



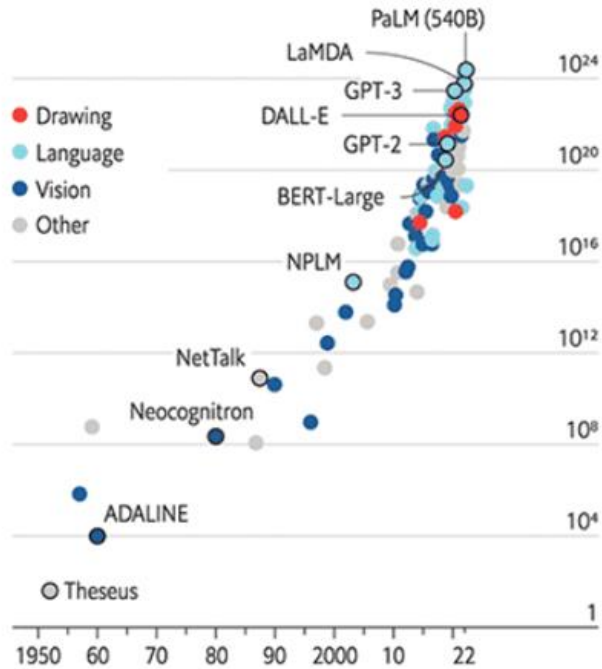
Universal Genomic Screening RNA Screening and AI for CVD

Santiago Miriuka, MD MSc PhD

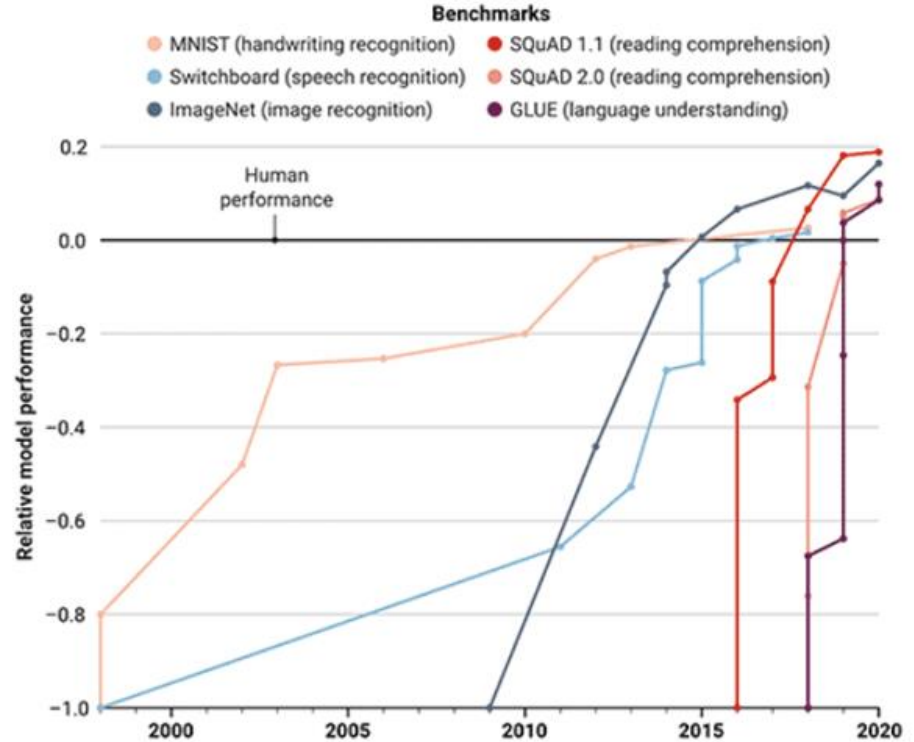
The future of disease detection and prevention is digital

- **Artificial Intelligence** - Data transformation and analysis
- **Genomic Sequencing** - Provides us with more data than ever before
- **Data management, processing and integration** - A vast amount of different sources of information providing insight on health and disease

