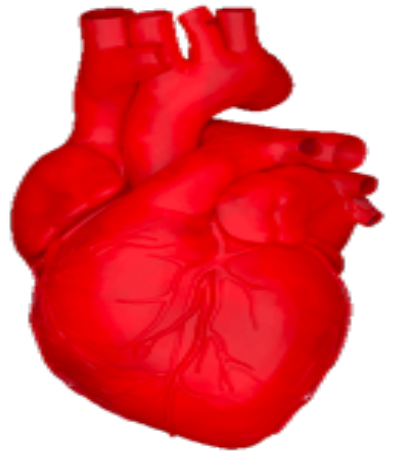

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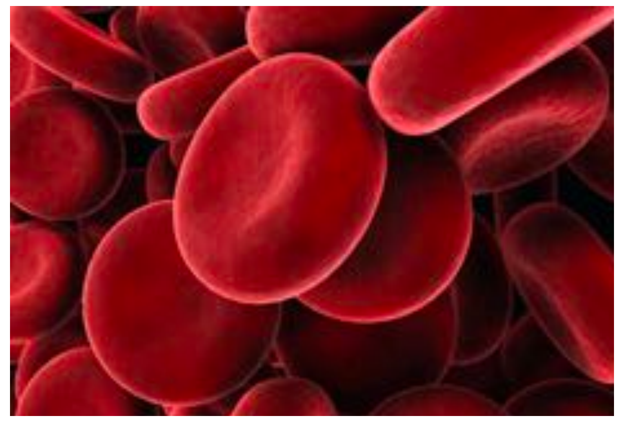
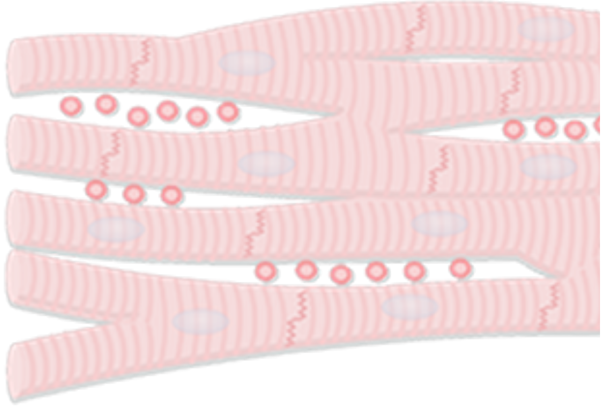
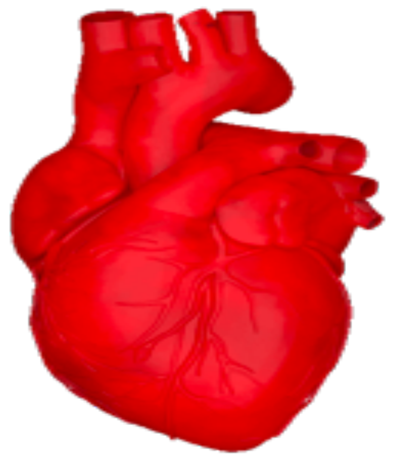
Standard tasks but computationally challenging…

Cell model X







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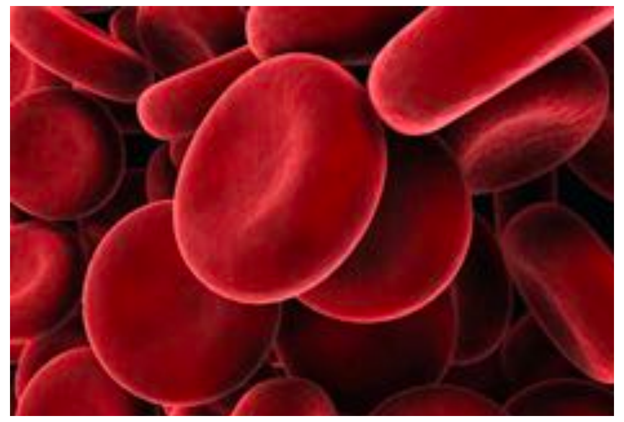
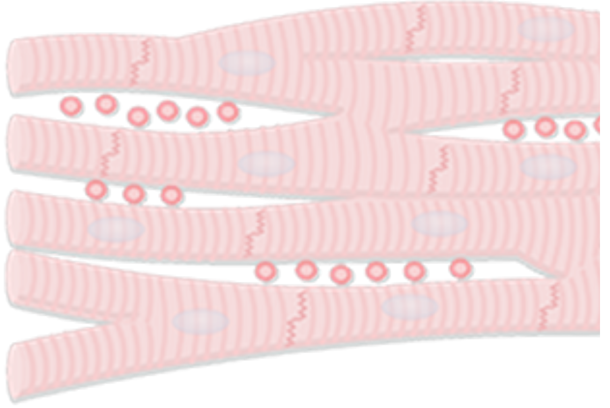
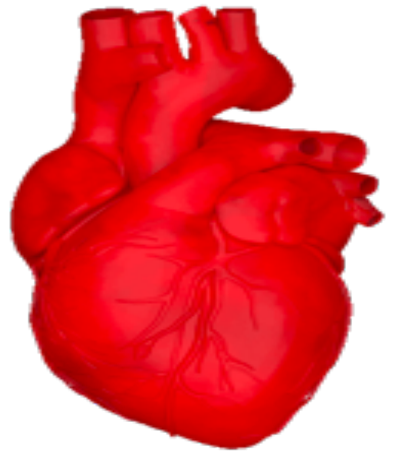
- $T = 10^6$ to explore $\mathbb{P}^\star$ well

Standard tasks but computationally challenging...

Cell model X







Heart model f

1. Random sampling via MCMC

$$X_1, \dots, X_T \sim \mathbb{P}^\star$$

(posterior in $\mathbb{R}^{38}$)

2. Uncertainty propagation via Monte Carlo integration (mean, variance,..)

$$\mathbb{P}^\star f \triangleq \int f(X) d\mathbb{P}^\star(X) \approx \frac{1}{T}\sum_{i=1}^{T} f(X_i)$$

- $T = 10^6$ to explore $\mathbb{P}^\star$ well
- Time to run **MCMC** **~ 2 CPU weeks**

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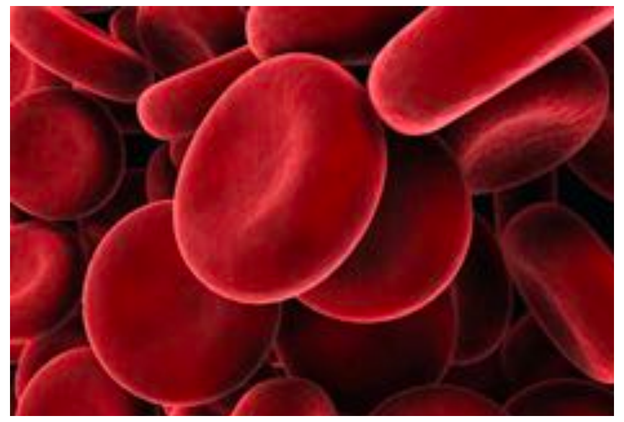

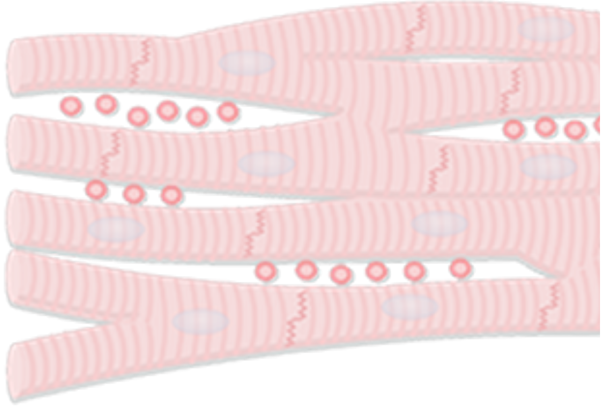

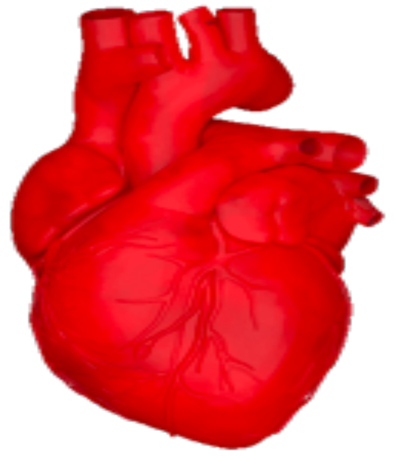

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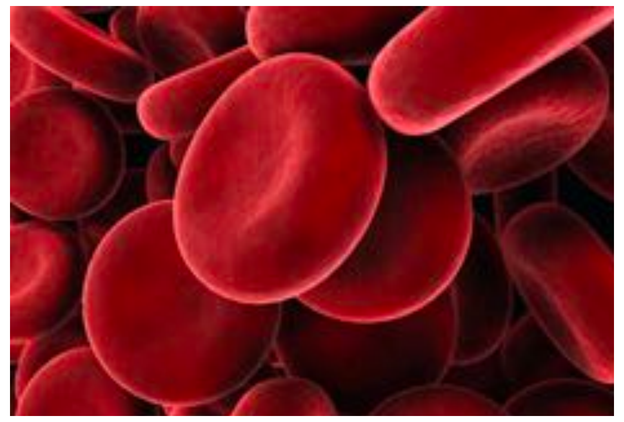

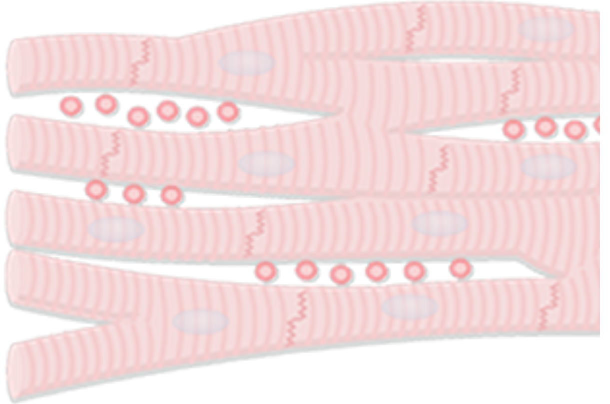

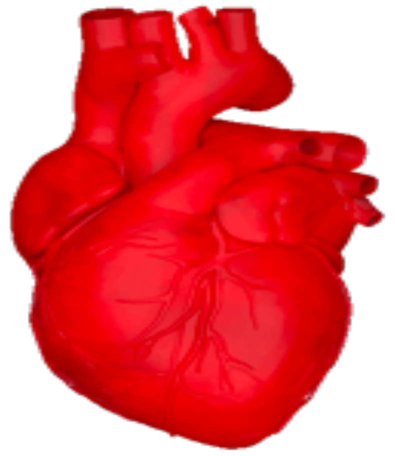

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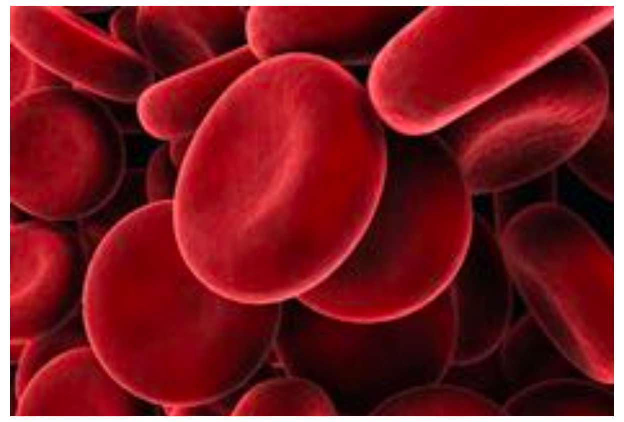

