

Advanced Cardiovascular Risk Detection For the Critical Decades

OCTOBER 30, 2025

Bayes through the Life Course
Sarah Urbut, MD PhD



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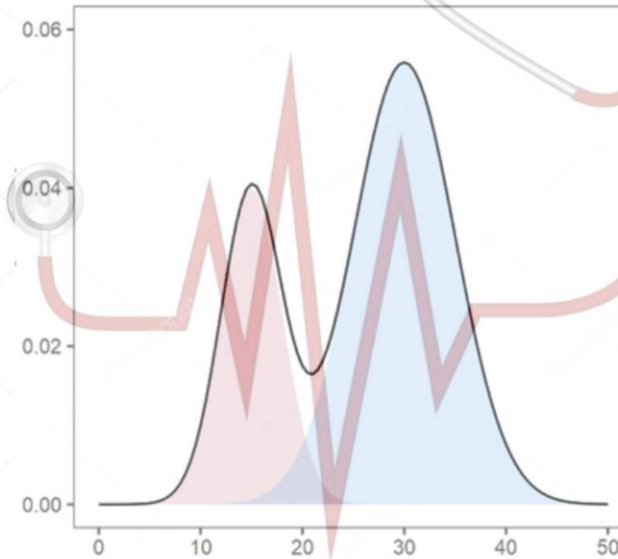
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Disclosures

No disclosures to report

Conflicts of Interest



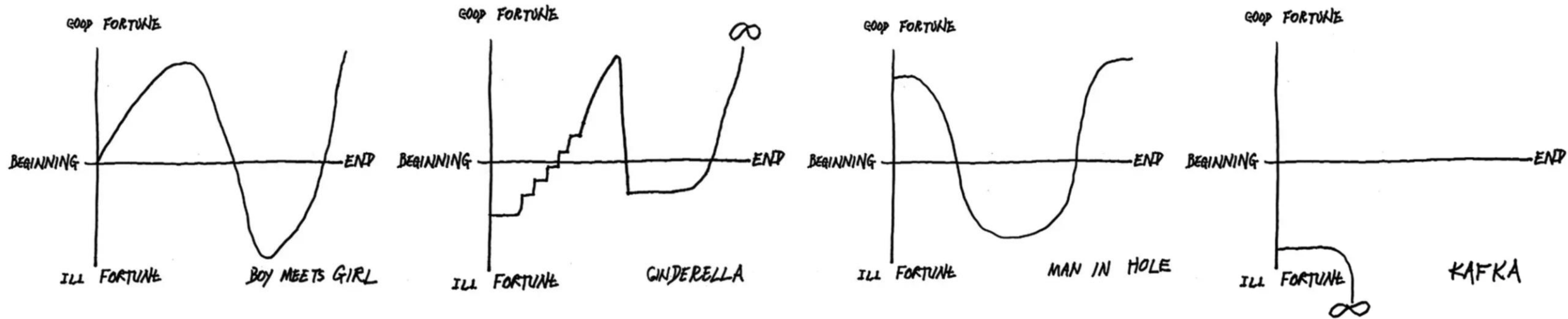
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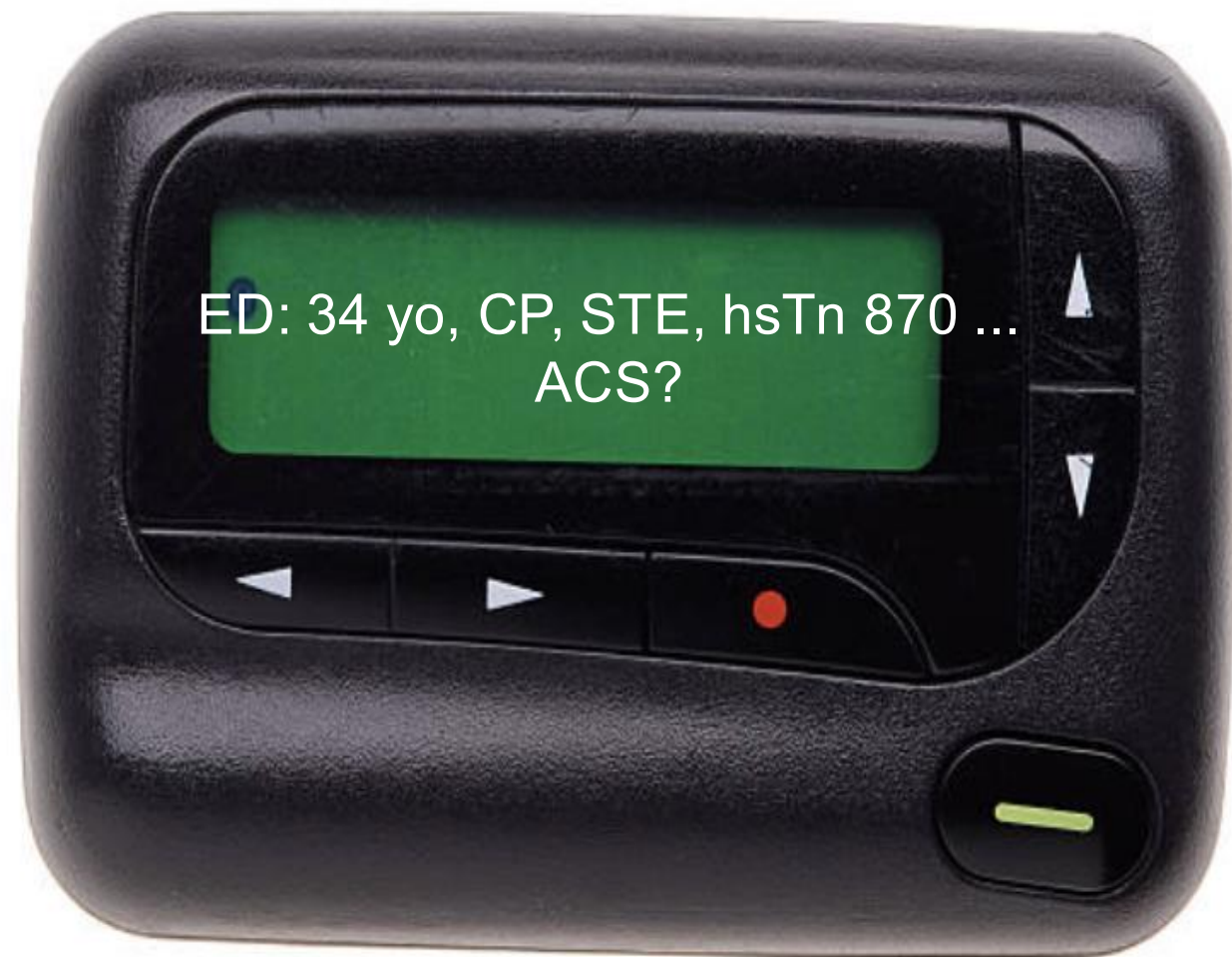
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Shape of a story



Vonnegut, 1955

Everything is moving



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EDITORIAL COMMENT

Polygenic Risk Scoring for Coronary Heart Disease

The First Risk Factor*

Pradeep Natarajan, MD, MMSc

Absolute risk assessment for coronary heart disease (CHD) based on a composite of risk factors is the foundation of contemporary CHD prevention (1). Risk scores serve: 1) to identify individuals at greater risk of CHD over a given time frame; and 2) to establish candidacy for pharmacological preventive strategies. In this issue of the *Journal*, Inouye et al. (2) describe a framework of using polygenic risk scoring to complement clinical risk scoring to identify both high- and low-risk individuals.

SEE PAGE 1883

A HISTORICAL PERSPECTIVE OF

(LDL) cholesterol lowering among individuals with multiple CHD risk factors (5).

In the 1990s, the Framingham risk score, incorporating multiple risk categories to predict the onset of CHD within 10 years, was incorporated into the ATP-III (6). Using largely the same risk categories, the Pooled Cohort Equations incorporated additional cohorts and non-European Americans to develop a 10-year risk estimator for atherosclerotic cardiovascular disease. The Pooled Cohort Equations was adopted by the 2013 American College of Cardiology/American Heart Association joint cholesterol guidelines and is widely used in practice (1).

However, among younger individuals, the ability

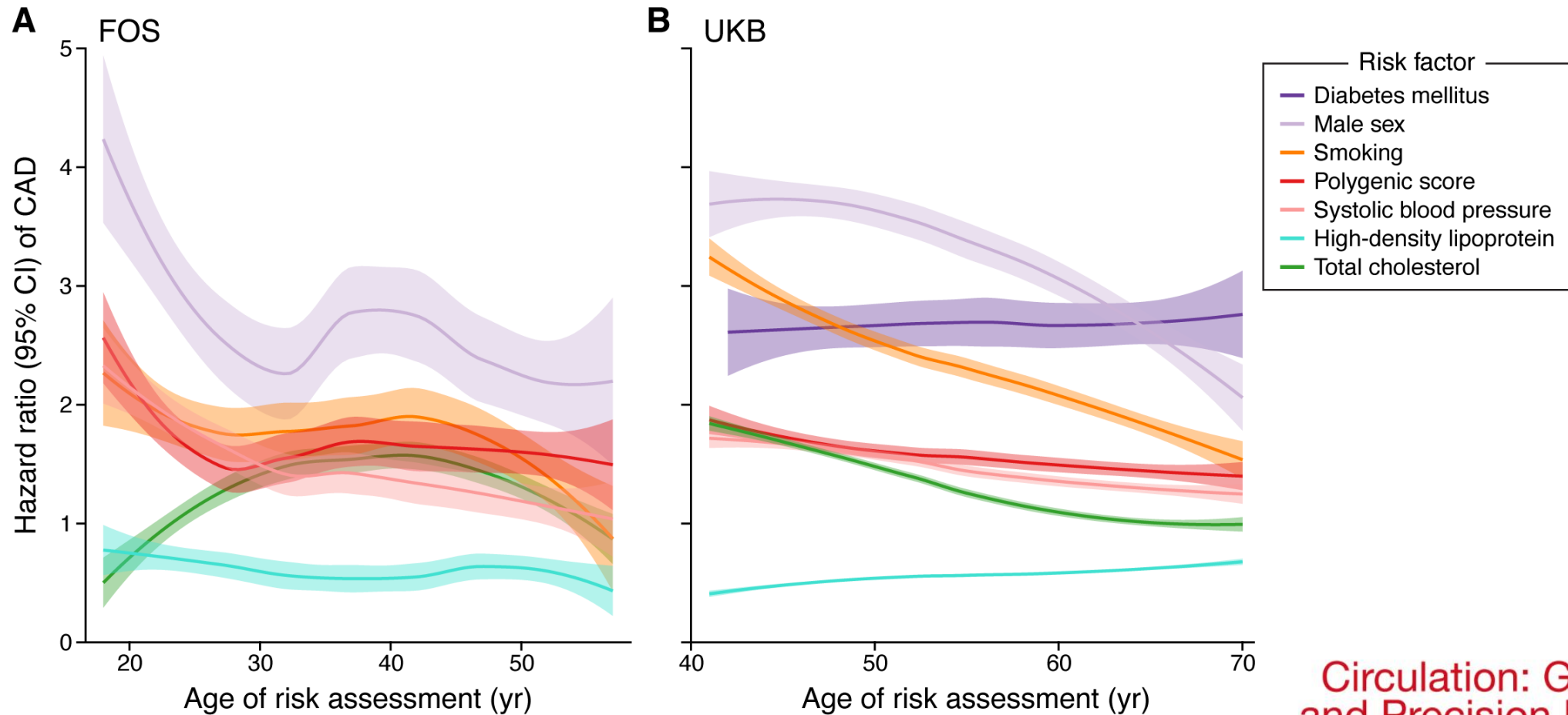
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Dynamic hazard – different roles, same end