The scaling up of Artificial Intelligence

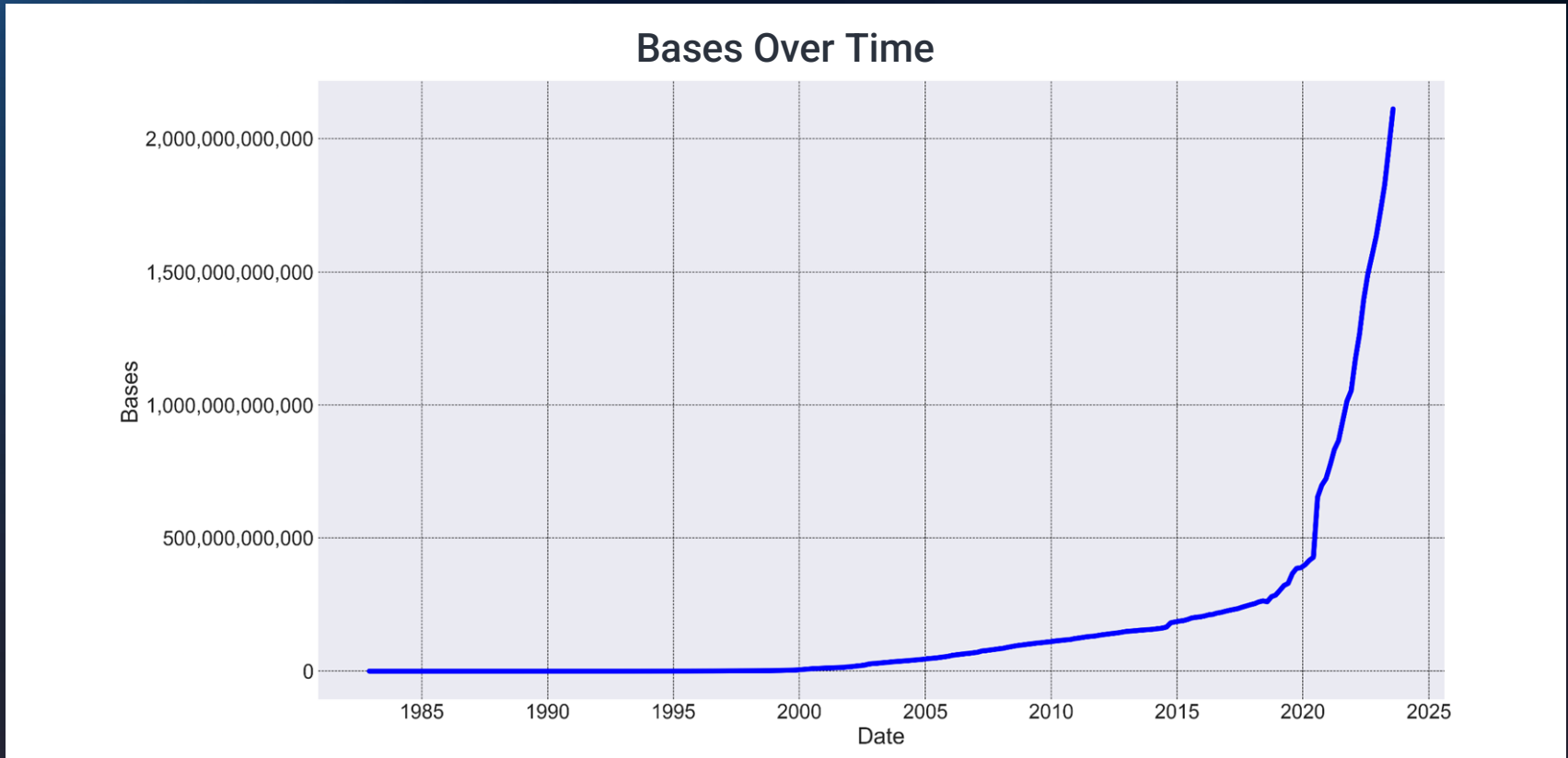


Sources: "Compute trends across three eras of machine learning", by J. Sevilla et al., arXiv, 2022; Our World in Data