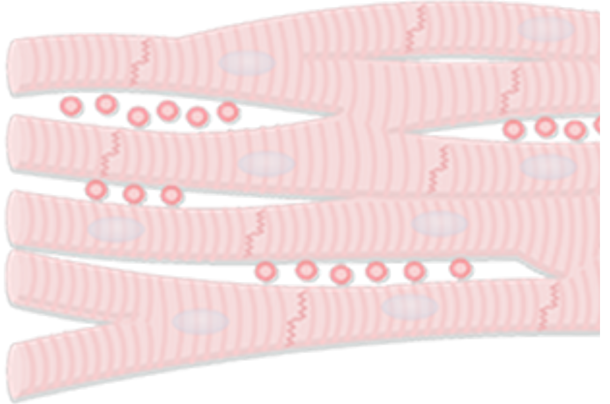

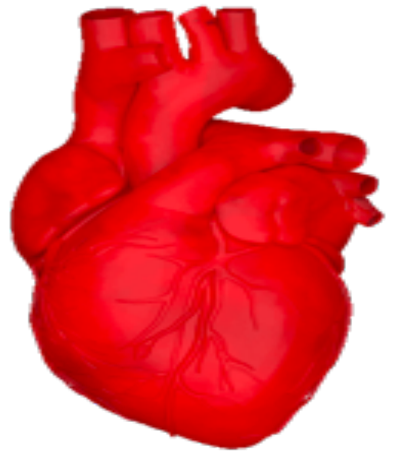

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??

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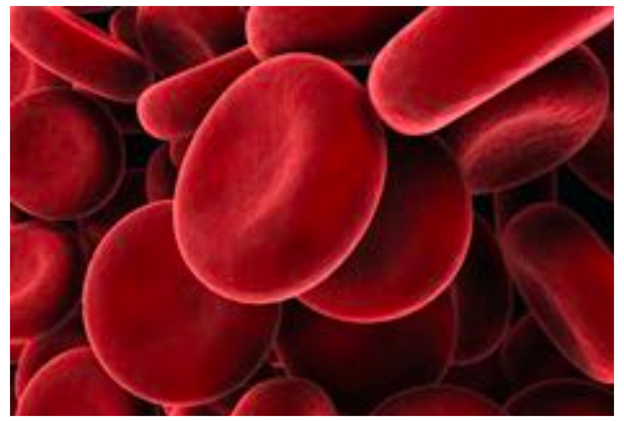
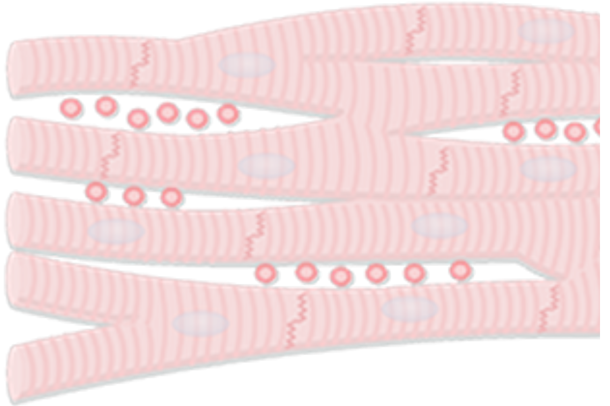
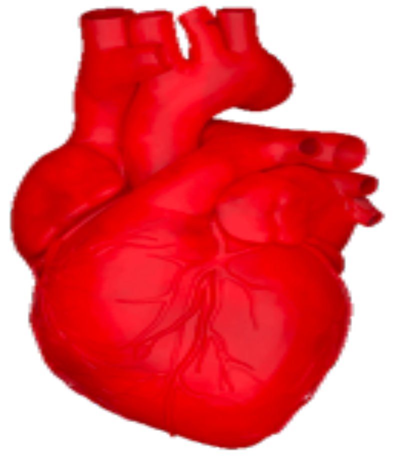
Part 2 overview: Computationally-efficient integration for high-dimensional models

Cell model X







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This talk

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Efficient integration via distribution compression

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T IID or MCMC points

$$X_1, \ldots, X_T$$

$$\mathbb{P}_T f \triangleq \frac{\Sigma_{i=1}^T f(X_i)}{T}$$

Efficient integration via distribution compression

T IID or MCMC points

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$$\mathbb{P}_T f \triangleq \frac{\Sigma_{i=1}^{T} f(X_i)}{T}$$

s output points (coreset)

Compress $\longrightarrow$

$$X'_1, \ldots, X'_s$$

$$\mathbb{P}_{out} f \triangleq \frac{\Sigma_{i=1}^{s} f(X'_i)}{s}$$

s (fewer) function evaluations

Efficient integration via distribution compression

T IID or MCMC points

s output points (coreset)

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s (fewer) function evaluations

$$|\mathbb{P}^\star f - \mathbb{P}_T f| = \Theta(T^{-1/2})$$

Efficient integration via distribution compression

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s (fewer) function evaluations

Standard thinning

(take every T/s-th point)

or iid thinning/

uniform **sub-sampling**

$$|\mathbb{P}^\star f - \mathbb{P}_T f| = \Theta(T^{-1/2})$$

$$|\mathbb{P}^\star f - \mathbb{P}_{out} f| = \Theta(s^{-1/2})$$

$$|\mathbb{P}^\star f - \mathbb{P}_{out} f| = \Theta(T^{-1/4})$$

when $s = T^{1/2}$

a million → a thousand

T IID or MCMC points

a million → a thousand

$T^{1/2}$ output points

Standard thinning

$$|\mathbb{P}^\star f - \mathbb{P}_T f| = \Theta(T^{-1/2}) \longrightarrow |\mathbb{P}^\star f - \mathbb{P}_{out} f| = \Theta(T^{-1/4})$$

What is the best error we can hope for?

T IID or MCMC points   a million → a thousand $T^{1/2}$ output points

$$|\mathbb{P}^\star f - \mathbb{P}_T f| = \Theta(T^{-1/2}) \xrightarrow{\text{Standard thinning}} |\mathbb{P}^\star f - \mathbb{P}_{out} f| = \Theta(T^{-1/4})$$

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$\Omega(T^{-1/2})$ minimax **lower bound**

- If output = $T^{1/2}$ points
- If input = T IID points (any estimator)

[Tolstikhin+ '17, Philips+ '20]

Prior strategies for efficient integration

T IID or MCMC points

a million → a thousand

$T^{1/2}$ output points

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Special $\mathbb{P}^\star$
- Uniform on $[0,1]^d$
- Bounded support & special function class

- $o(T^{-1/4})$ error guarantee: Quasi Monte Carlo, Bayesian quadrature, determinantal point processes, Haar thinning
[O'Hagan '91, Hickernell '98, Novak+'10, Liu+ '18, Karvonen+'18, **Dwivedi+**'19, Belhadji+ '20]

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Generic $\mathbb{P}^\star$ & rich function class

$\tilde{O}(T^{-1/4})$ error guarantee:
Kernel herding, greedy sign selection, Stein points MCMC, support points, supersampling
[Chen+ '10, Lacoste+ '15, Paige+ '16, Tolstikhin+ '17, Mak+ '18, Chen '19, Karnin '19]

$\Omega(T^{-1/2})$ minimax lower bound

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A new practical & provably near-optimal procedure

T IID or MCMC points   a million → a thousand $T^{1/2}$ output points

Standard thinning

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Kernel thinning

Dwivedi and Mackey '21, '22

$$|\mathbb{P}^\star f - \mathbb{P}_{out} f| = \tilde{O}(T^{-1/2})$$

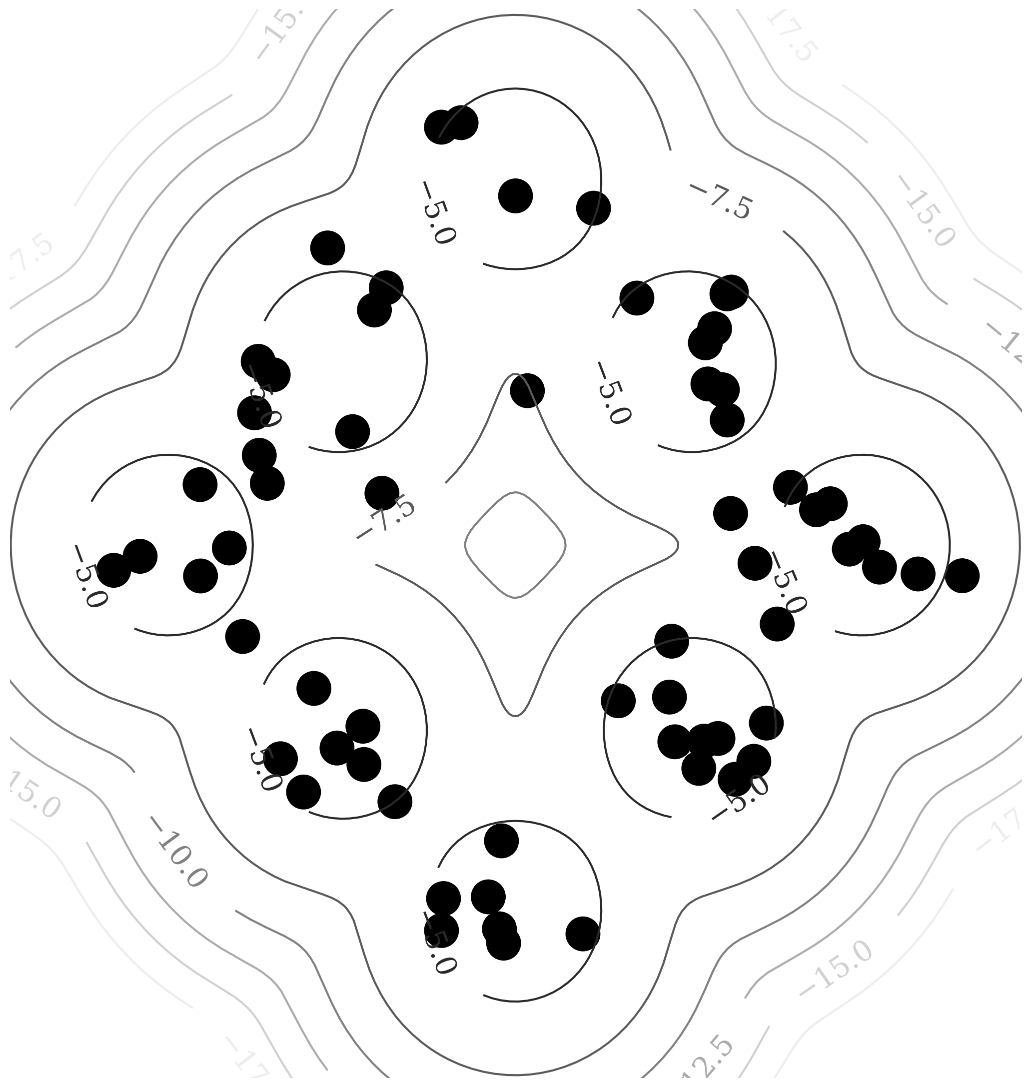
✓ for generic $\mathbb{P}^\star$ on generic domains
✓ for rich function classes