**Circulation: Genomic
and Precision Medicine**

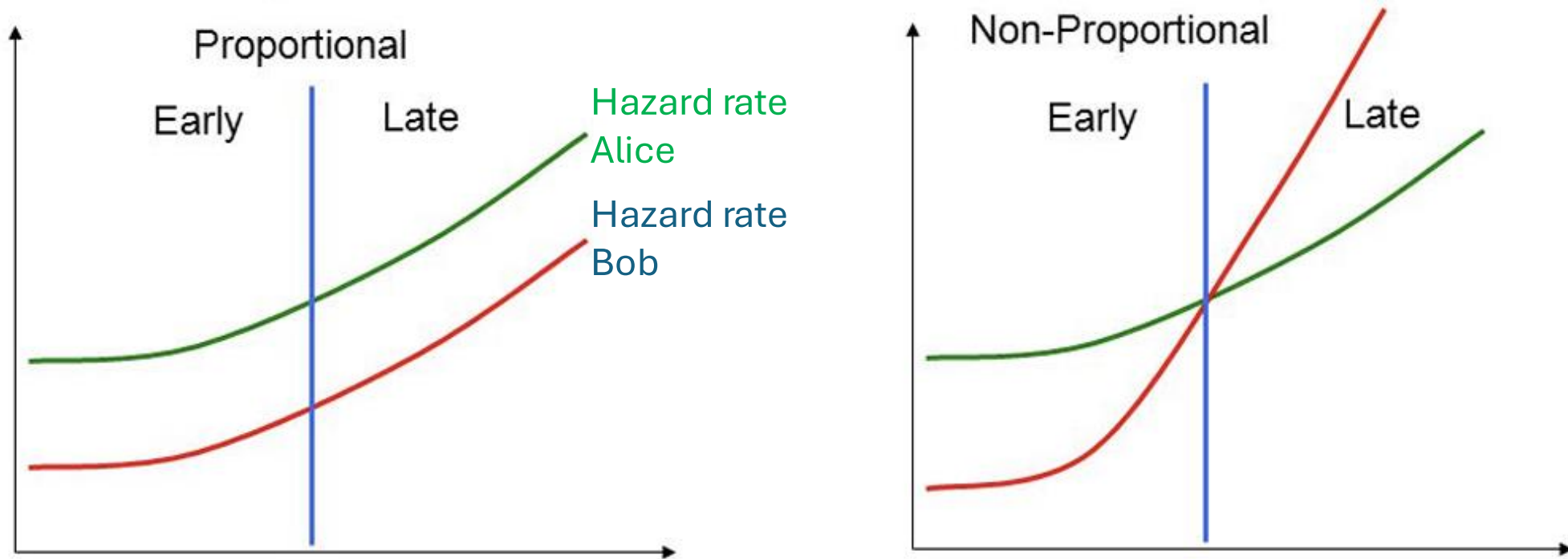
ORIGINAL ARTICLES

Lipoprotein(a) Atherosclerotic Cardiovascular Disease Risk Score Development and Prediction in Primary Prevention From Real-World Data
Random Survival Forest Machine Learning for the Prediction of Cardiovascular Events Among Patients With

ORIGINAL ARTICLES

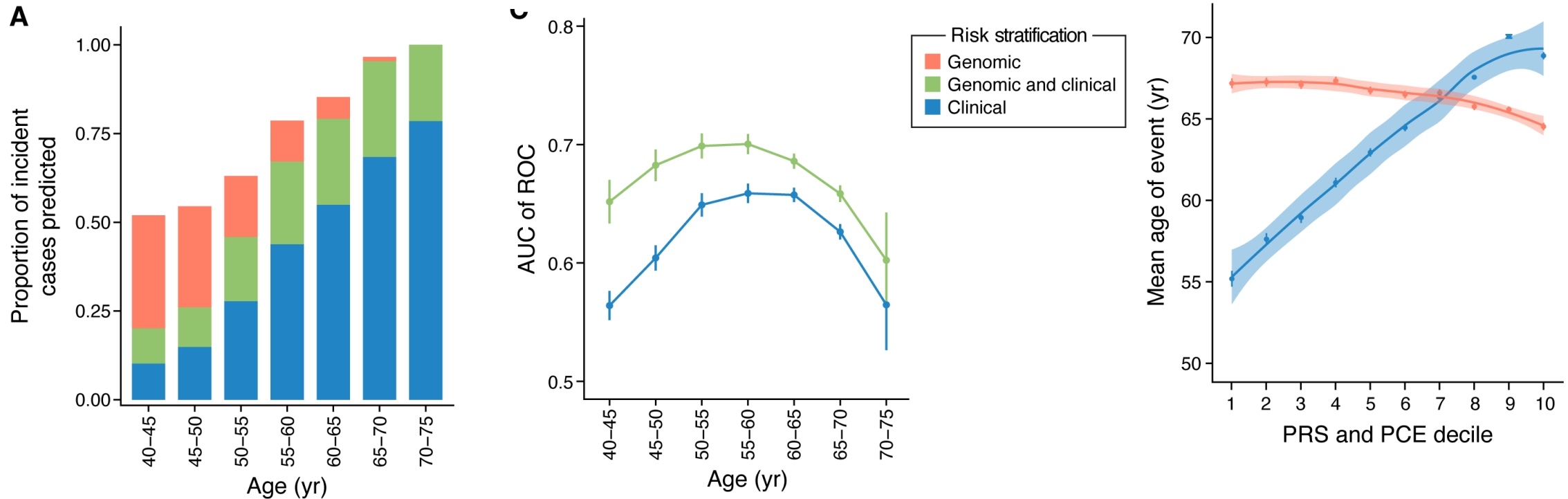
Dynamic Importance of Genomic and Clinical Risk for Coronary Artery Disease Over the Life Course
Sex-Specific Clinical and Genetic Factors Associated With Adverse Outcomes in Hypertrophic Cardiomyopathy

How does lifetime risk depend on ... when the question is asked?



The Cox Proportional Hazards model assumes the hazards are **proportional**: the **relative** hazard ratio remains **constant over time**

Predicts not only early disease, which we sometimes think of, but also all disease early



POWER

DISCRIMINATION

*Urbut et al,
Circulation GPM
2025*

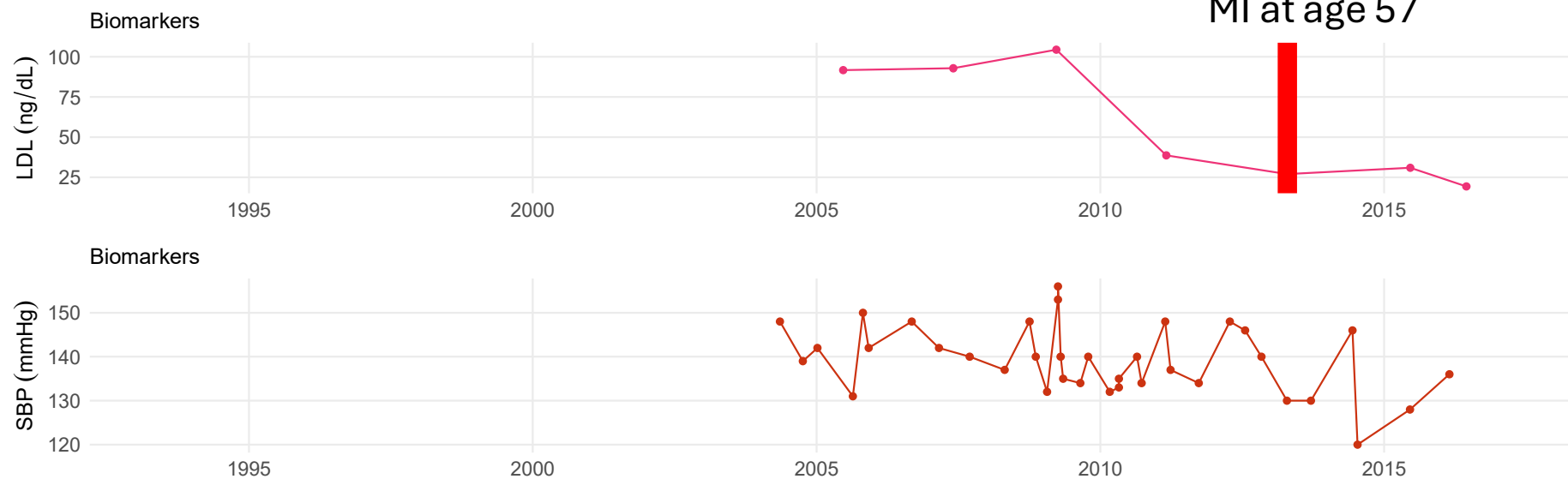
Using the EHR to be predictive, not responsive

Participant summary: ID 1002769

Inferred period of data collection



MI at age 57



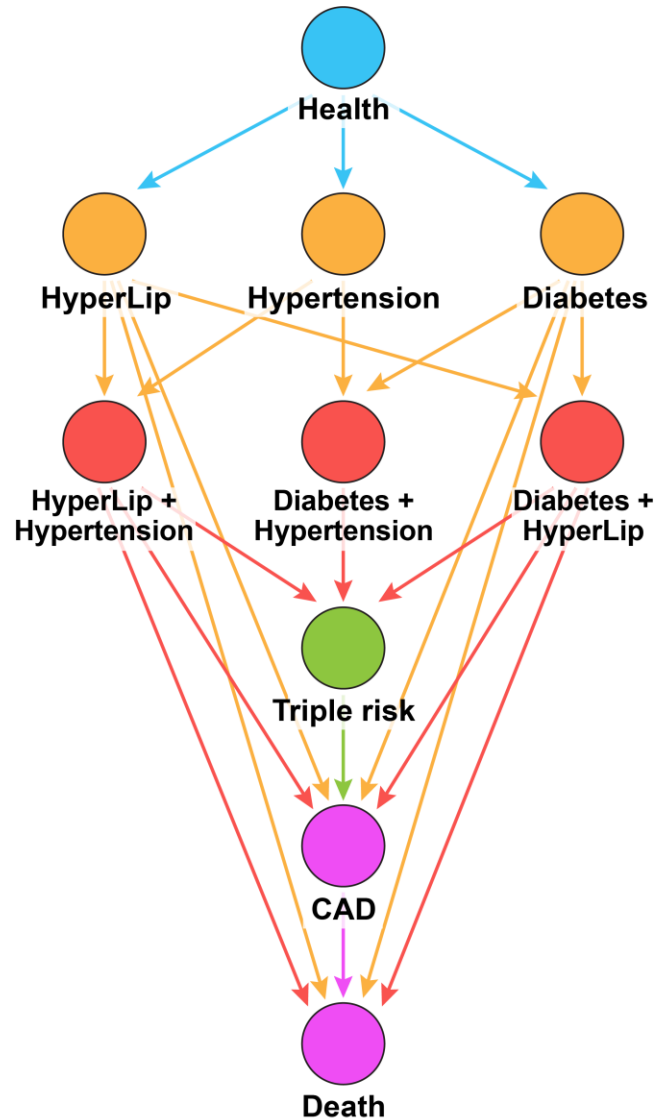
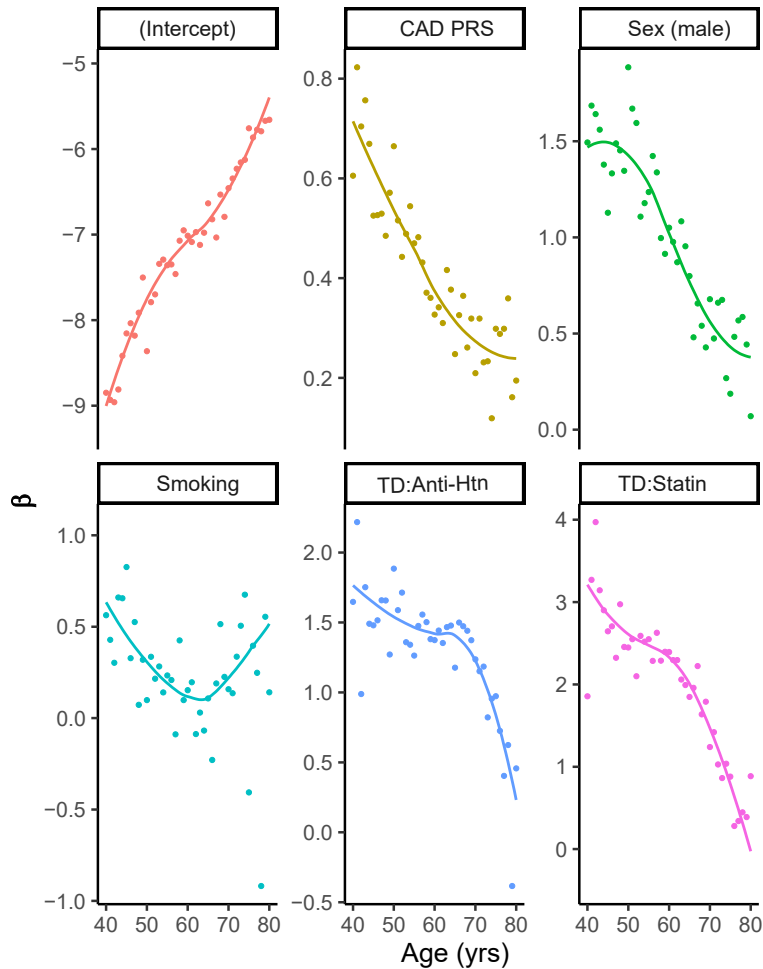
Using the EHR to be predictive, not responsive