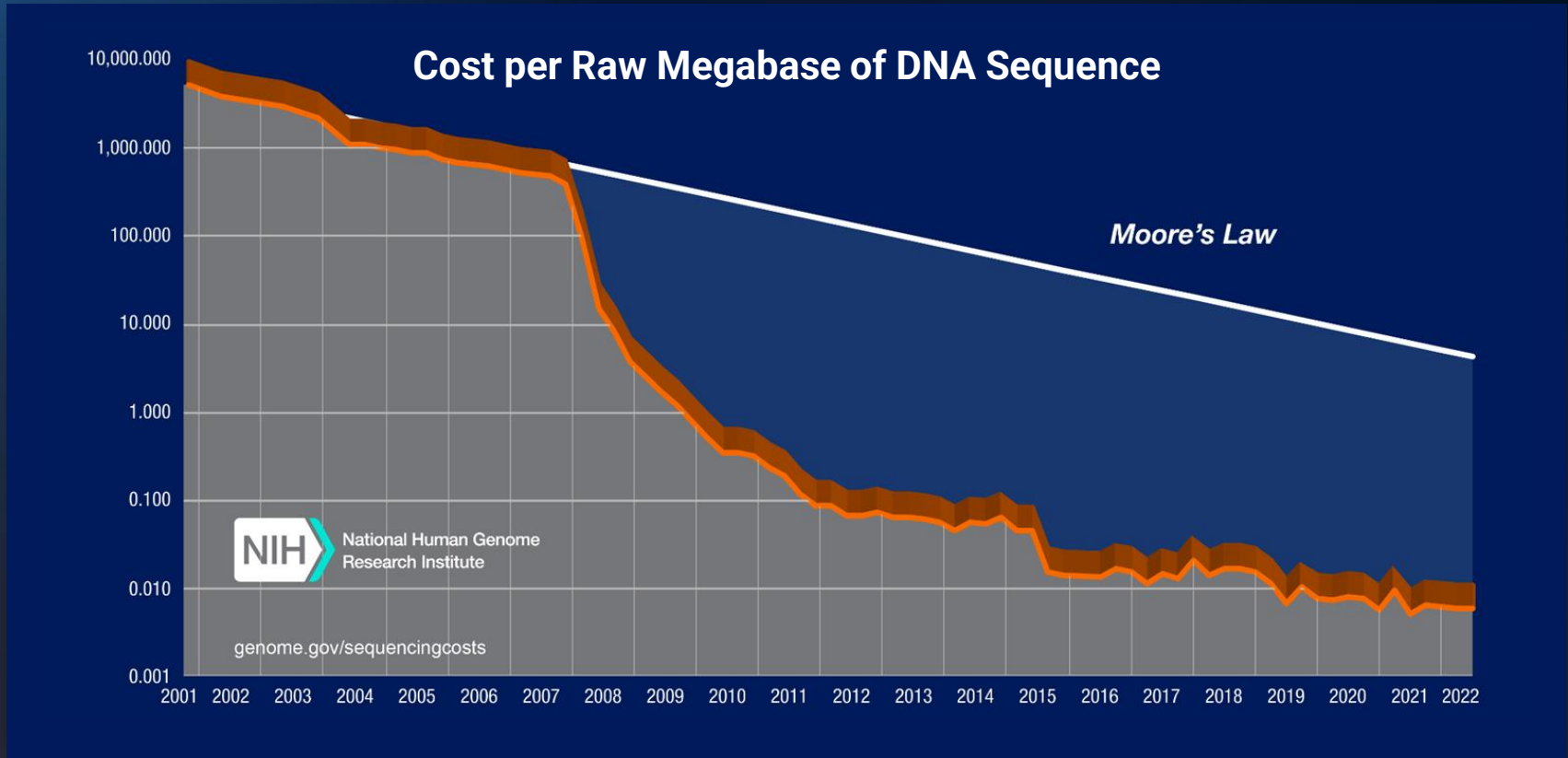


(GRAPHIC) K. FRANKLIN/SCIENCE; (DATA) D. KIELA ET AL., DYNABENCH: RETHINKING BENCHMARKING IN NLP, DOI:10.48550/ARXIV.2104.14337