$\Omega(T^{-1/2})$ minimax lower bound

Visual comparison on $\mathbb{P}^\star$ = 8 mixture of Gaussian

64 iid input points

8 output points



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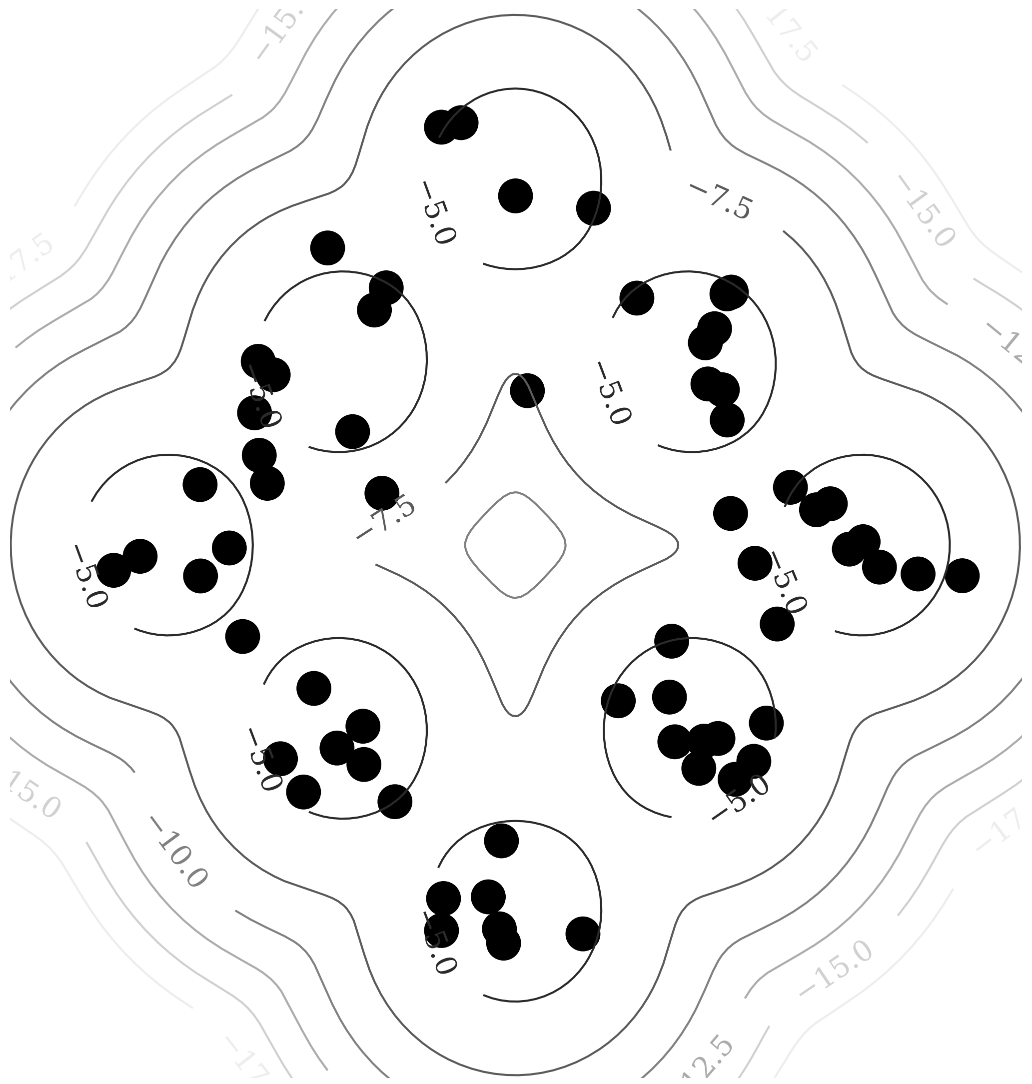

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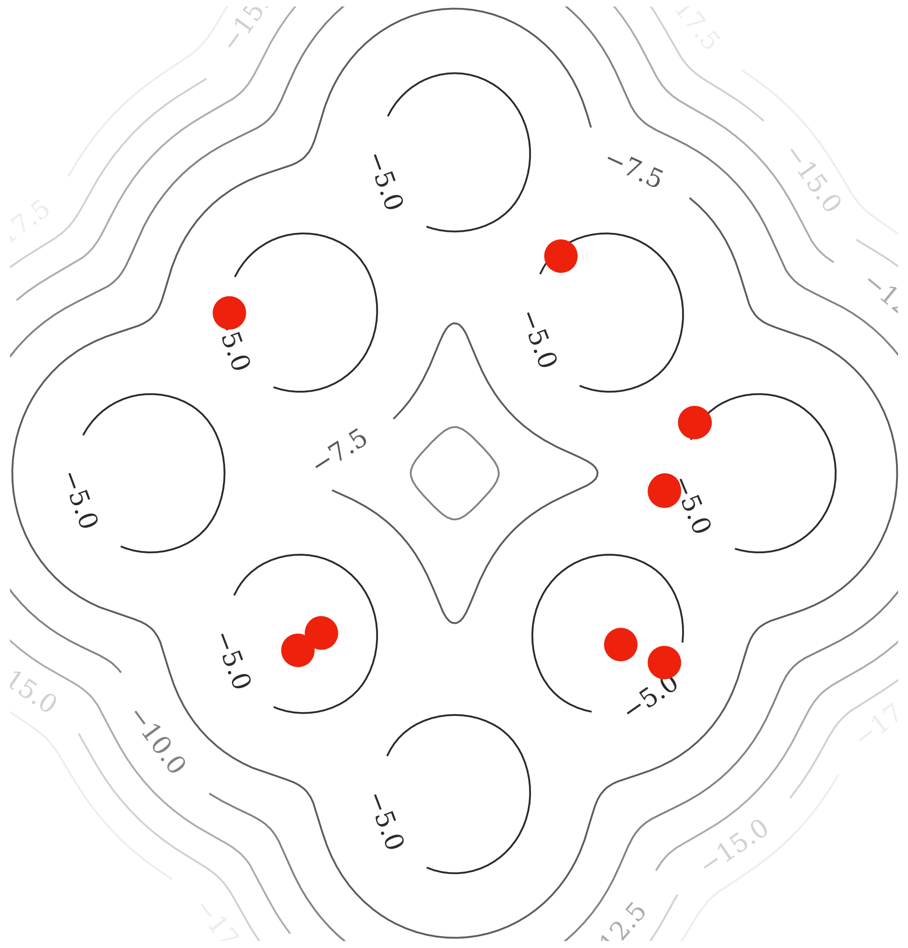

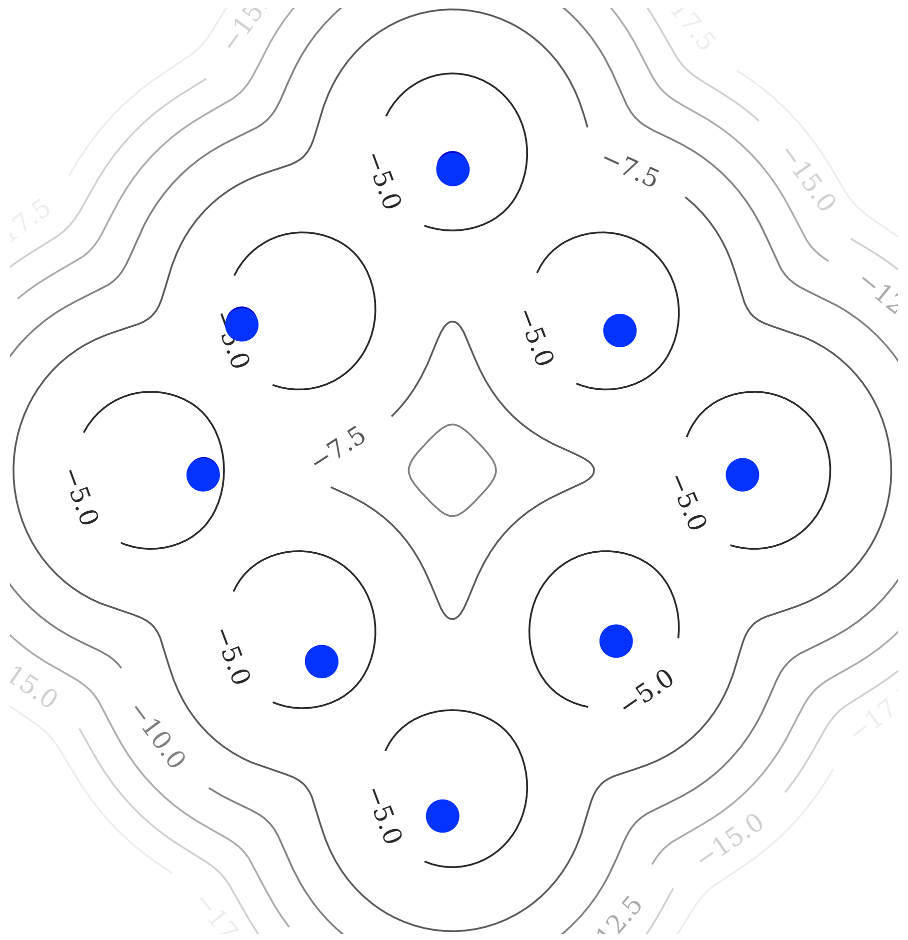

Quantitative measure: Worst-case error over a rich class

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Namely, over the unit ball of a reproducing kernel Hilbert space (RKHS)

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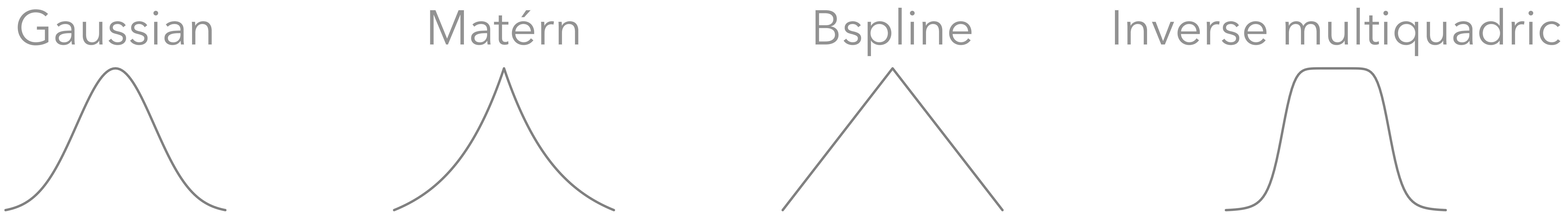
  any symmetric ($\mathbf{k}(x, y) = \mathbf{k}(y, x)$) and positive semidefinite function