Participant summary: ID 1002769

Inferred period of data collection



MSGene: multistate model using genetics for dynamic prediction



- 1 Health
- 2 Single risk factor
- 3 Double risk
- 4 Triple risk
- 5 Absorbing

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Article | [Open access](#) | Published: 07 June 2024

MSGene: a multistate model using genetic risk and the electronic health record applied to lifetime risk of coronary artery disease

Sarah M. Urbut, Ming Wai Yeung, Shaan Khurshid, So Mi Jemma Cho, Art Schuermans, Jakob German, Kogil Taraszka, Kaavya Paruchuri, Aki C. Fahed, Patrick T. Ellinger, Ludovic Trinquart, Giovanni Parmigiani, Alexander Gusev & Pradeep Natarajan

Nature Communications 15, Article number: 4884 (2024) | [Cite this article](#)

Chapter 4: Everyone is a Bayesian

2^n = a numbers problem

Urbut et al, 2025 medRxiv

medRxiv
THE PREPRINT SERVER FOR HEALTH SCIENCES

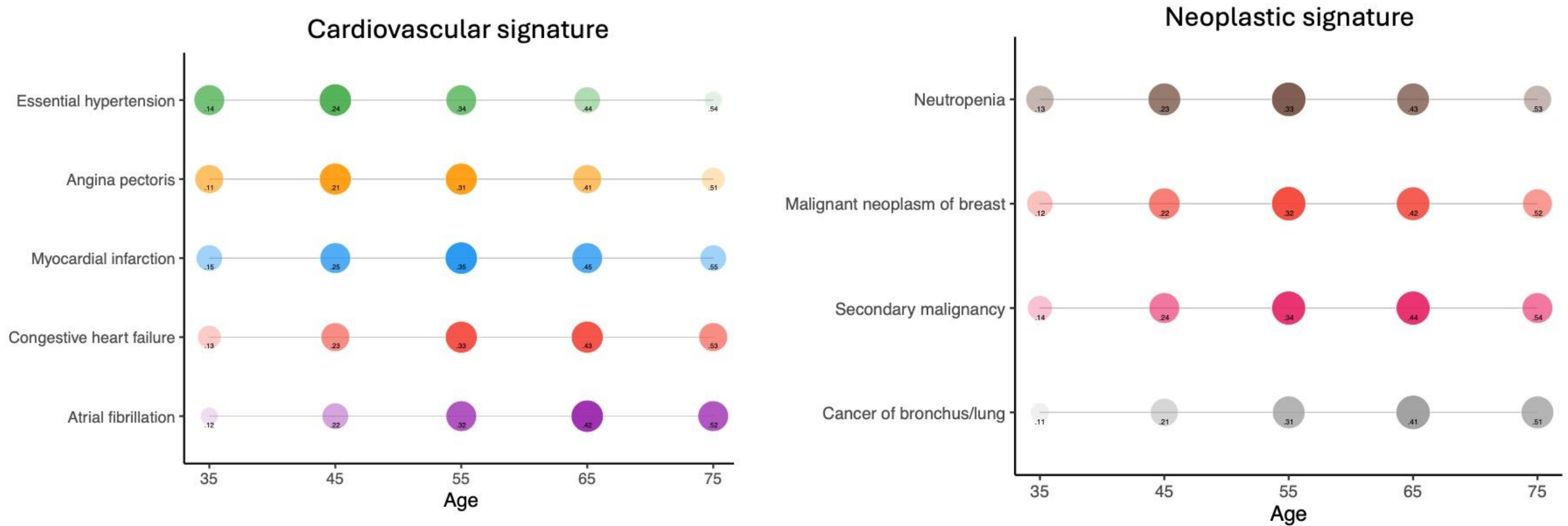
 [Follow this preprint](#)

ALADYNOULLI: A Bayesian approach to disease progression modeling for genomic discovery and clinical prediction

 Sarah M. Urbut,  Yi Ding,  Tetsushi Nakao,  Xilin Jiang, Leslie Gaffney,  Anika Misra, Whitney Hornsby,  Jordan W. Smoller,  Alexander Gusev,  Pradeep Natarajan,  Giovanni Parmigiani
[doi: https://doi.org/10.1101/2024.09.29.24314557](https://doi.org/10.1101/2024.09.29.24314557)



Latent patterns of disease . . . within a signature



Signatures: patterns of disease co-occurrence that vary in time

Time varying trajectories ...

But when do you ask the question?

Patient A: Metabolic → Cancer

Classic metabolic syndrome evolving into malignancy

| Condition | Age 35 | Age 40 | Age 45 | Age 50 | Age 55 | Age 60 | Age 65 |
|-----------------|--------|--------|--------|--------|--------|--------|--------|
| Hypertension | 0 | 1 | 1 | 1 | 1 | 1 | 1 |
| Type 2 Diabetes | 0 | 0 | 1 | 1 | 1 | 1 | 1 |
| CAD | 0 | 0 | 0 | 1 | 1 | 1 | 1 |
| Colon Polyps | 0 | 0 | 0 | 0 | 1 | 1 | 1 |
| Colon Cancer | 0 | 0 | 0 | 0 | 0 | 1 | 1 |
| Metastasis | 0 | 0 | 0 | 0 | 0 | 0 | 1 |

Patient B: Inflammatory → CVD → Neuro

Inflammatory disease followed by cardiovascular complications and neurological issues

| Condition | Age 35 | Age 40 | Age 45 | Age 50 | Age 55 | Age 60 | Age 65 |
|----------------------|--------|--------|--------|--------|--------|--------|--------|
| Rheumatoid Arthritis | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| IBD | 0 | 1 | 1 | 1 | 1 | 1 | 1 |
| CAD | 0 | 0 | 0 | 1 | 1 | 1 | 1 |
| Heart Failure | 0 | 0 | 0 | 0 | 1 | 1 | 1 |
| Depression | 0 | 0 | 0 | 0 | 1 | 1 | 1 |
| Cognitive Decline | 0 | 0 | 0 | 0 | 0 | 1 | 1 |

Patient C: Early CVD → GI → Metabolic

Early cardiovascular disease with later digestive and metabolic complications

| Condition | Age 35 | Age 40 | Age 45 | Age 50 | Age 55 | Age 60 | Age 65 |
|-----------------|--------|--------|--------|--------|--------|--------|--------|
| CAD | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| Heart Failure | 0 | 1 | 1 | 1 | 1 | 1 | 1 |
| GERD | 0 | 0 | 1 | 1 | 1 | 1 | 1 |
| IBD | 0 | 0 | 0 | 1 | 1 | 1 | 1 |
| Type 2 Diabetes | 0 | 0 | 0 | 0 | 1 | 1 | 1 |
| Obesity | 0 | 0 | 0 | 0 | 1 | 1 | 1 |

Metabolic CVD Cancer Inflammatory GI Neuro

- *ACC Statement on Inflammation:*
Inflammation is therapeutic target in CVD
- CANTOS trial: IL-1 β inhibition reduces CV events

Urbut et al, 2024

Everyone is Bayesian

Joint consideration: discovery and prediction

$$P(\Pi|Diagnoses) \propto P(Diagnoses|\Pi) p(\Pi)$$

Continuously
updated posteriors

Individual data
likelihood (EHR,
clinical data)

Individual
predilection to
a signature

Population level
signatures



BIG DATA

*Thomas Bayes, History of life
insurance in its formative years*
American Conservation Co., 1936,
Chicago

This is statistics: Aladynoulli

$$\phi \sim N(\mu_d + \psi_{kd}, K)$$

$$\lambda \sim N(\gamma_k g_i + r_i, K)$$

Hazard for individual i of disease d at time t :

$$\pi_{idt} = \sum_k \underbrace{f(\lambda_{idt})}_{\text{Individual}} \underbrace{f(\phi_{idt})}_{\text{Population}}$$

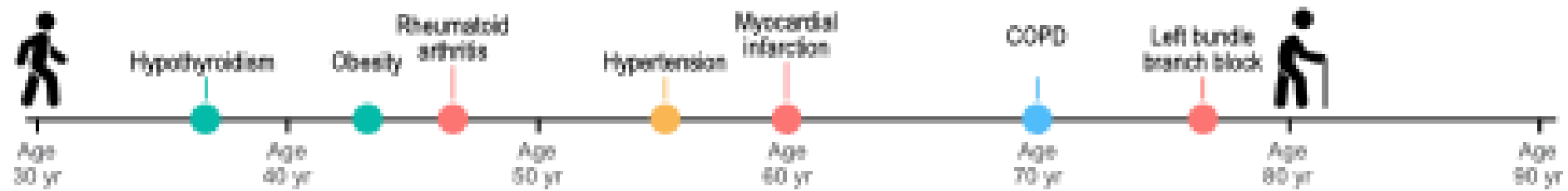
We are all Bayesians

$$l_{id} = \underbrace{\sum_{t < E_{id}} \log(1 - \pi_{idt})}_{\text{At risk}} + \underbrace{Y_{idt} \pi_{idt}}_{\text{Event}} + \underbrace{(1 - Y_{idE_{id}})(1 - \pi_{idt})}_{\text{Censored}}$$

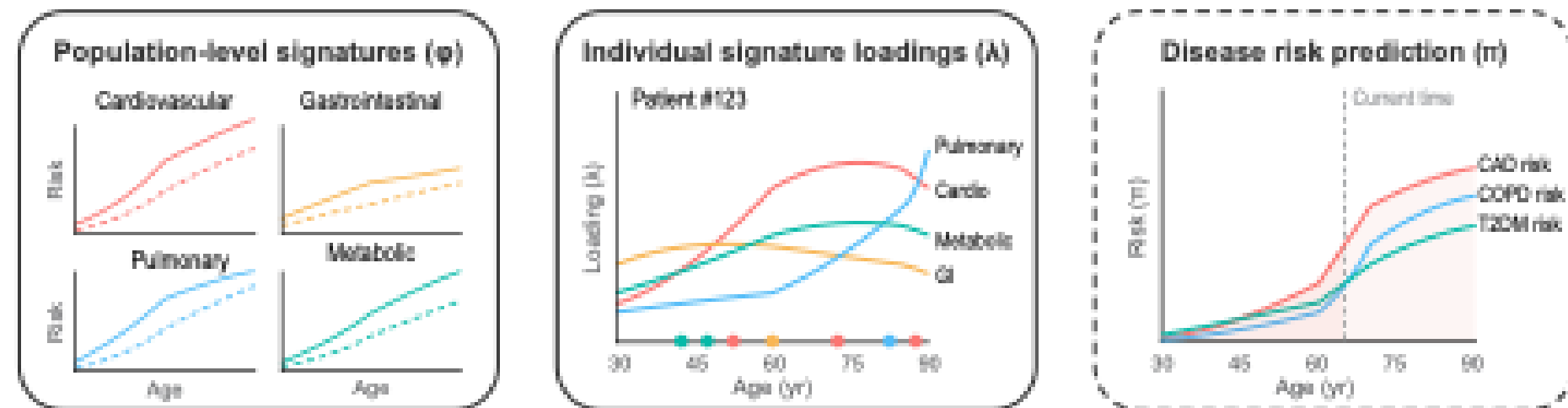
$$\text{Post} \propto \textit{Likelihood} \times \textit{Prior}$$

$$P(\textit{Model} | \textit{Diagnoses}) = P(\textit{Diagnoses} | \textit{Model}) \cdot P(\textit{Model})$$