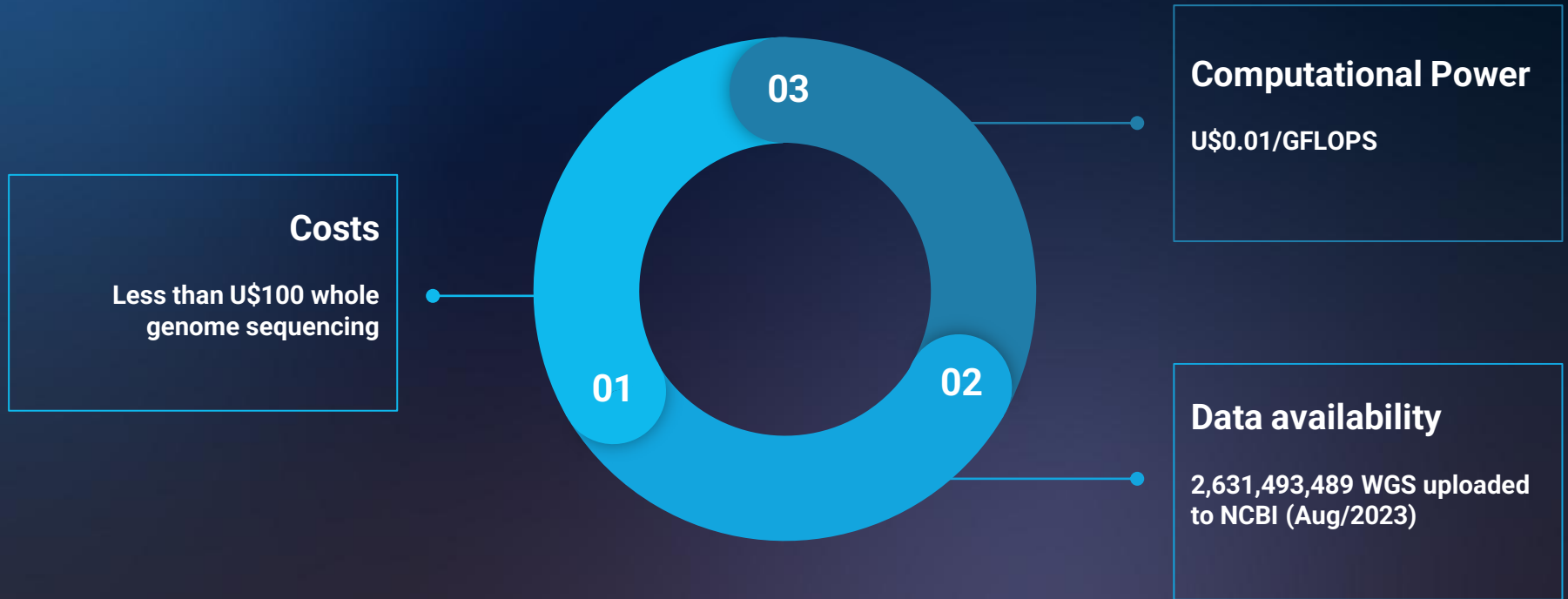
Genomic sequencing is exploding



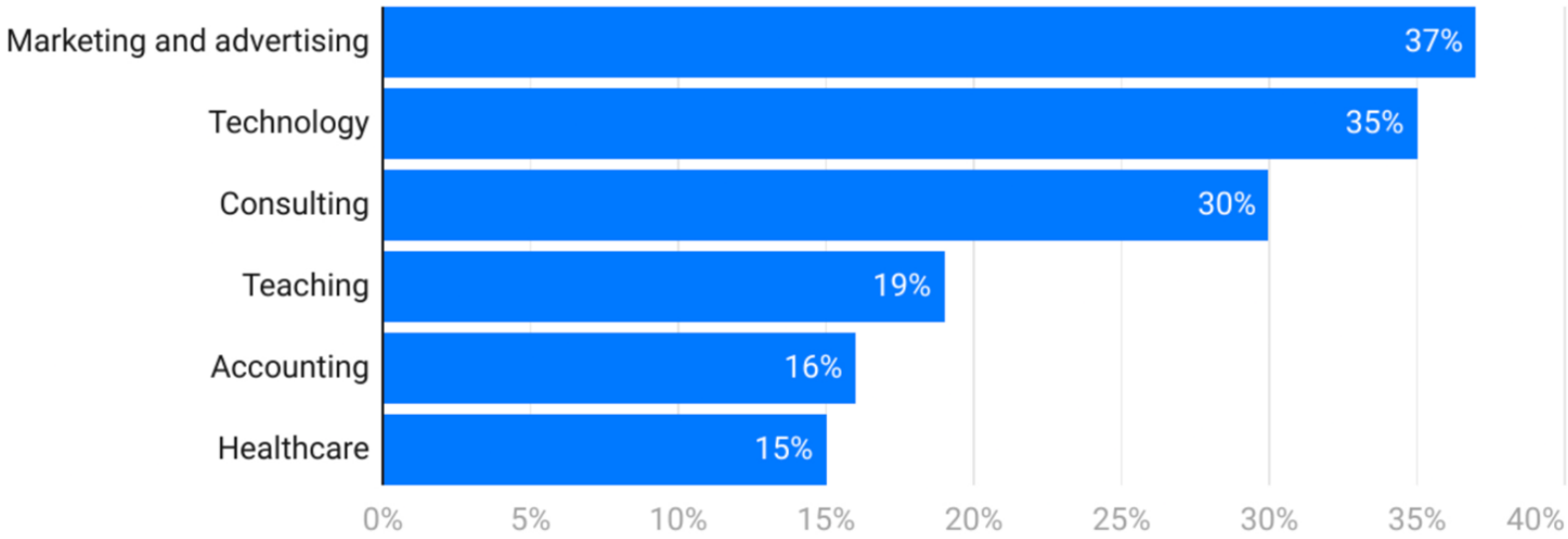
Cost of genomic sequencing



The Virtuous Circle of AI Genomics



Rate of generative AI adoption in the workplace in the United States 2023, by industry



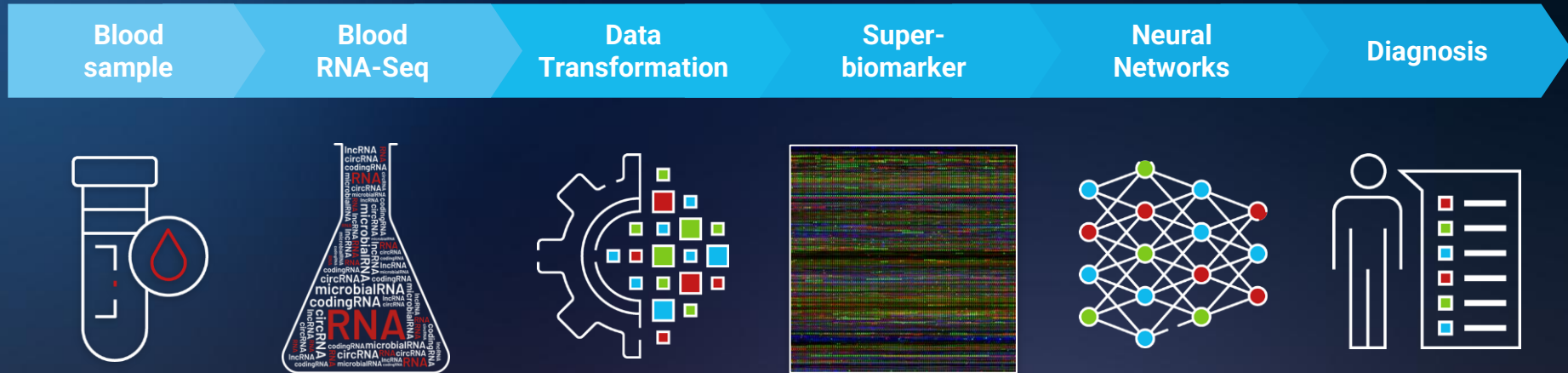
MultiplAI empowers anyone and their doctors to safely, privately, and accurately assess disease risk exceptionally early to proactively manage long-term health outcomes.

Next-gen
genomic sequencing

Proprietary data
transformation

AI
neural networks

Digitizing the blood



DNA mainly offers insights into **hereditary** conditions
RNA also reflects **environmental factors**

We express the **complexity of RNA** in a format **ideal for algorithmic analysis**