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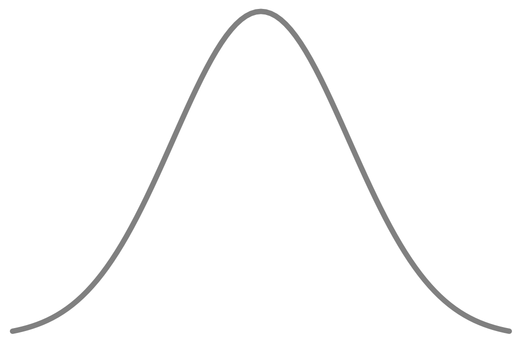
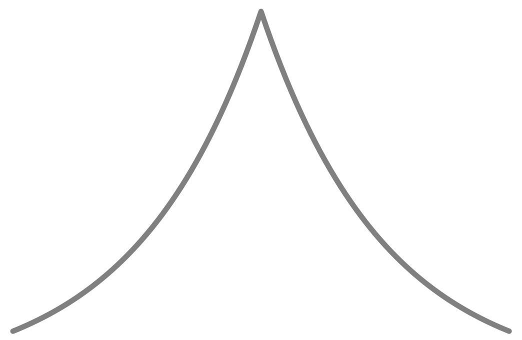
- **Metrizes convergence in distribution** for popular infinite-dimensional $\mathbf{k}$

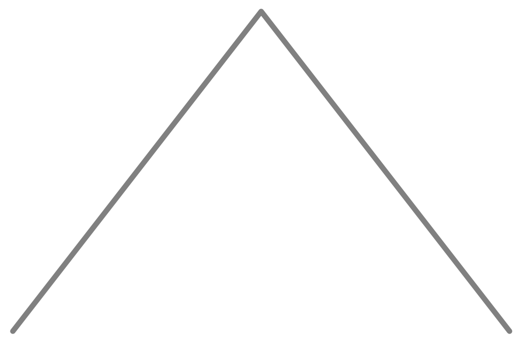
Gaussian



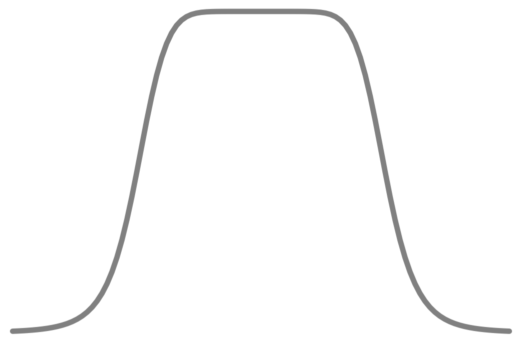
Matérn



Bspline



Inverse multiquadric



Main result: A high probability bound for generic $\mathbb{P}^\star$ and $\mathbf{k}$

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- $\sup_{\|f\|_{\mathbf{k}} \leq 1} |\mathbb{P}^\star f - \mathbb{P}_{out} f| \lesssim \sqrt{\frac{\log^{d/2+1} T}{T}}$ Sub-gaussian $\mathbb{P}^\star$ and $\mathbf{k}$ on $\mathbb{R}^d$ (Gaussian)

  $\lesssim \sqrt{\frac{\log^{d+1} T}{T}}$ Sub-exponential $\mathbb{P}^\star$ and $\mathbf{k}$ on $\mathbb{R}^d$ (Matérn)

- A near-quadratic gain over $T^{-1/4}$ standard thinning error
- Matches minimax lower bounds $T^{-1/2}$ up to log factors

Kernel thinning

points $X_1, X_2, \ldots, X_T$ $\rightarrow$ **Kernel** $\rightarrow$ functions in RKHS $v_1, v_2, \ldots, v_T$

Kernel thinning ≡ Recursive halving via kernel evaluations

points $X_1, X_2, \ldots, X_T$ → **Kernel** → functions in RKHS $v_1, v_2, \ldots, v_T$

Input T points → Kernel halving → $T/2$ points → $O(\log T)$ Kernel halving rounds → **Output** $\sqrt{T}$ points + some refinement

Kernel halving

points $X_1, X_2, \ldots, X_T$ → Kernel → functions in RKHS $v_1, v_2, \ldots, v_T$

$v_1, v_2, \ldots, v_T$ → Kernel halving → $v'_1, v'_2, \ldots, v'_{T/2}$

$$\left| \frac{\Sigma_{i=1}^{T} v_i}{T} - \frac{\Sigma_{i=1}^{T/2} v'_i}{T/2} \right| = \text{small}$$

Kernel halving $\equiv$ Discrepancy minimization problem

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$\equiv$

Assign $\varepsilon_i \in \{-1,1\}$ to v_i
such that $|\Sigma_{i=1}^{T} \varepsilon_i v_i|$ is small
& output points with $\varepsilon_i = -1$

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$$\sum_{i=1}^{T} \varepsilon_i v_i = \sum_{\varepsilon_i=+1} v_i - \sum_{\varepsilon_i=-1} v_i$$

Kernel halving ≡ Discrepancy minimization problem

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$$= \sum_{i=1}^{T} v_i - 2 \sum_{\varepsilon_i=-1} v_i$$

Kernel halving ≡ Discrepancy minimization problem

points

$X_1, X_2, \ldots, X_T$ → Kernel → functions in RKHS

$v_1, v_2, \ldots, v_T$

$v_1, v_2, \ldots, v_T$ → Kernel halving → $v'_1, v'_2, \ldots, v'_{T/2}$

Assign $\varepsilon_i \in \{-1,1\}$ to v_i

such that $|\sum_{i=1}^T \varepsilon_i v_i|$ is small

& output points with $\varepsilon_i = -1$

$$\left| \frac{\sum_{i=1}^T v_i}{T} - \frac{\sum_{i=1}^{T/2} v'_i}{T/2} \right| = \text{small}$$

$$\equiv$$

$$\sum_{i=1}^{T} \varepsilon_i v_i = \sum_{\varepsilon_i=+1} v_i - \sum_{\varepsilon_i=-1} v_i$$

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$$\frac{\sum_{i=1}^T \varepsilon_i v_i}{T} = \frac{\sum_{i=1}^T v_i}{T} - \frac{\sum_{\varepsilon_i=-1} v_i}{T/2}$$

Kernel halving ≡ Discrepancy minimization problem

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≡

Assign $\varepsilon_i \in \{-1, 1\}$ to v_i

such that $|\sum_{i=1}^{T} \varepsilon_i v_i|$ is small

& output points with $\varepsilon_i = -1$

$$\sum_{i=1}^{T} \varepsilon_i v_i = \sum_{\varepsilon_i = +1} v_i - \sum_{\varepsilon_i = -1} v_i$$

$$= \sum_{i=1}^{T} v_i - 2 \sum_{\varepsilon_i = -1} v_i$$

$$\frac{\sum_{i=1}^{T} \varepsilon_i v_i}{T} = \frac{\sum_{i=1}^{T} v_i}{T} - \frac{\sum_{\varepsilon_i = -1} v_i}{T/2}$$

Error — Input mean — Halve output mean

KT intuition: IID vs correlated signs

$|\sum_{i=1}^{T} \varepsilon_i v_i|$ is small

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$|\sum_{i=1}^{T} \varepsilon_i v_i|$ is small

$\varepsilon_i = \pm 1$ with equal probability

KT intuition: IID vs correlated signs

$|\Sigma_{i=1}^{T} \varepsilon_i v_i|$ is small

$\varepsilon_i = \pm 1$ with equal probability

$\sigma_T^2 = \sigma_{T-1}^2 + v_T^2$

$|\Sigma_{i=1}^{T} \varepsilon_i v_i| = O(\sigma_T) = O(T^{1/2})$

Standard thinning

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ε_i negatively correlated with $\Sigma_{j=1}^{i-1} \varepsilon_j v_j$

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ε_i negatively correlated with $\Sigma_{j=1}^{i-1}\varepsilon_j v_j$

$\sigma_T^2 \leq \beta\, \sigma_{T-1}^2 + v_T^2$ for $\beta < 1$

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Kernel thinning

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Kernel thinning

Discrepancy minimization

[… Spencer '77, Banaszczyk '98, '12, Eldan+ '18, …
Bansal+ '16, '18, '19, '20, **Dwivedi**+ '19, Alweiss+ '21, …]