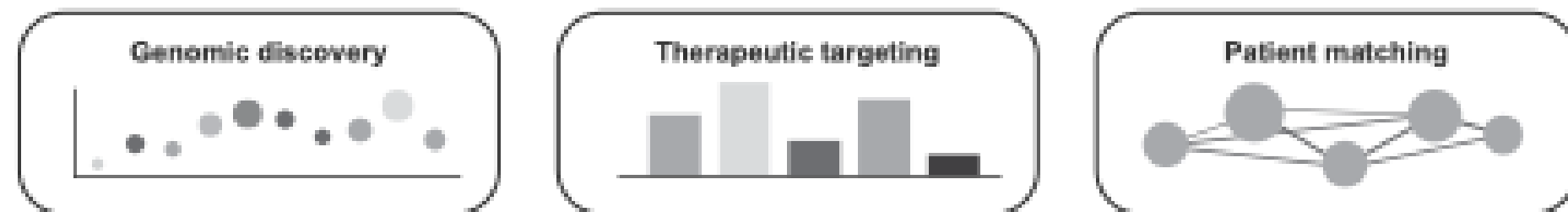
Life journey with diagnoses (patient #123)



Aladynou model components



Applications



Signatures: Characteristic patterns of incidence and timing

$f(\phi_{idt})$

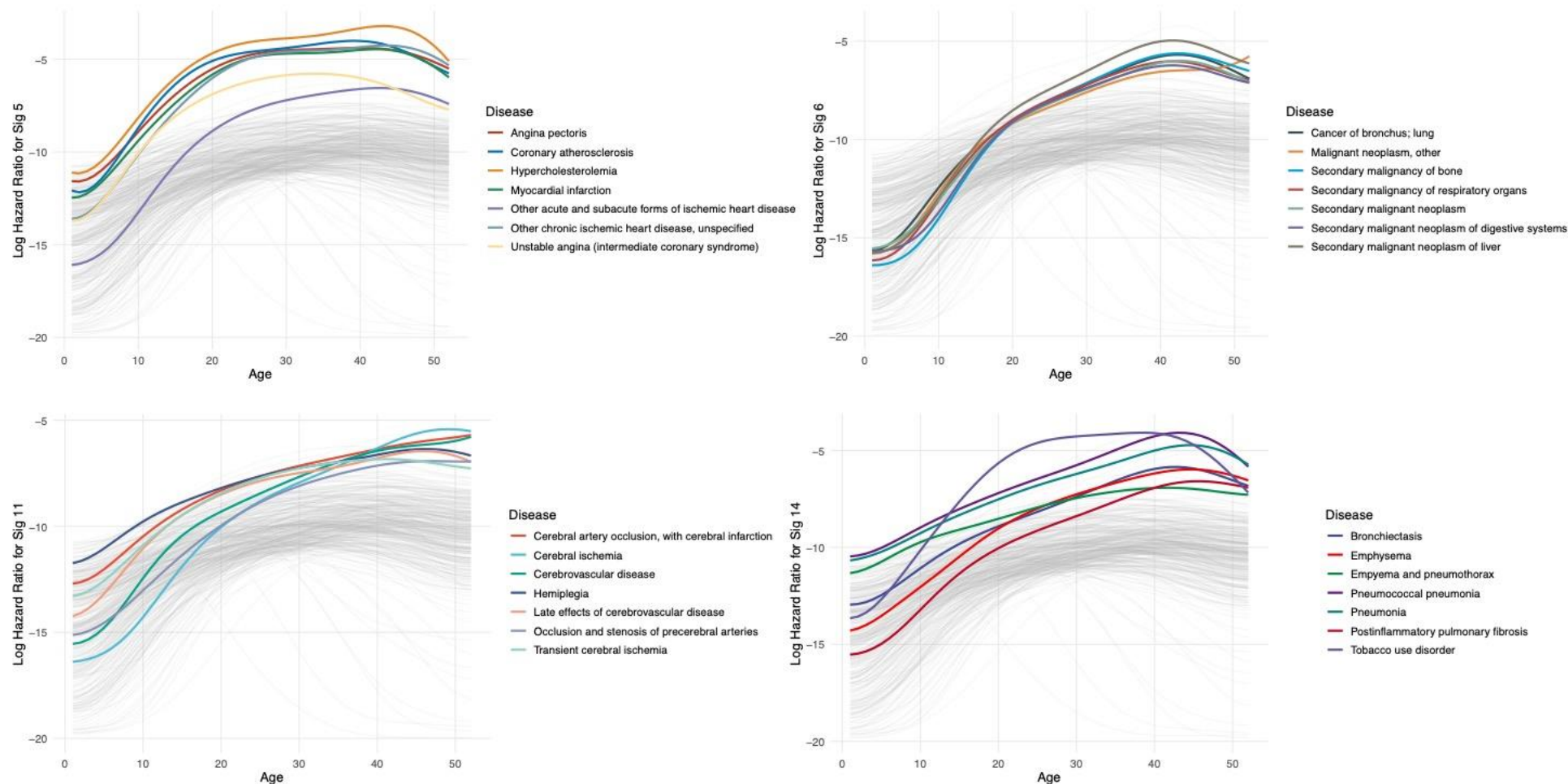
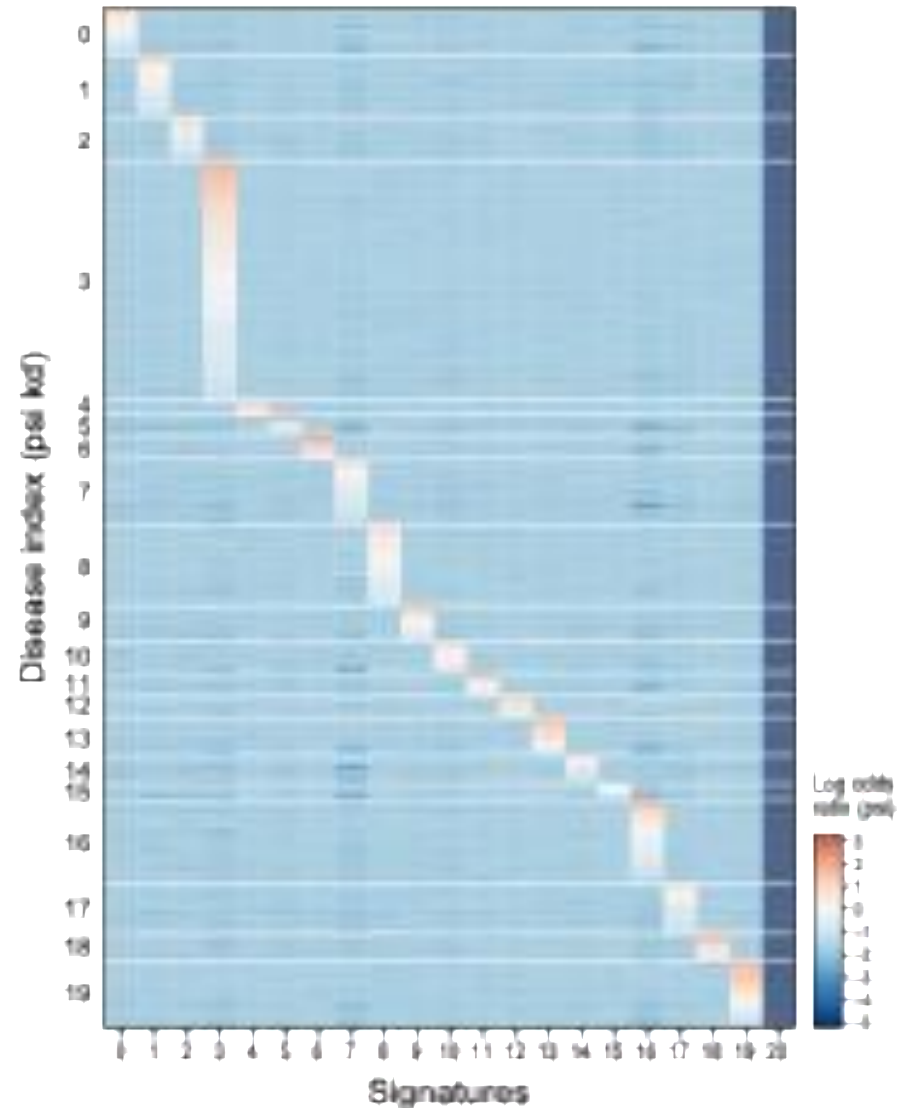
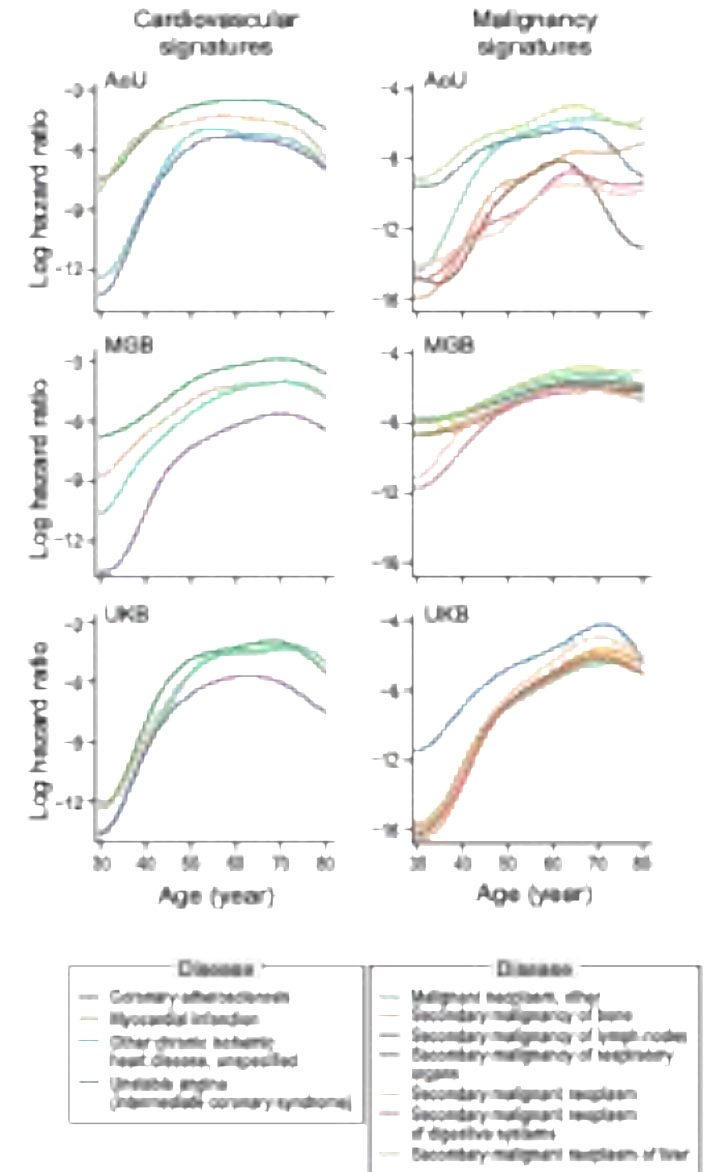
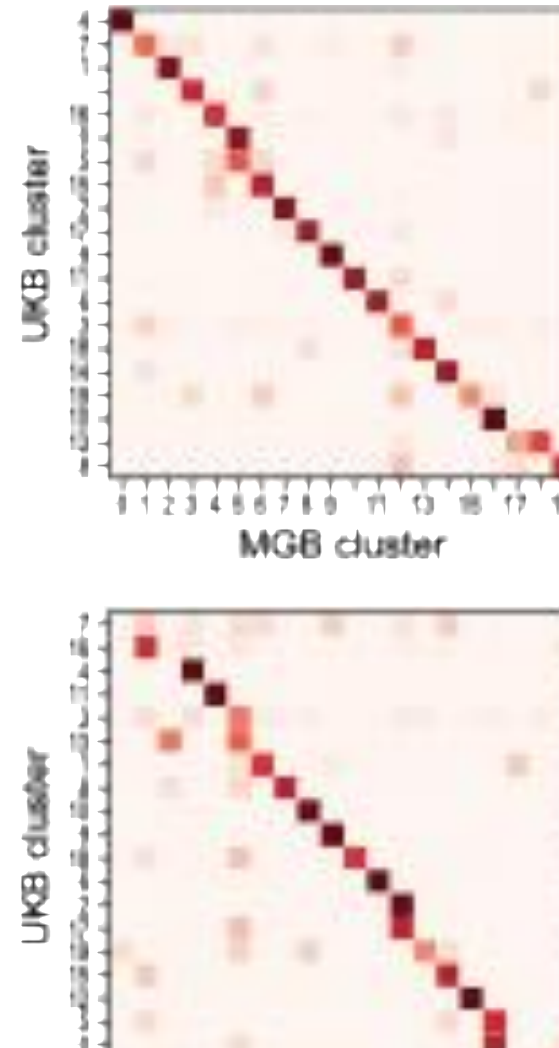