Is KT better practically? Gaussian $\mathbb{P}^\star$ in $\mathbb{R}^d$

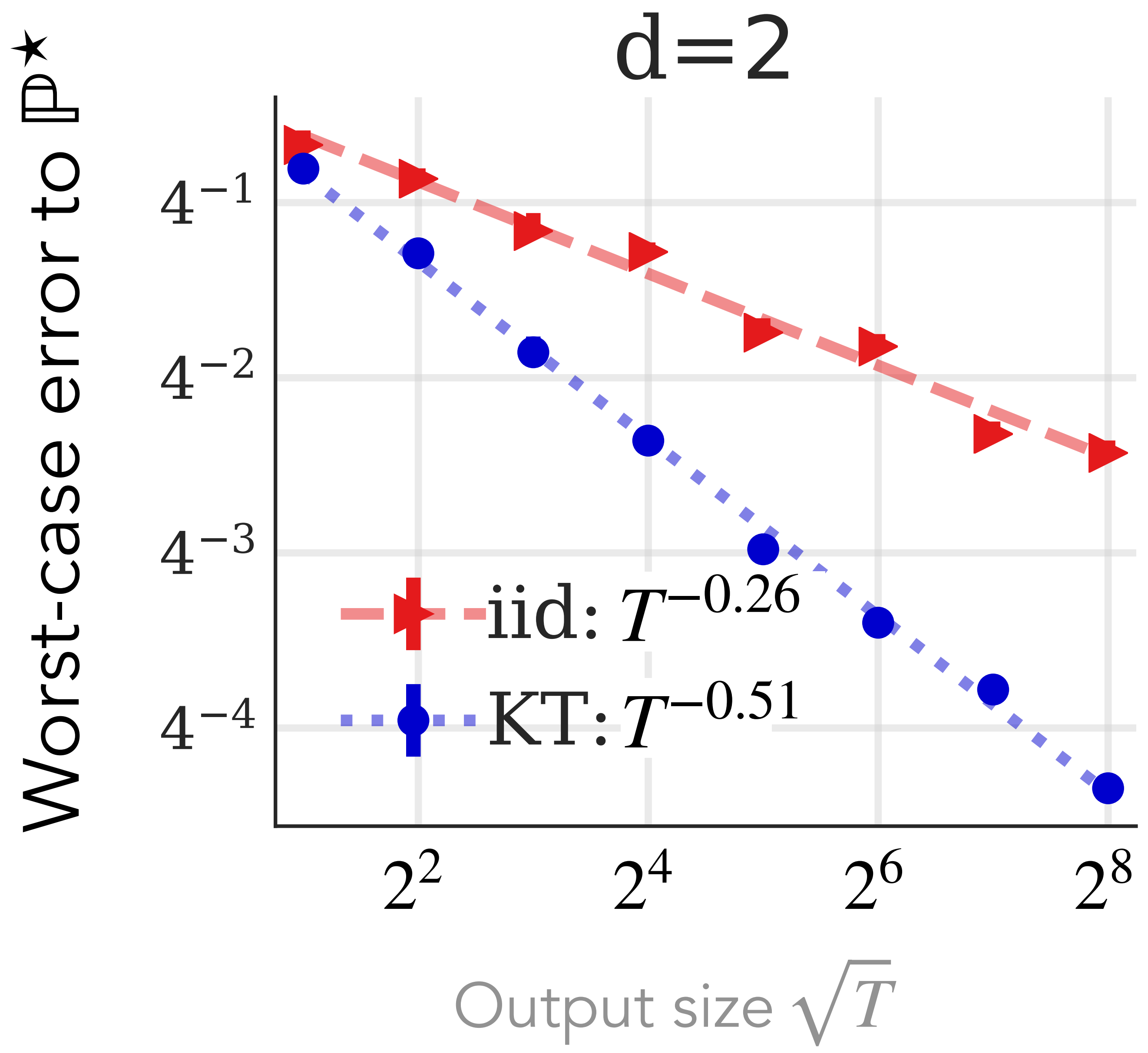
iid input, Gaussian kernel

Worst-case error to $\mathbb{P}^\star$

Output size $\sqrt{T}$

Is KT better practically? Gaussian $\mathbb{P}^\star$ in $\mathbb{R}^d$

iid input, Gaussian kernel



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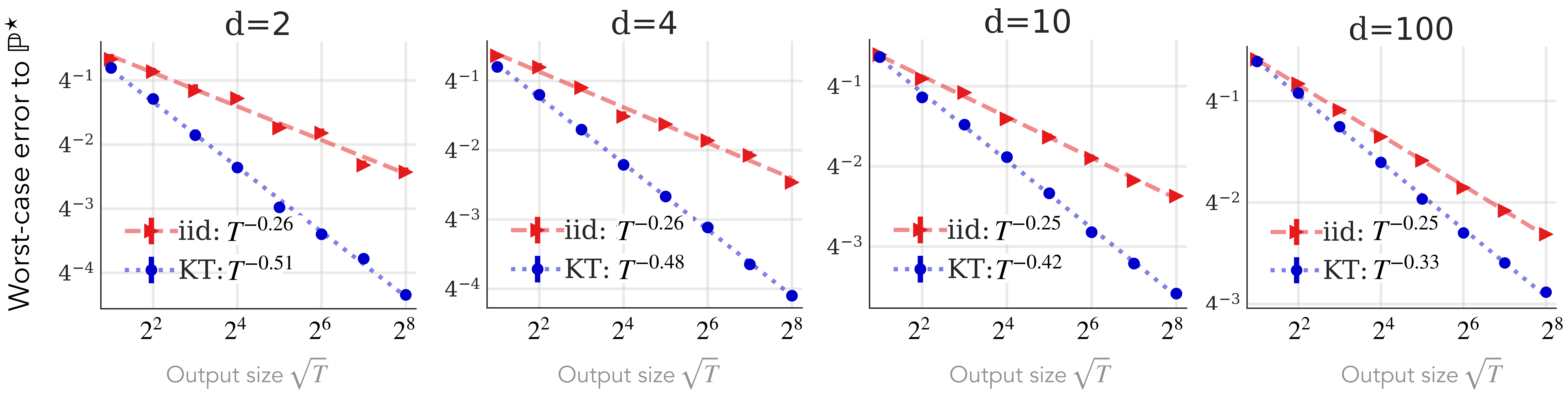

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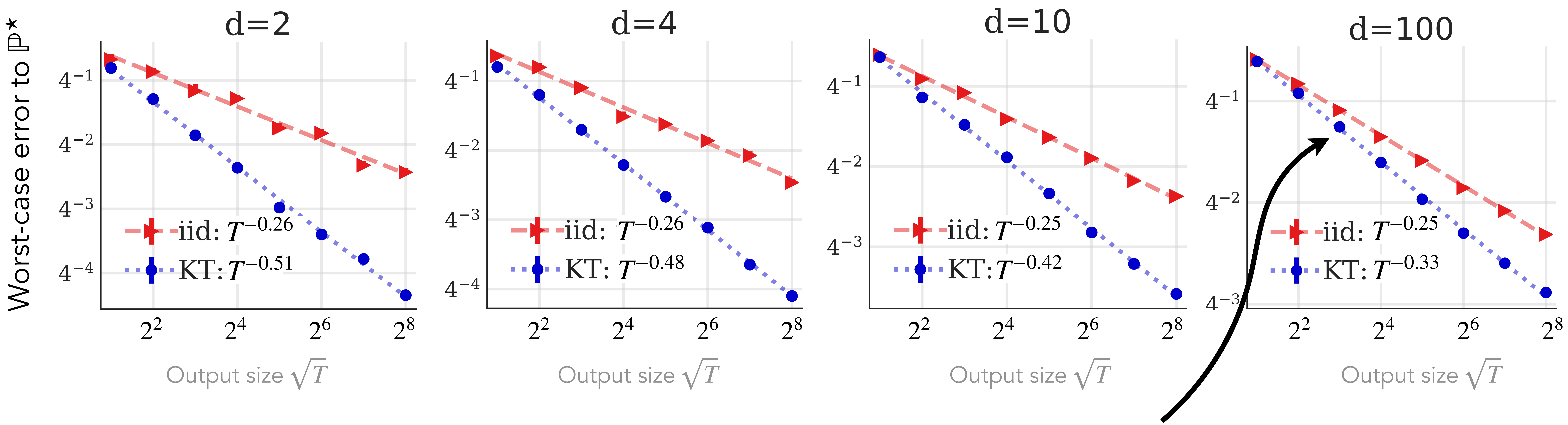
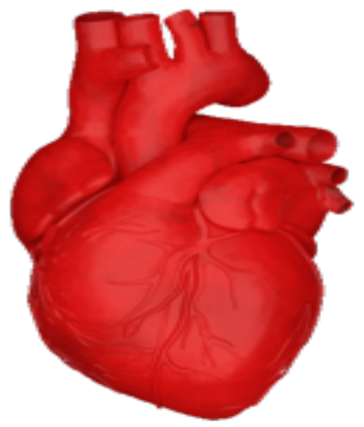


Significant gains in $d = 100$

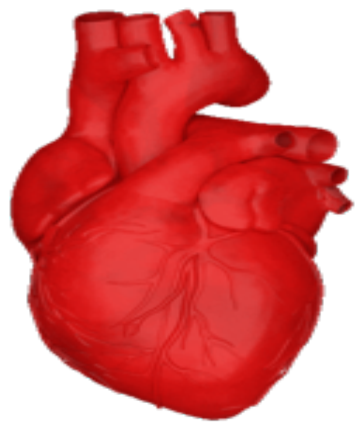
with just 8 output points

KT on MCMC points for $\mathbb{P}^\star$ in  experiments $(d = 38)$

†Input = 2 MCMC runs on 2 posteriors $\mathbb{P}^\star$, Gaussian kernel

Worst-case error to $\mathbb{P}_T$

[†MCMC data from Riabiz-Chen-Cockayne-Swietach-Niederer-Mackey-Oates '21]

KT on MCMC points for $\mathbb{P}^\star$ in  experiments ($d = 38$)

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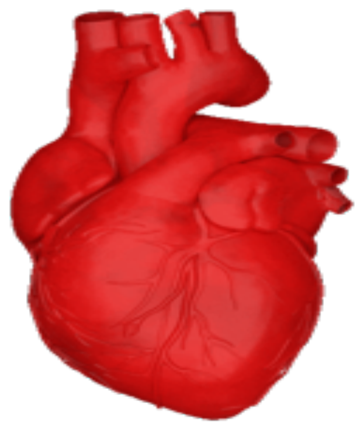
KT on MCMC points for $\mathbb{P}^\star$ in  experiments ($d = 38$)

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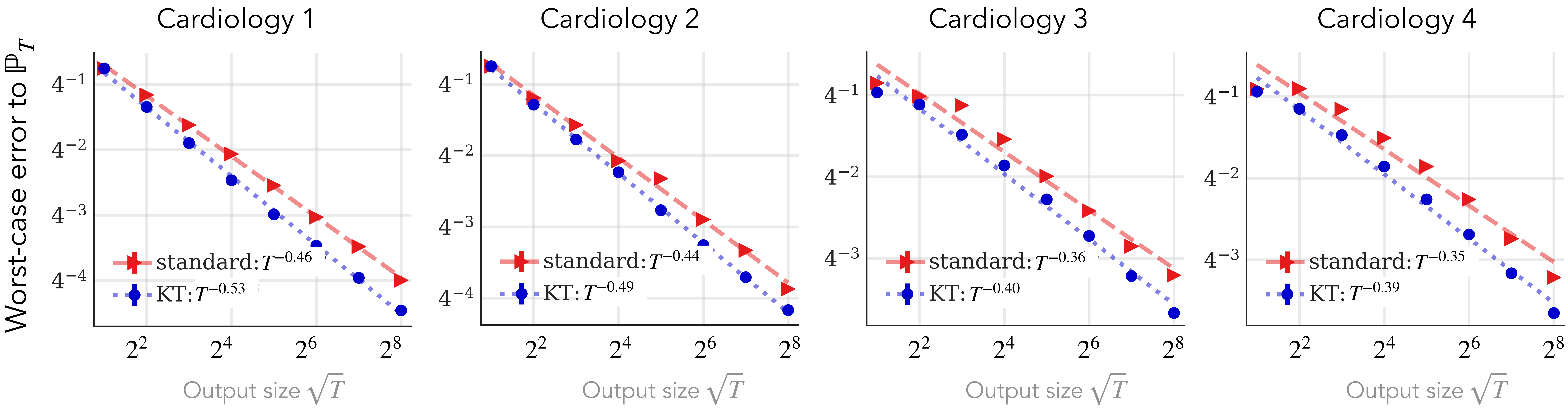


Standard thinning does well but **KT provides further improvement**
& offers **50% computational savings** (each point ~ 4 CPU weeks)

[†MCMC data from Riabiz-Chen-Cockayne-Swietach-Niederer-Mackey-Oates ’21]

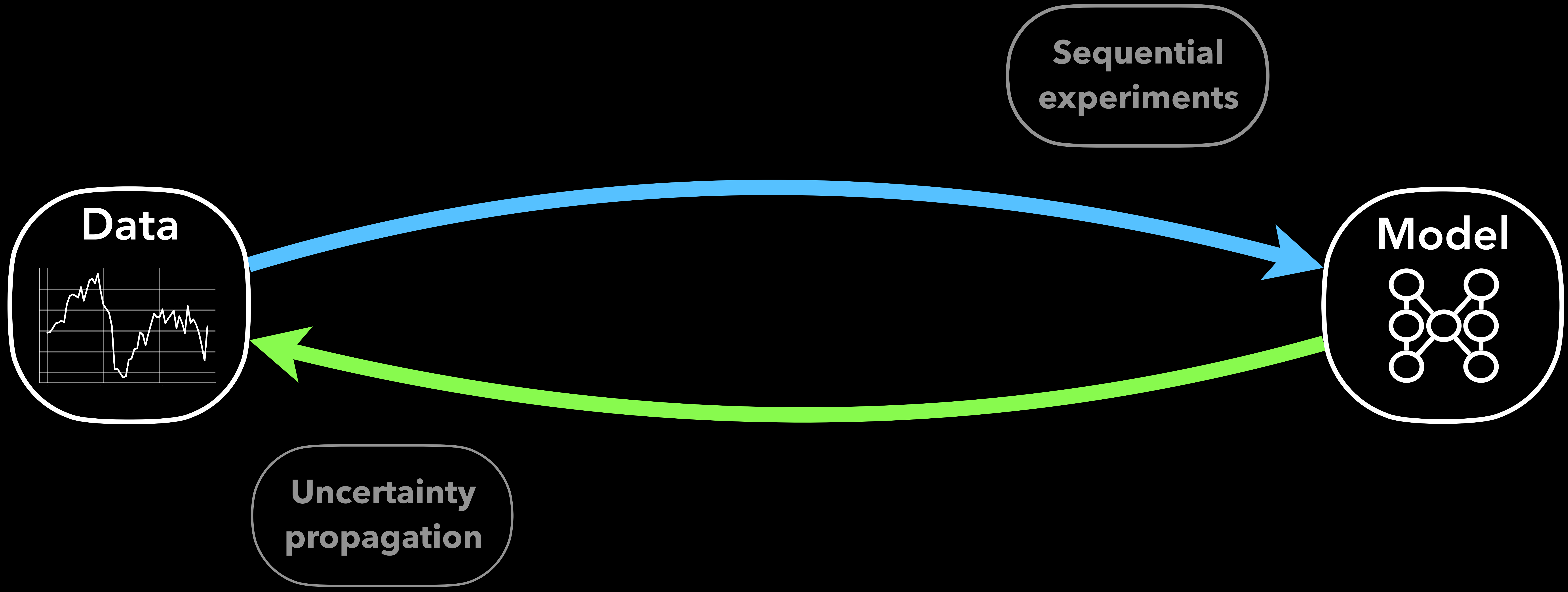
Kernel thinning: Near-optimal compression in near-linear time

Kernel thinning: Near-optimal compression in near-linear time

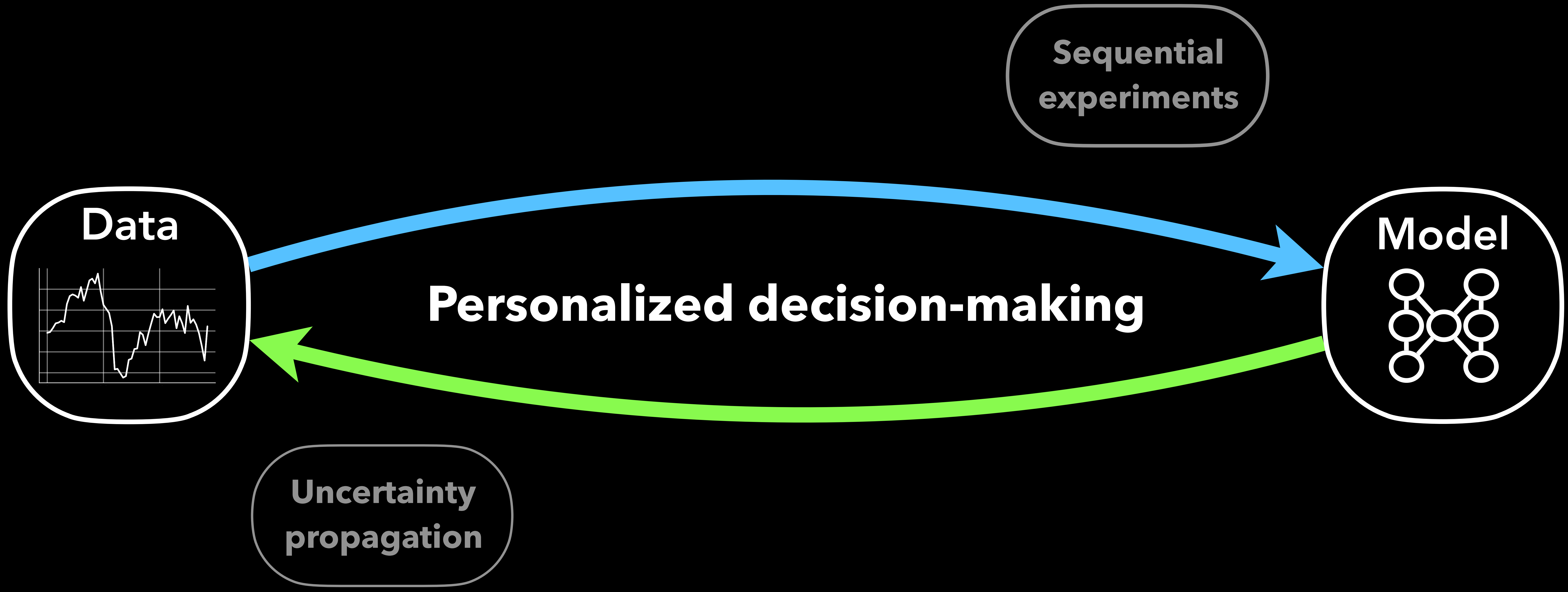
python™ **pip install goodpoints**

Thin 100k points in 100 dimensions in 10mins

talk summary



talk summary



From HeartSteps

Personalized inference by **averaging neighborhoods**

Sequential experiments

Uncertainty propagation

talk summary

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Personalized inference by **averaging neighborhoods**

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Personalized simulations by **thinning neighborhoods**

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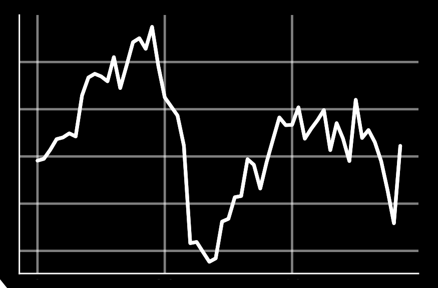
Sequential experiments
Data
Model
Uncertainty propagation

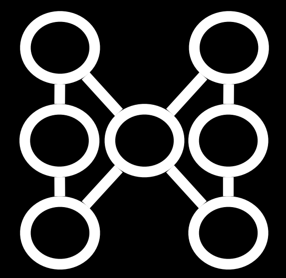
Sequential experiments

Data



Data and computation efficient methods
for personalized decision-making

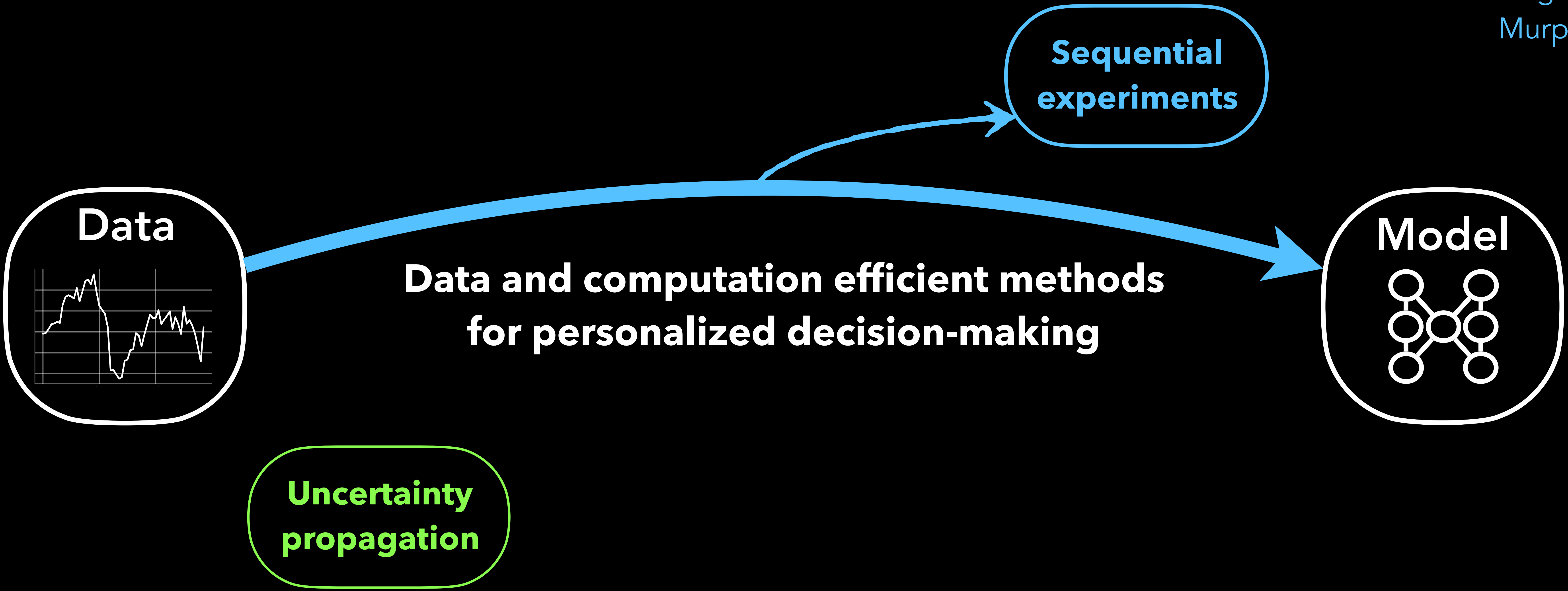
Model



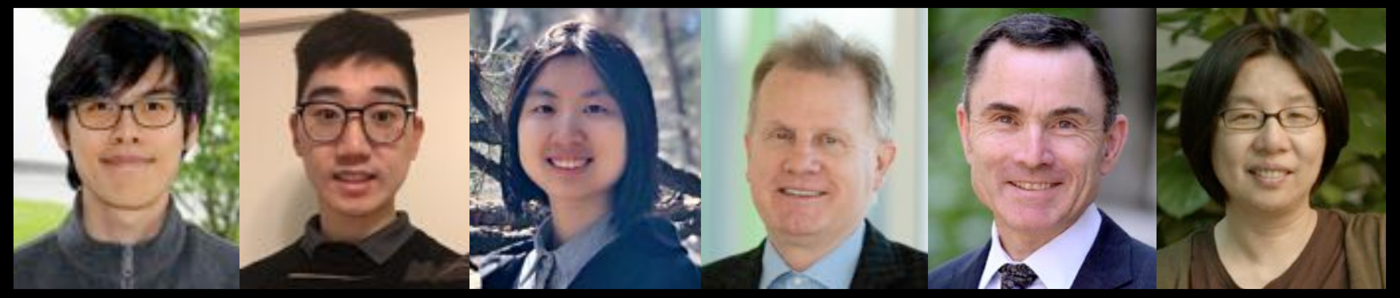
Uncertainty propagation

Deep dive into personalization by a reinforcement learning algorithm

Dwivedi*-Zhang*-Chhabria-Klasnja-Murphy '23



Stable discovery of interpretable subgroups in randomized studies via calibration