Figure: Temporal evolution of disease probabilities across signatures

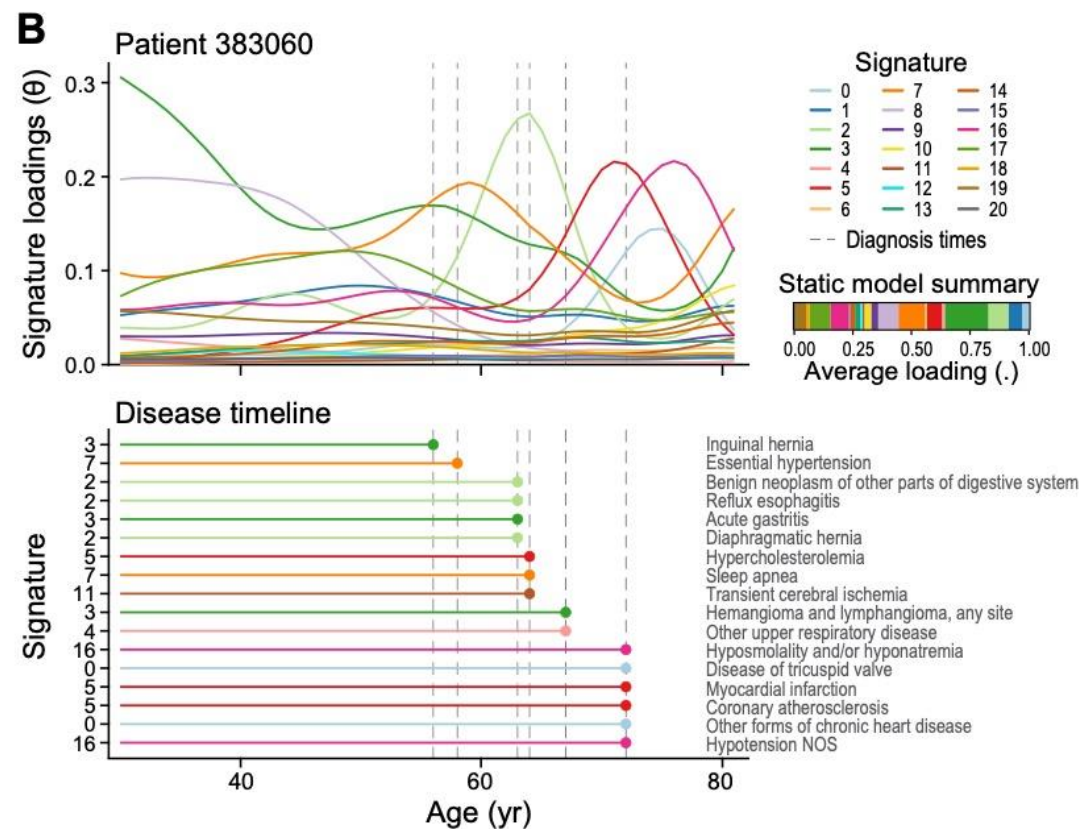
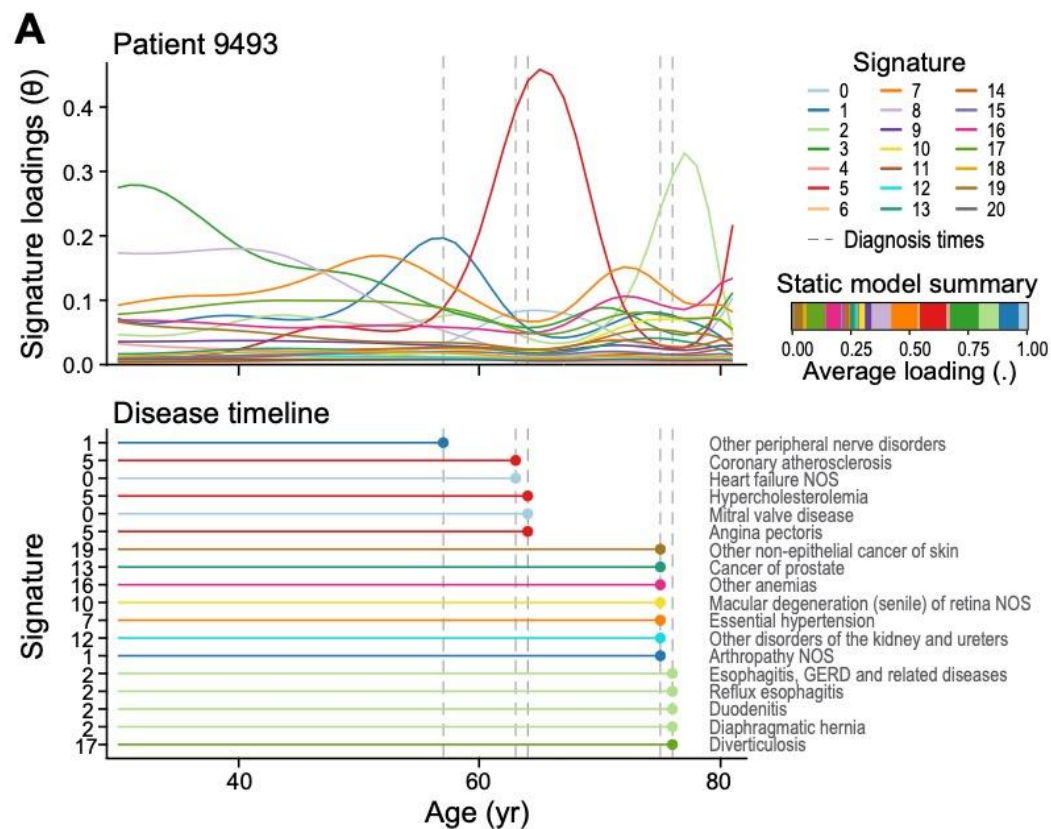
Varying degree of allegiance