Dwivedi*-Tan*-Park-Wei-Horgan-Madigan-Yu '20

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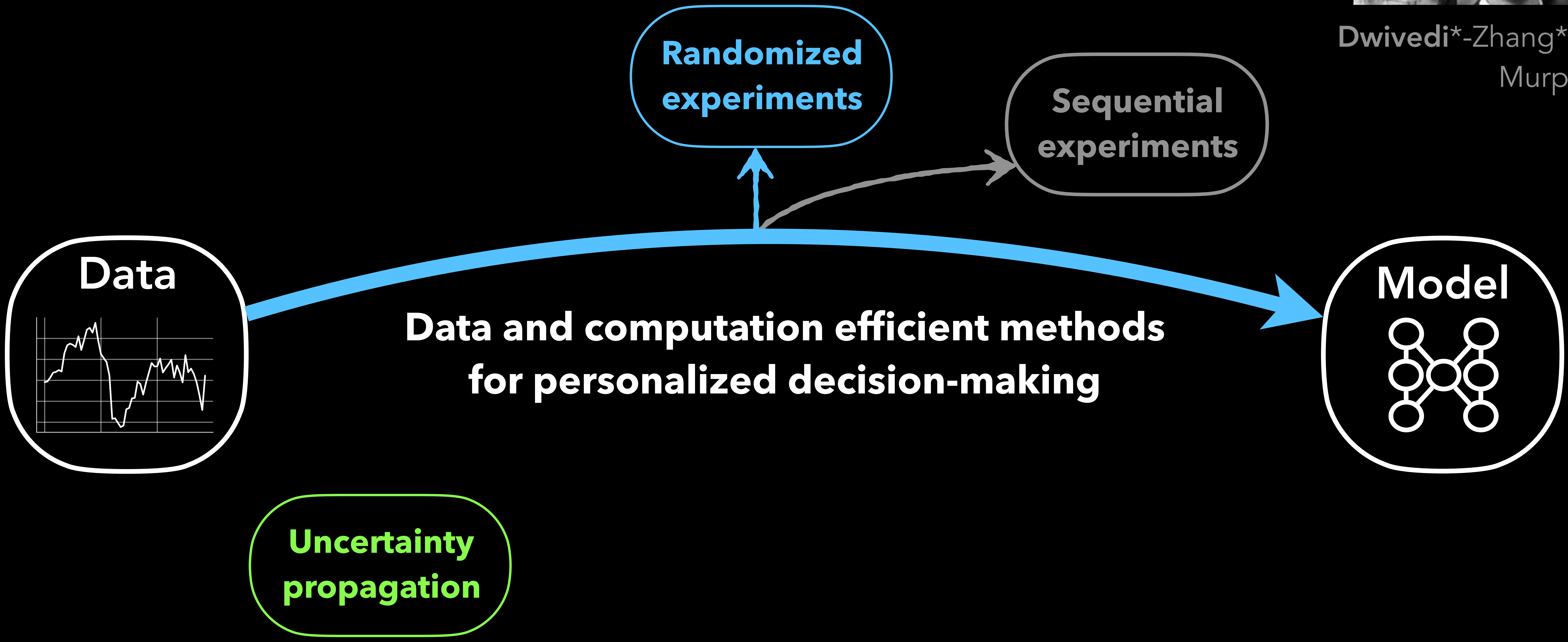

On counterfactual inference with unobserved confounding via exponential family

Shah-**Dwivedi**-Shah-Wornell '22

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Observational studies

Randomized experiments

Sequential experiments

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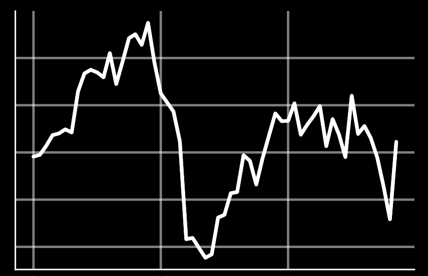

Data and computation efficient methods for personalized decision-making

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research overview

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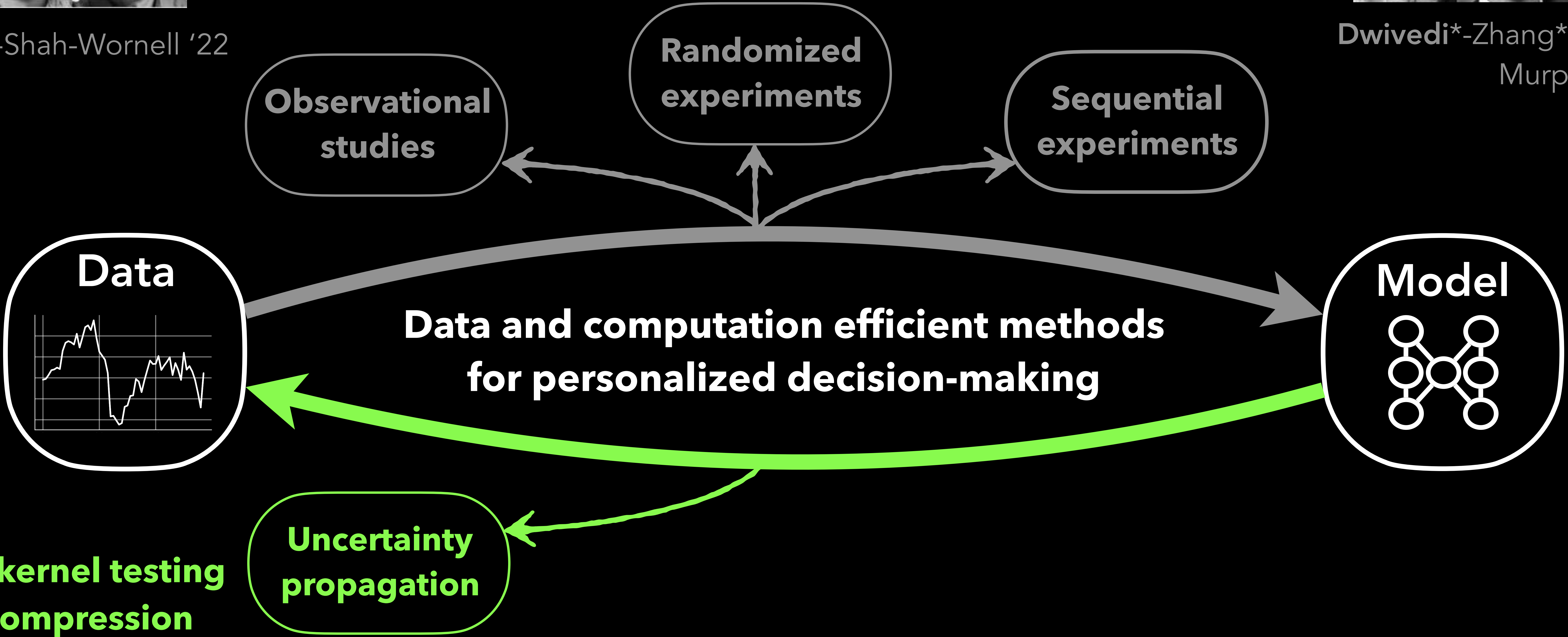

Fast and powerful kernel testing via distribution compression

Shetty-**Dwivedi**-Mackey '22,
Domingo Enrich-**Dwivedi**-Mackey '23

research overview

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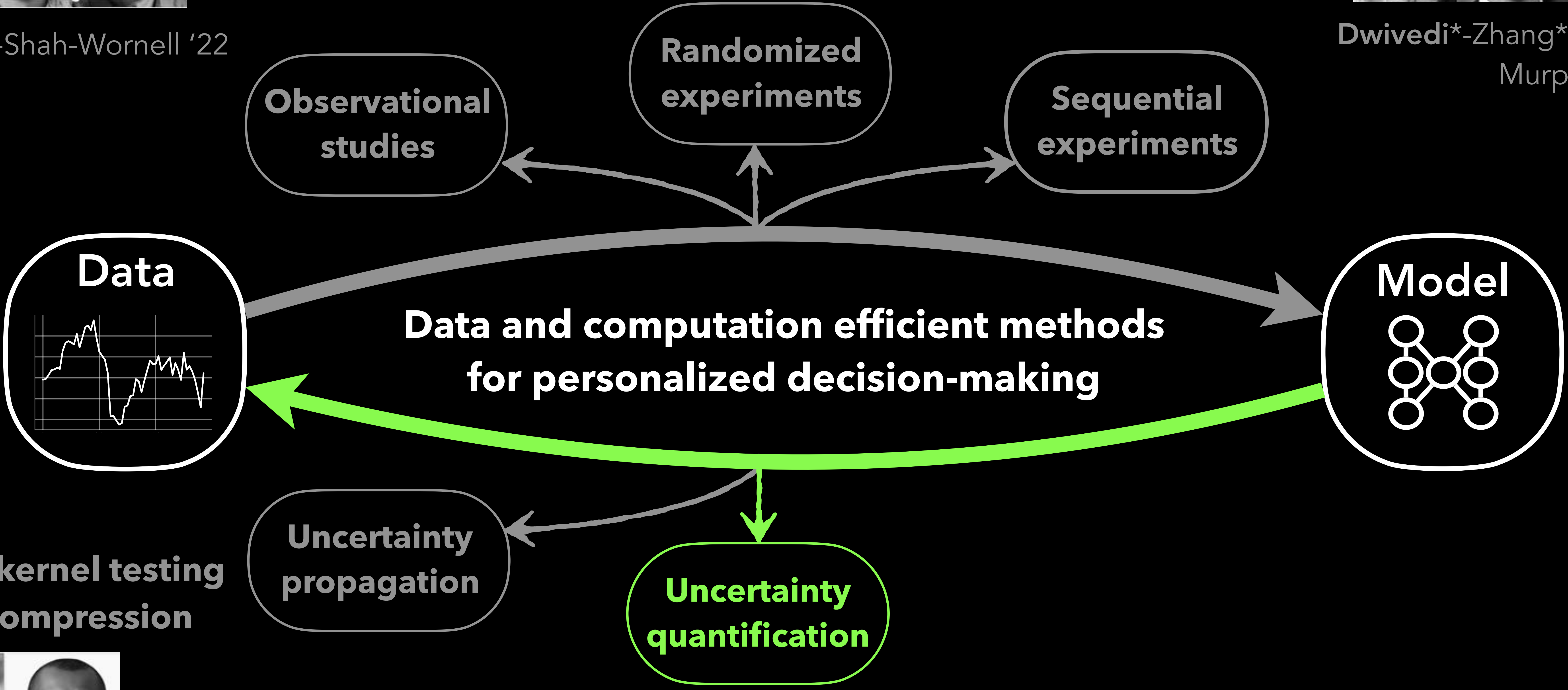

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Mixing time guarantees for MCMC algorithms in high dimensions

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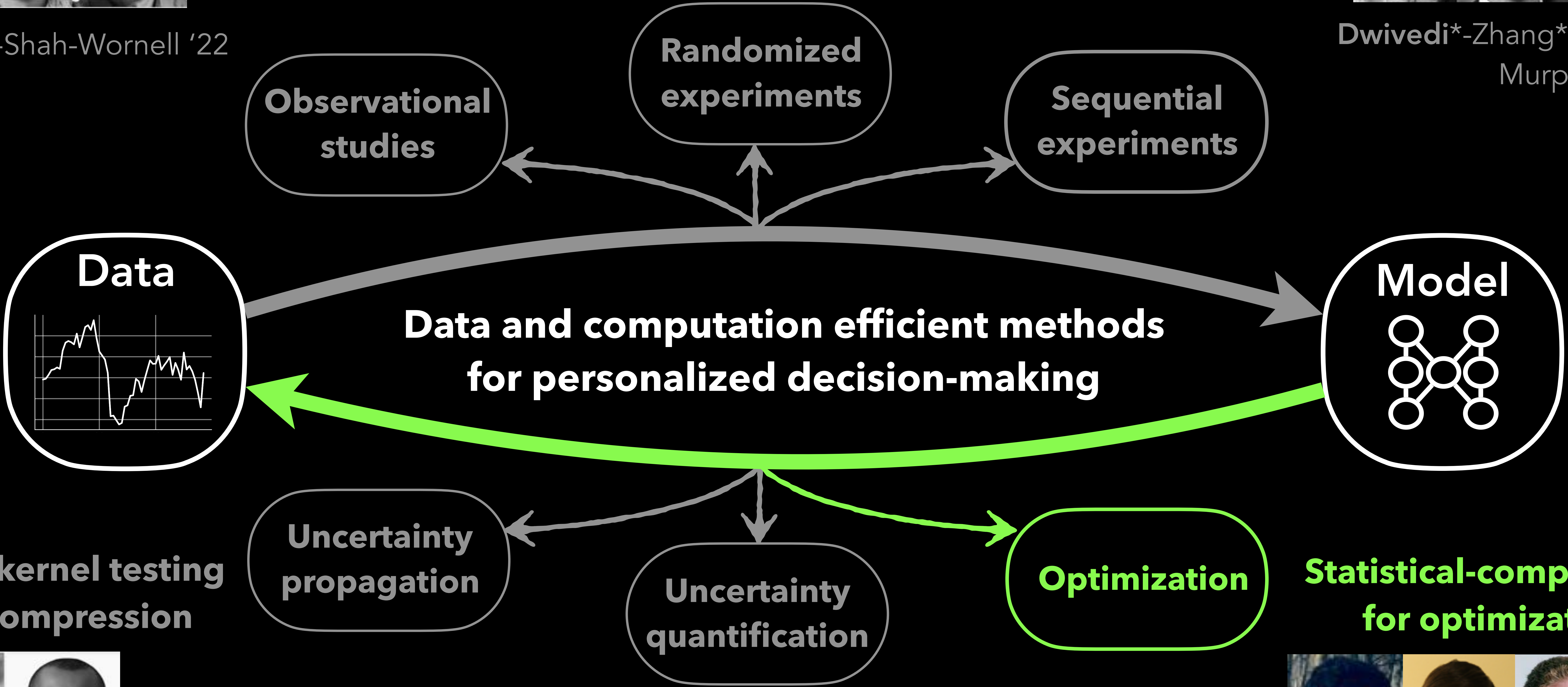

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Statistical-computational tradeoffs for optimization algorithms

Dwivedi*-Ho*-Khamaru*-Wainwright-Jordan-Yu ’19, ’20, ’21, ’22+

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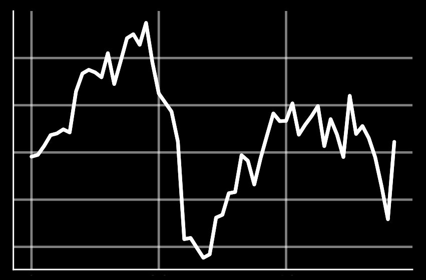

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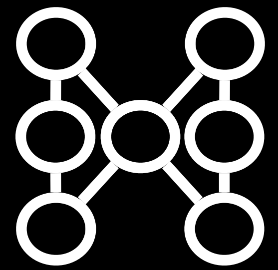

Uncertainty propagation

Uncertainty quantification

Optimization

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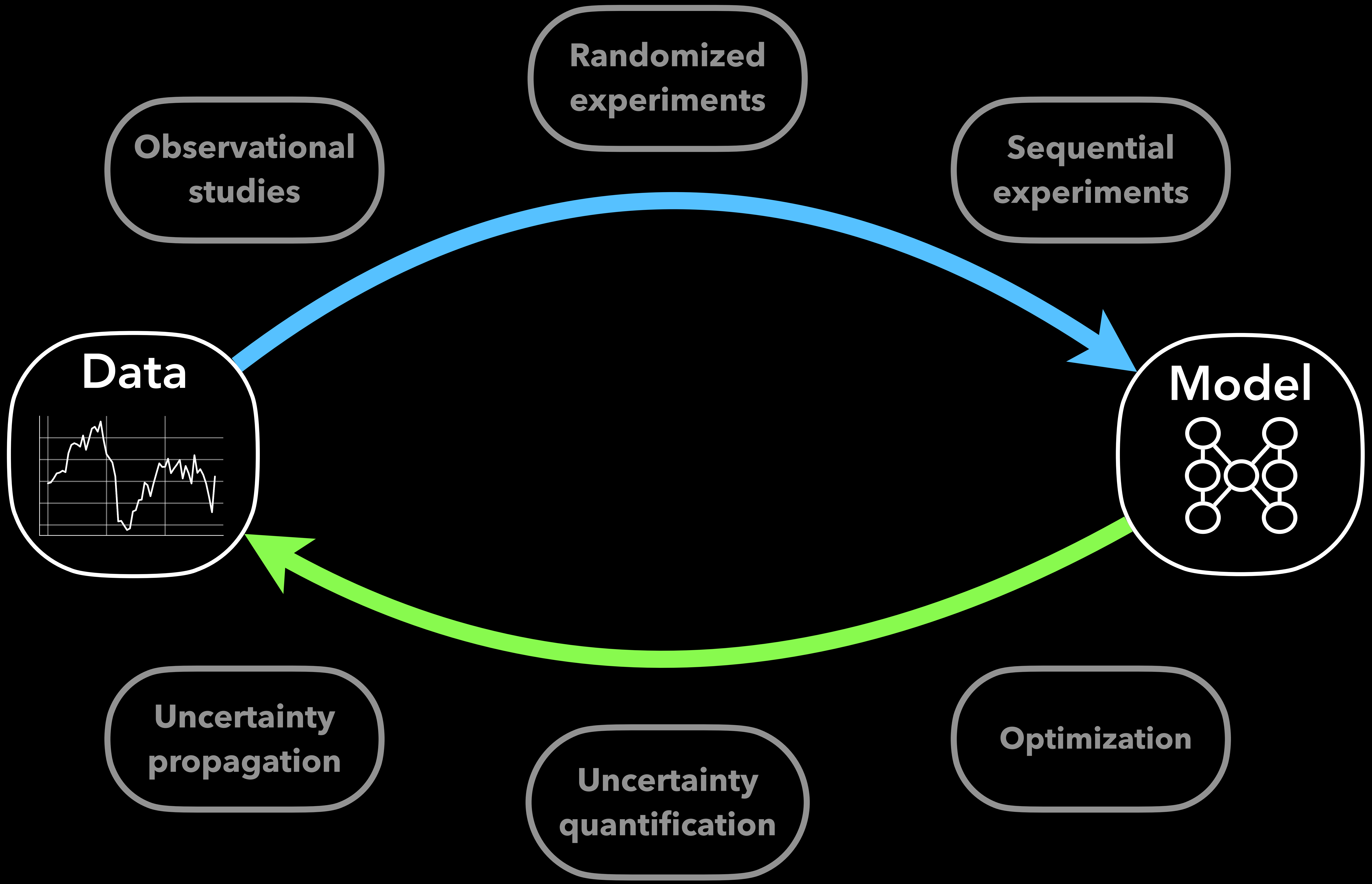
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Model
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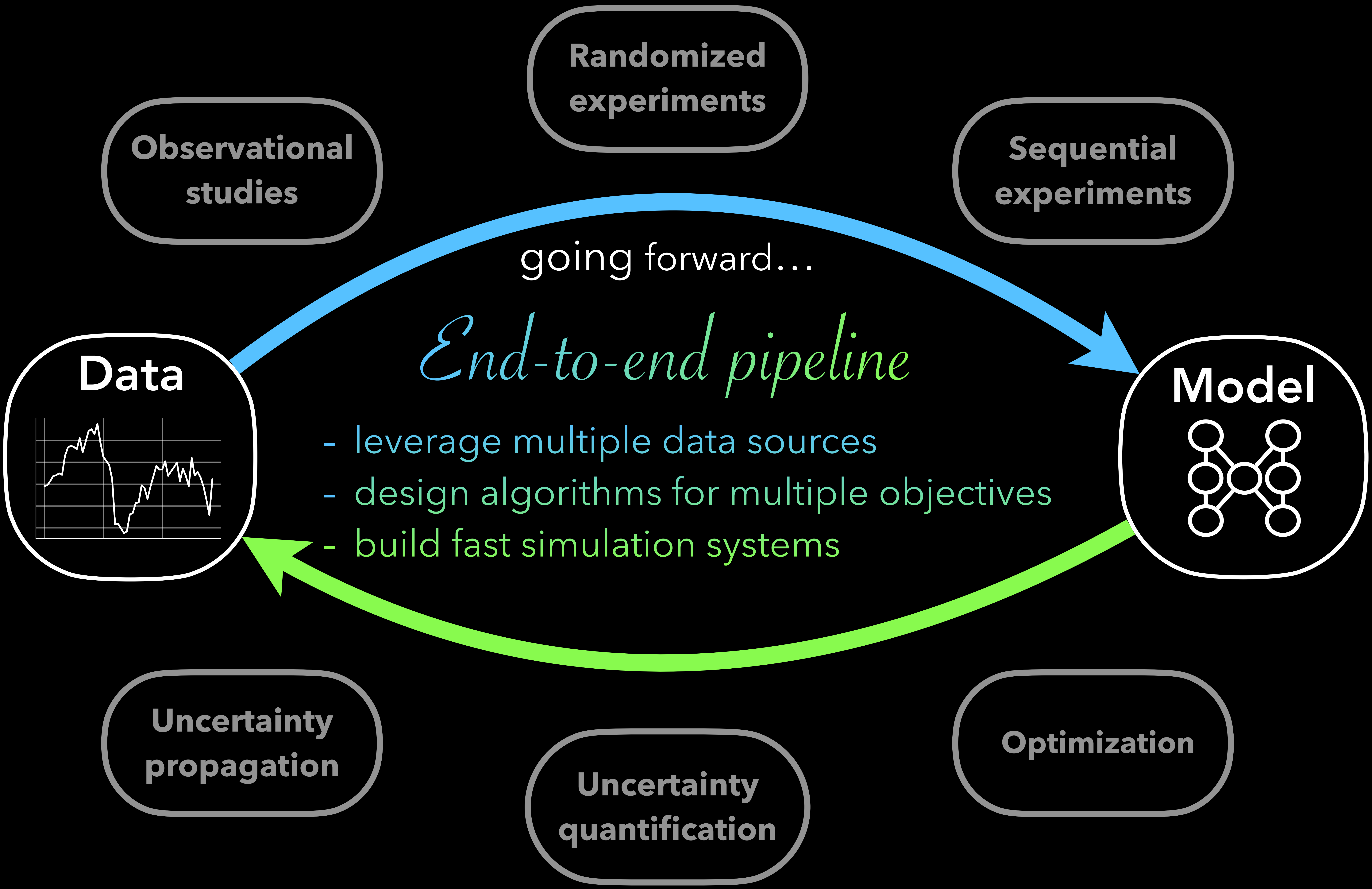

Thank you!

raazdwivedi.github.io