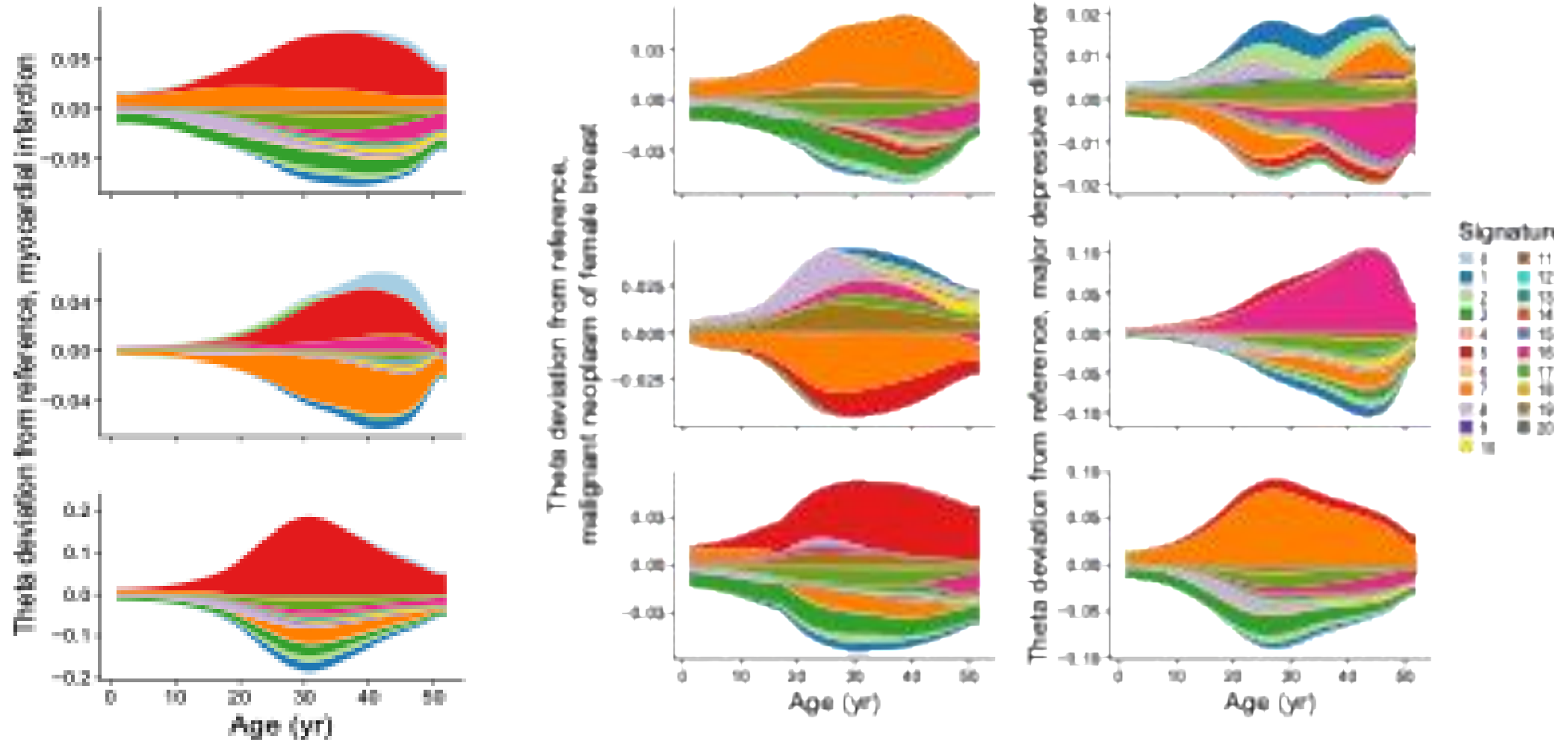
Consistency across biobanks



Walking the time-line... $f(\lambda_{idt})$

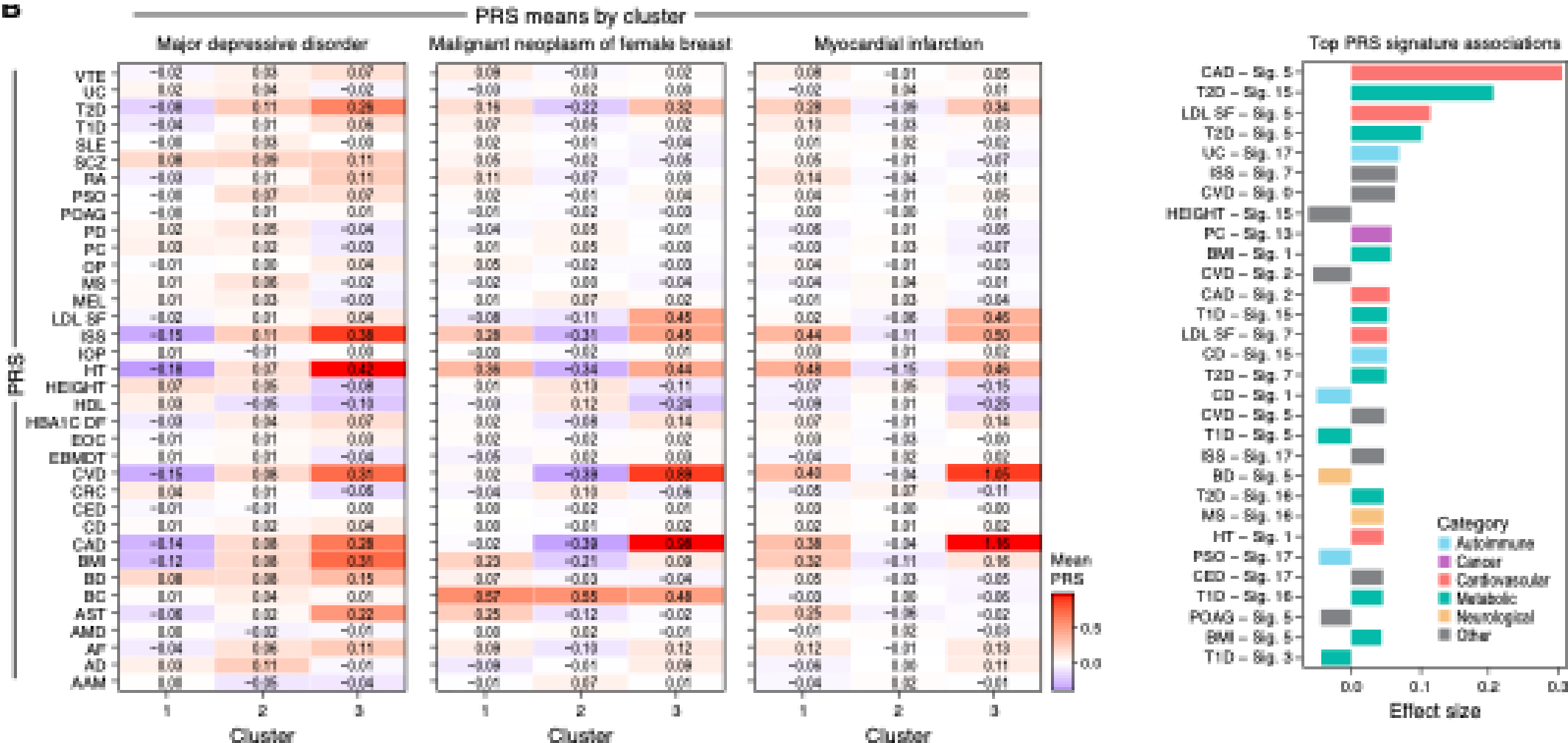


Heterogeneity within disease: Revealing biology



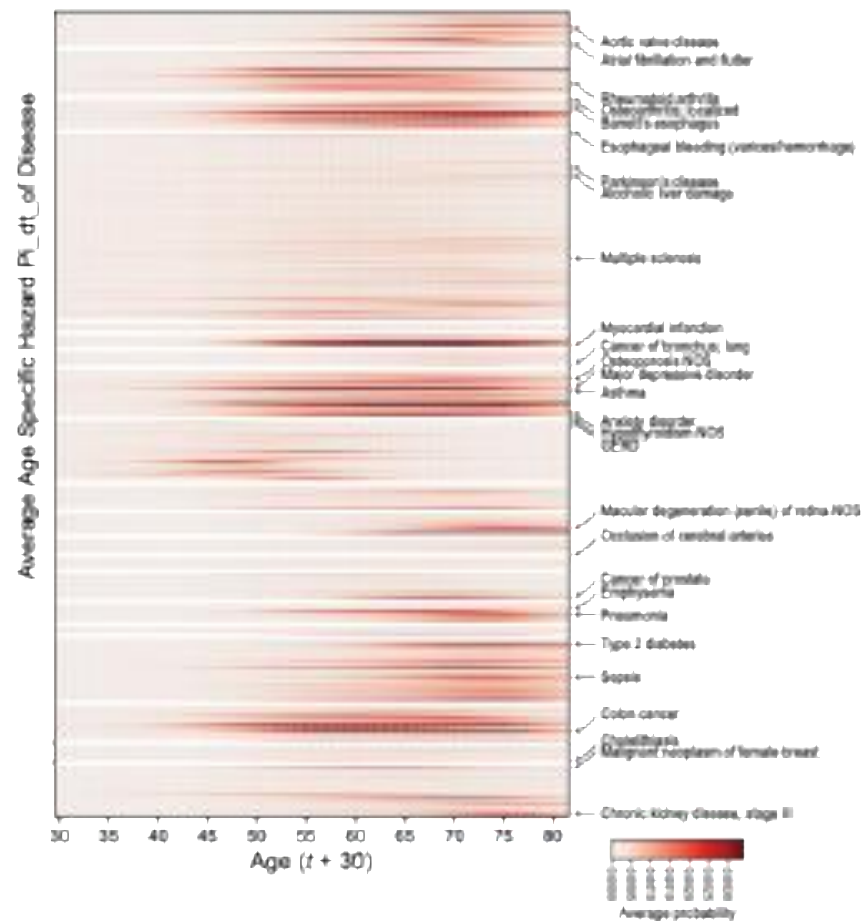
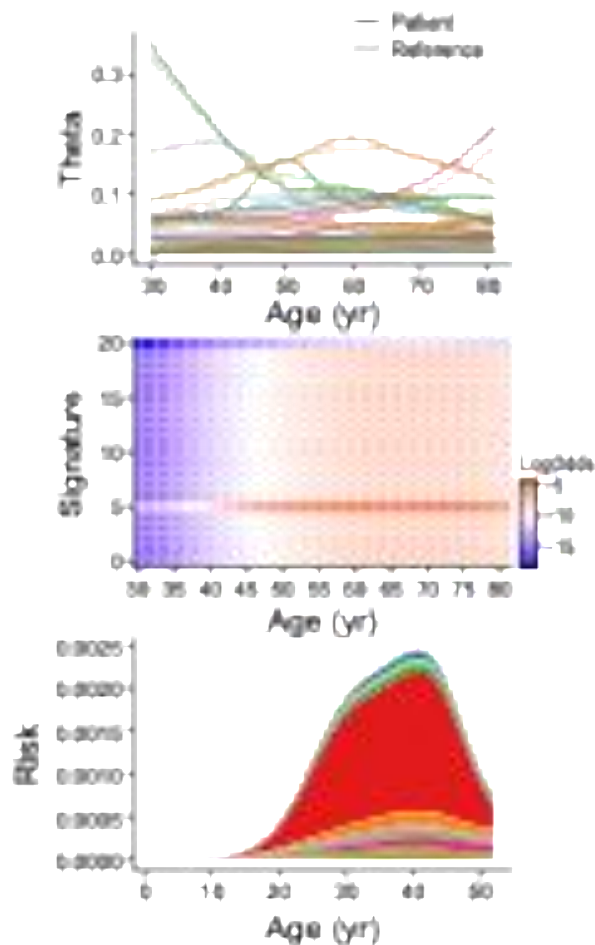
- Ference et al., *NEJM* 2012: Genetic variants that lower cholesterol even moderately over lifetime dramatically reduce CAD
- Khera et al., *Nat Genet* 2018: PRS identifies high-risk individuals missed by traditional screening

Heterogeneity in phenotype matches Genotype

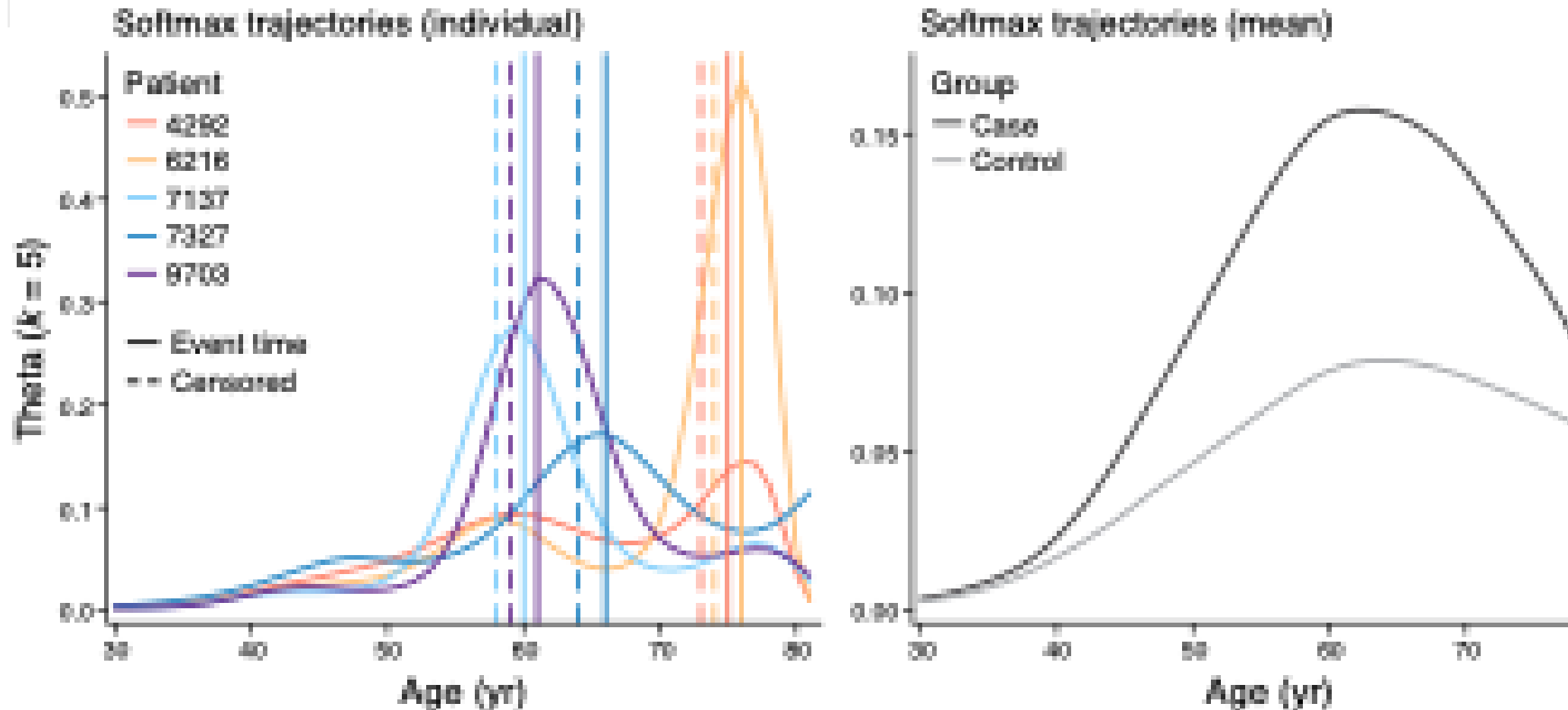


Making predictions is only part ...

$$\pi_{idt} = \sum_k f(\lambda_{idt}) f(\phi_{idt})$$

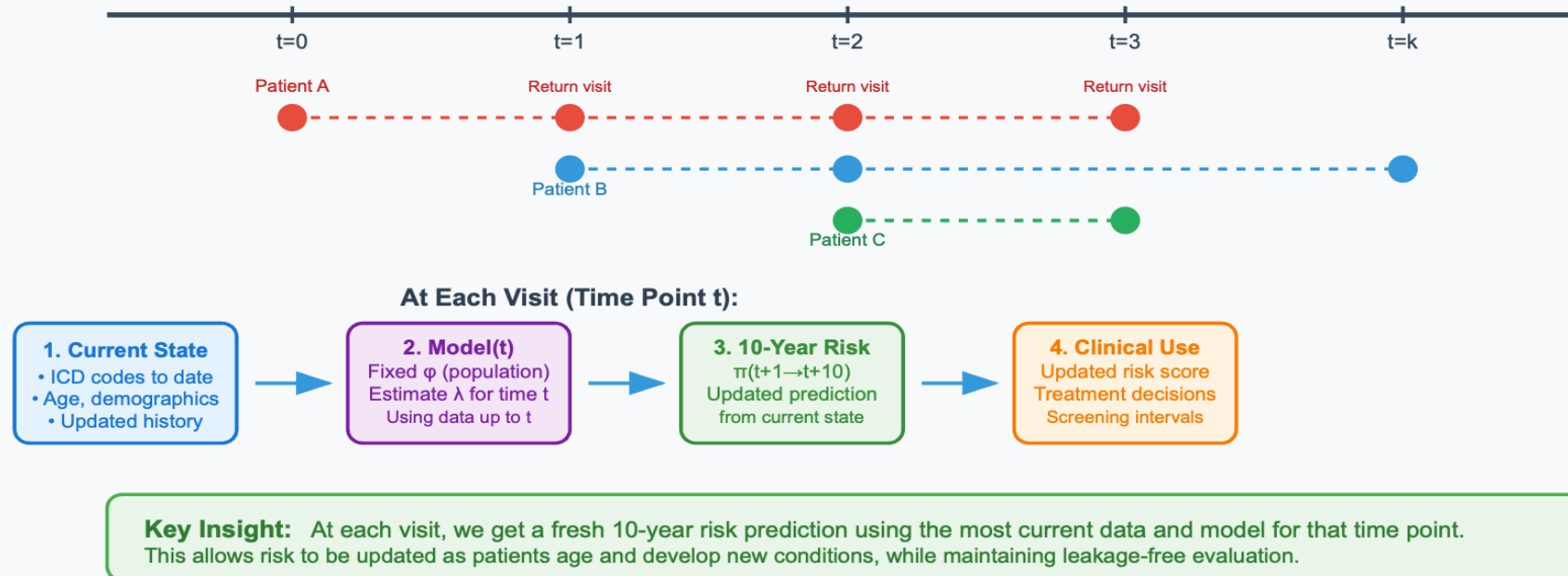


Aladynoulli: the genie works with a blindfold!

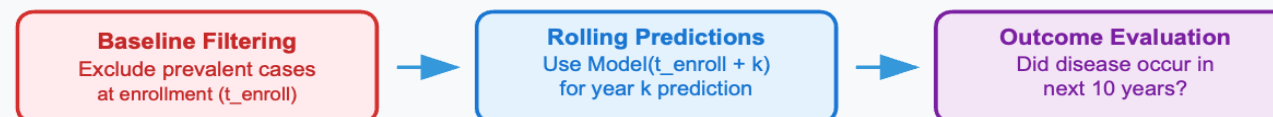


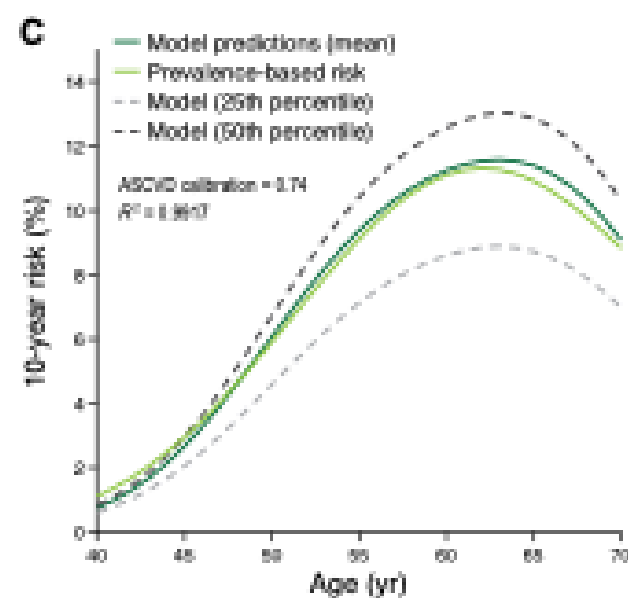
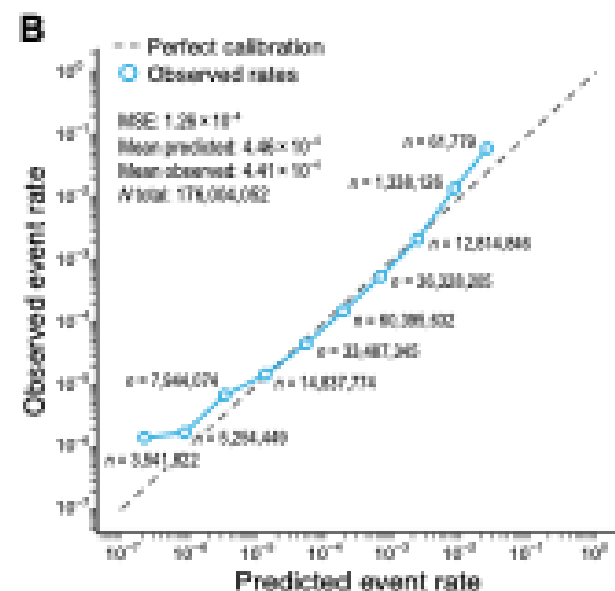
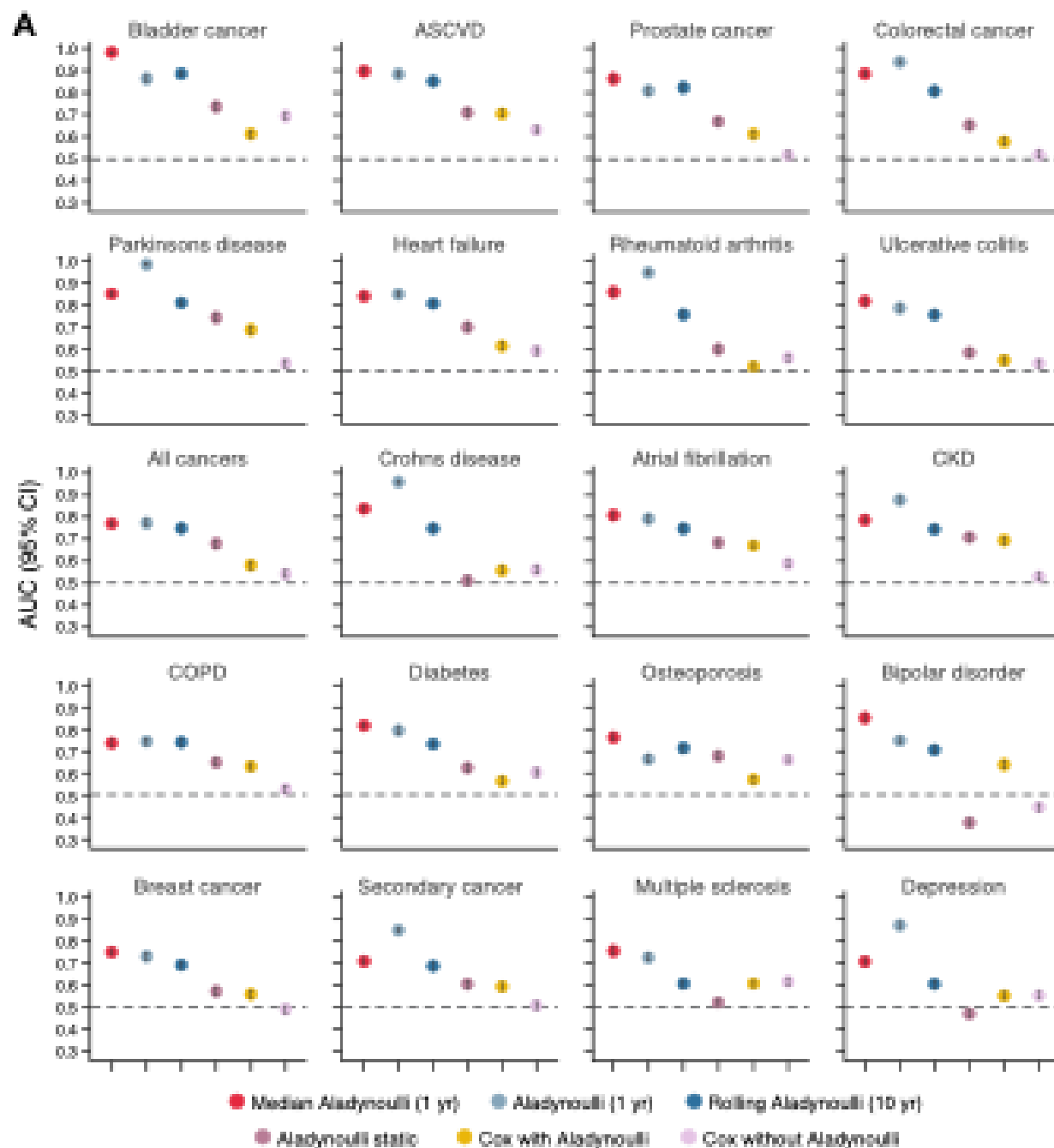
Performance assessment: Changing the paradigm

Dynamic Risk Prediction with Repeated Evaluations



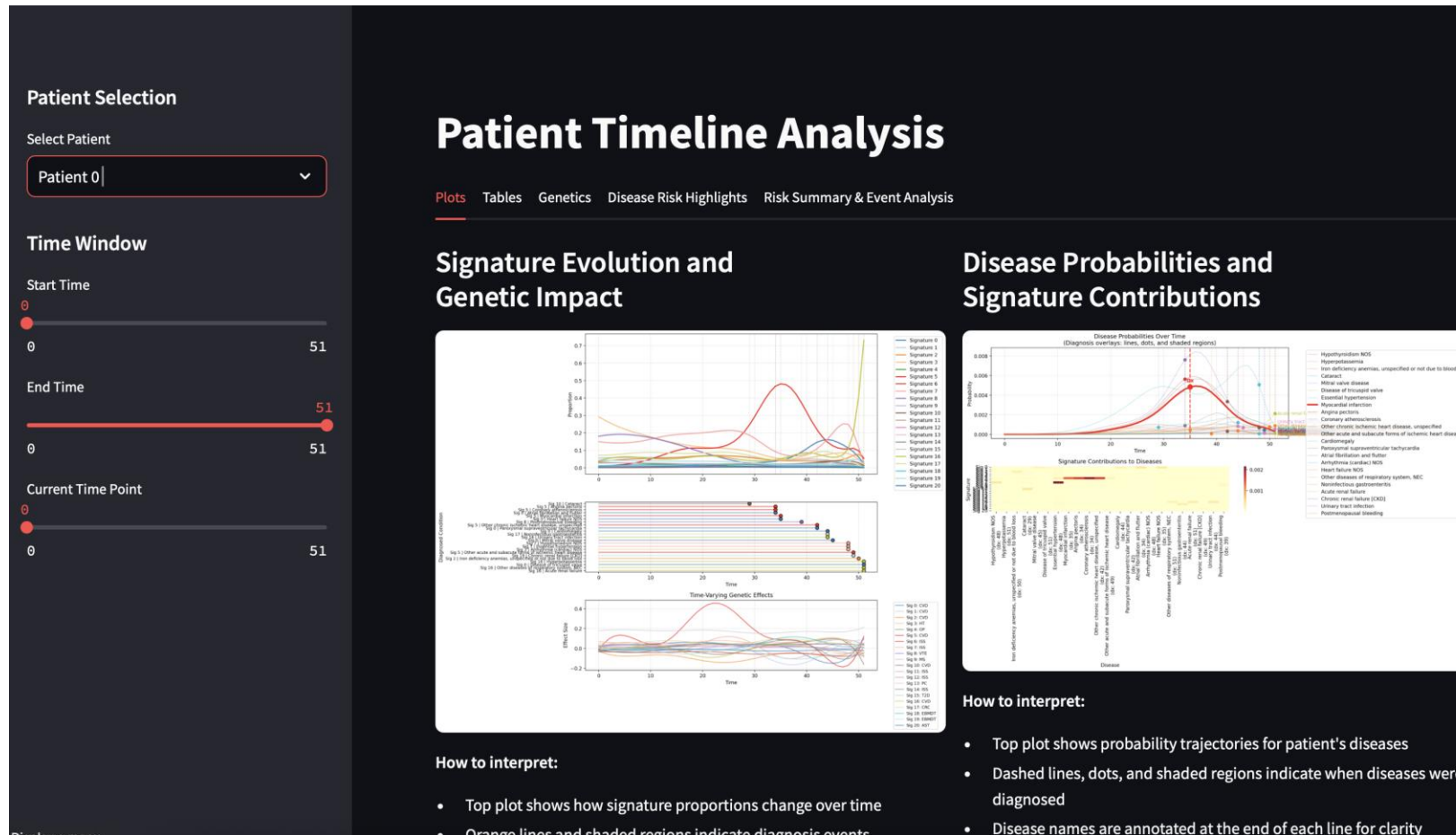
AUC Evaluation Methodology:

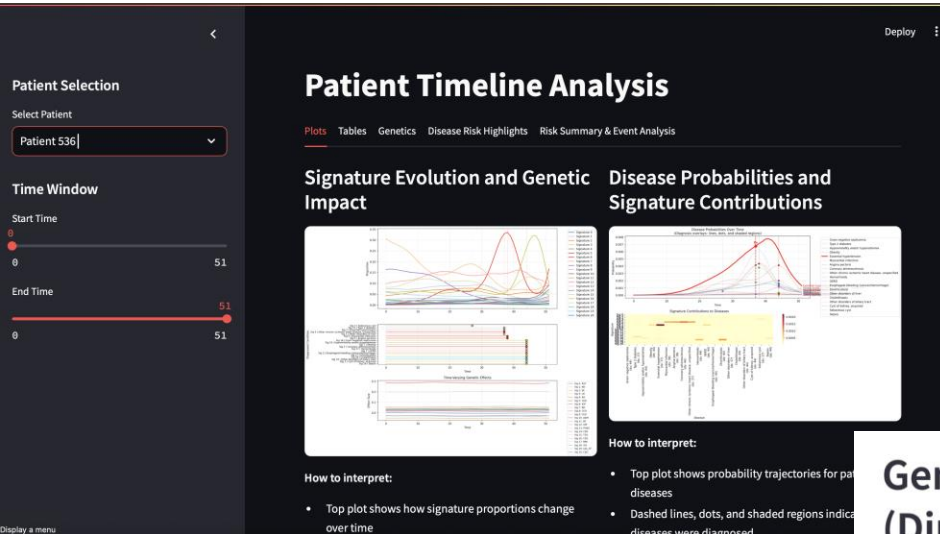




PREVENT: AUC
0.65
Aladynoulli
median dynamic
0.901

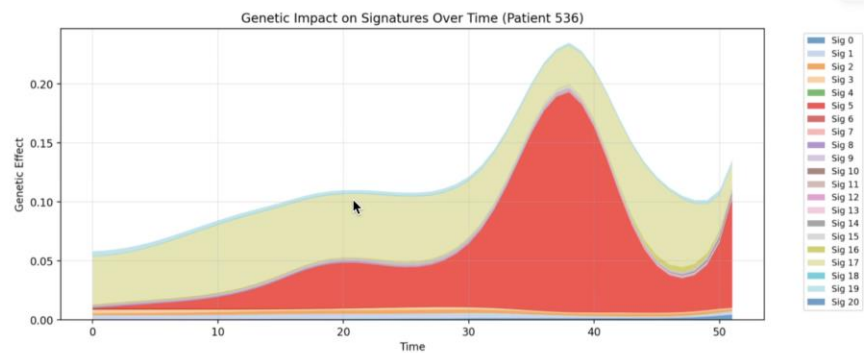
What does **Dynamic** Look like?





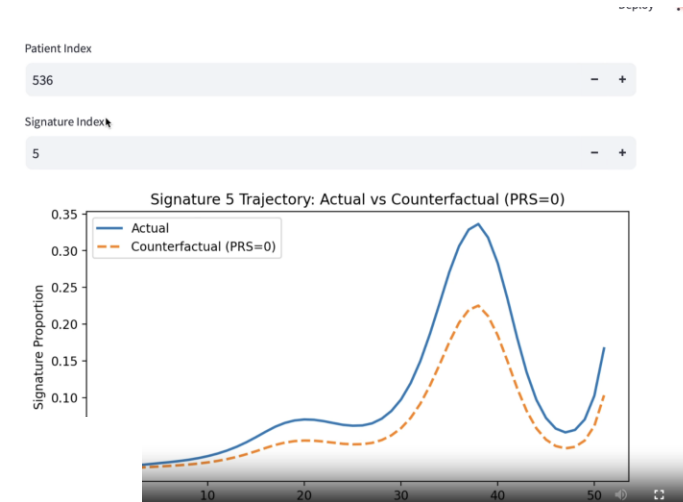
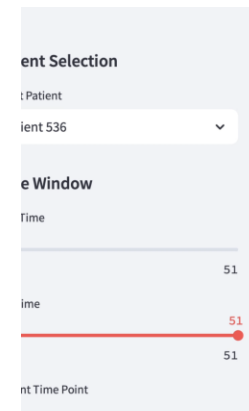
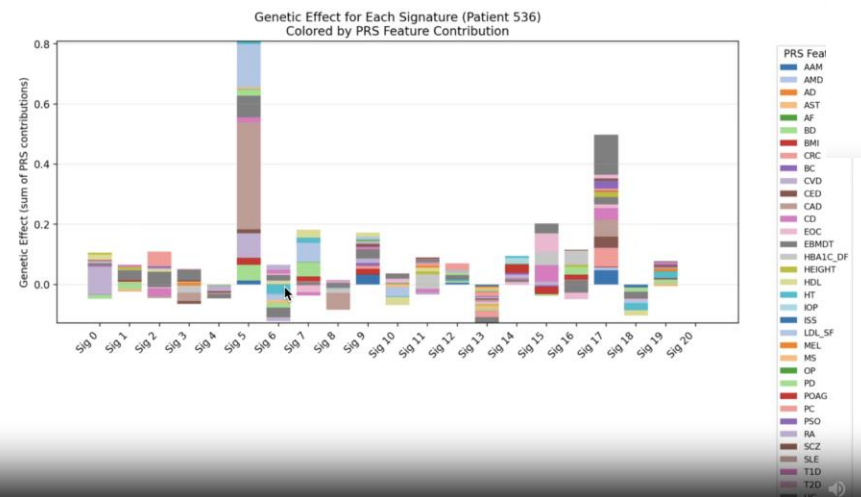
Genetic Impact on Signatures Over Time (Direct Weighting, Selected Patient)

This plot shows the direct genetic impact on each signature over time for the currently selected patient, using their PRS and the model's gamma weights.

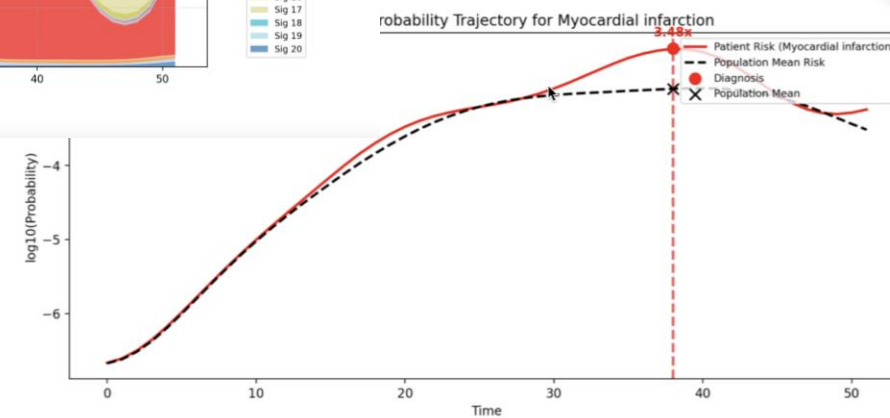


Patient, Stacked by PRS Feature

This stacked barplot shows the overall genetic effect for each signature for the currently selected patient with each bar colored by the fractional contribution of each PRS feature (G^* gamma).



[38]



Fold Enrichment of Disease Probability at Diagnosis

Discussion Points

- Tradeoffs of models where its highly precise, but difficult to communicate or models where it could potentially operate entirely behind the scenes and integrate into the EHR
 - Methods vs BlackBox

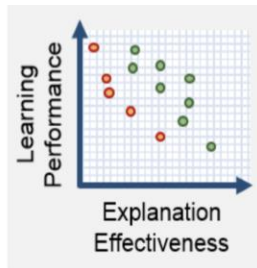


Figure 1: A fictional depiction of the “accuracy-interpretability trade-off,” taken from the DARPA XAI (Explainable Artificial Intelligence) Broad Agency Announcement [18].

Stop Explaining Black Box Machine Learning Models for High Stakes Decisions and Use Interpretable Models Instead

Cynthia Rudin
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Abstract

Implementing Machine Learning in Health Care — Addressing Ethical Challenges

Danton S. Char, M.D., Nigam H. Shah, M.B., B.S., Ph.D., and David Magnus, Ph.D.

Department of Anesthesiology, Division of Pediatric Cardiac Anesthesia (D.S.C.), the Center for Biomedical Ethics (D.S.C., D.M.), and the Center for Biomedical Informatics Research (N.S.), Stanford University School of Medicine, Stanford, CA

- What would be required to have a model that includes genomics, opportunistic imaging like CAC/CT-coronary, and AI image processing (like subtle signals on ECG and TTE)


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Article | [Open access](#) | Published: 12 September 2023

A deep learning-based electrocardiogram risk score for long term cardiovascular death and disease

[J. Weston Hughes](#) , [James Tooley](#), [Jessica Torres Soto](#), [Anna Ostropelets](#), [Tim Poterucha](#), [Matthew Kai Christensen](#), [Neal Yuan](#), [Ben Ehlert](#), [Dhamanpreet Kaur](#), [Guson Kang](#), [Albert Rogers](#), [Sanjiv Narayan](#), [Pierre Elias](#), [David Ouyang](#), [Euan Ashley](#), [James Zou](#) & [Marco V. Perez](#)

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Opinion

EDITORIAL

AI IN MEDICINE

AI in Medicine—JAMA's Focus on Clinical Outcomes, Patient-Centered Care, Quality, and Equity

Rohan Khera, MD, MS; Atul J. Butte, MD, PhD; Michael Berkwits, MD, MSCE; Yulin Hsuen, ScD, MPH; Annette Flanagan, RN, MA; Hannah Park; Gregory Curfman, MD; Kirsten Bibbins-Domingo, PhD, MD, MAS

Thank you!!!

Pradeep Natarajan, MD MMSc
Romit Bhattacharya, MD,
Giovanni Parmigiani, PhD
Sasha Gusev, PhD



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