MultiplAI: Current Results



More than 1,000 samples sequenced

~100m reads

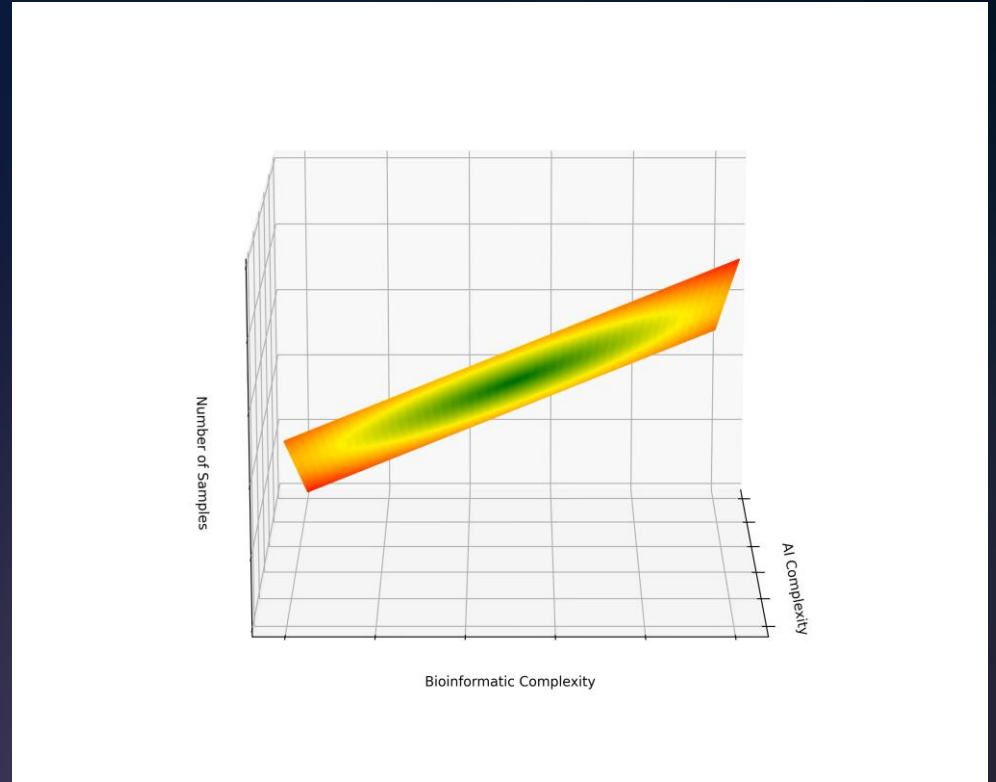
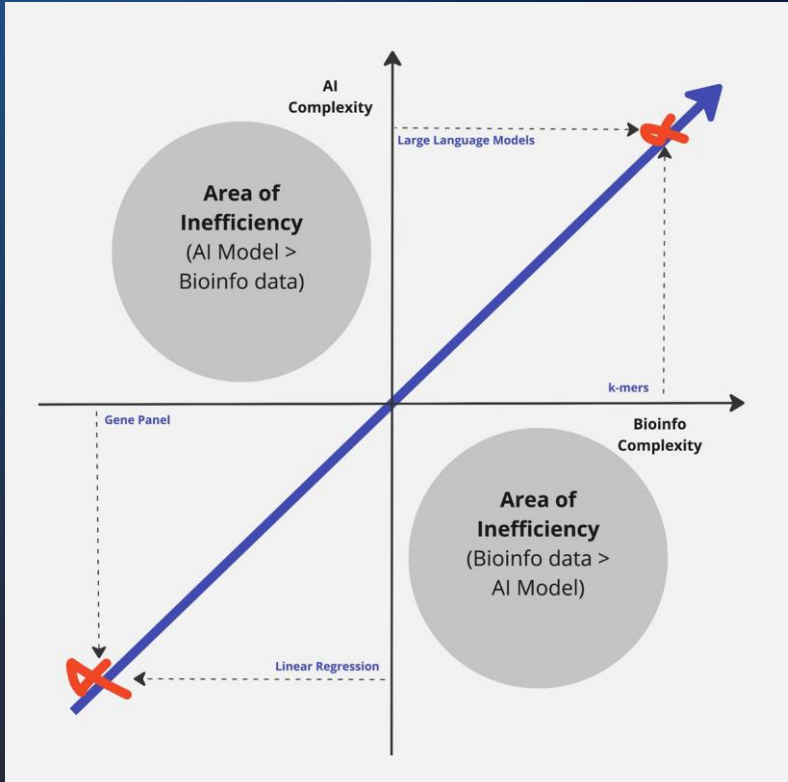
30-40k out of 65k genes detected

120k transcripts

7000 circRNAs

350k exons

The complex landscape of AI and Genomics



Cardiovascular disease (CVD) is the #1 cause of mortality globally

18m

deaths / year

15x

more women die of it
than **breast cancer**

#1

reason for waste in
healthcare spending

Almost **1 in 3 annual deaths globally** are caused by CVDs.

Traditional risk stratification **does not accurately detect vascular disease early enough**

High Risk




Died of stroke at 91

Low Risk



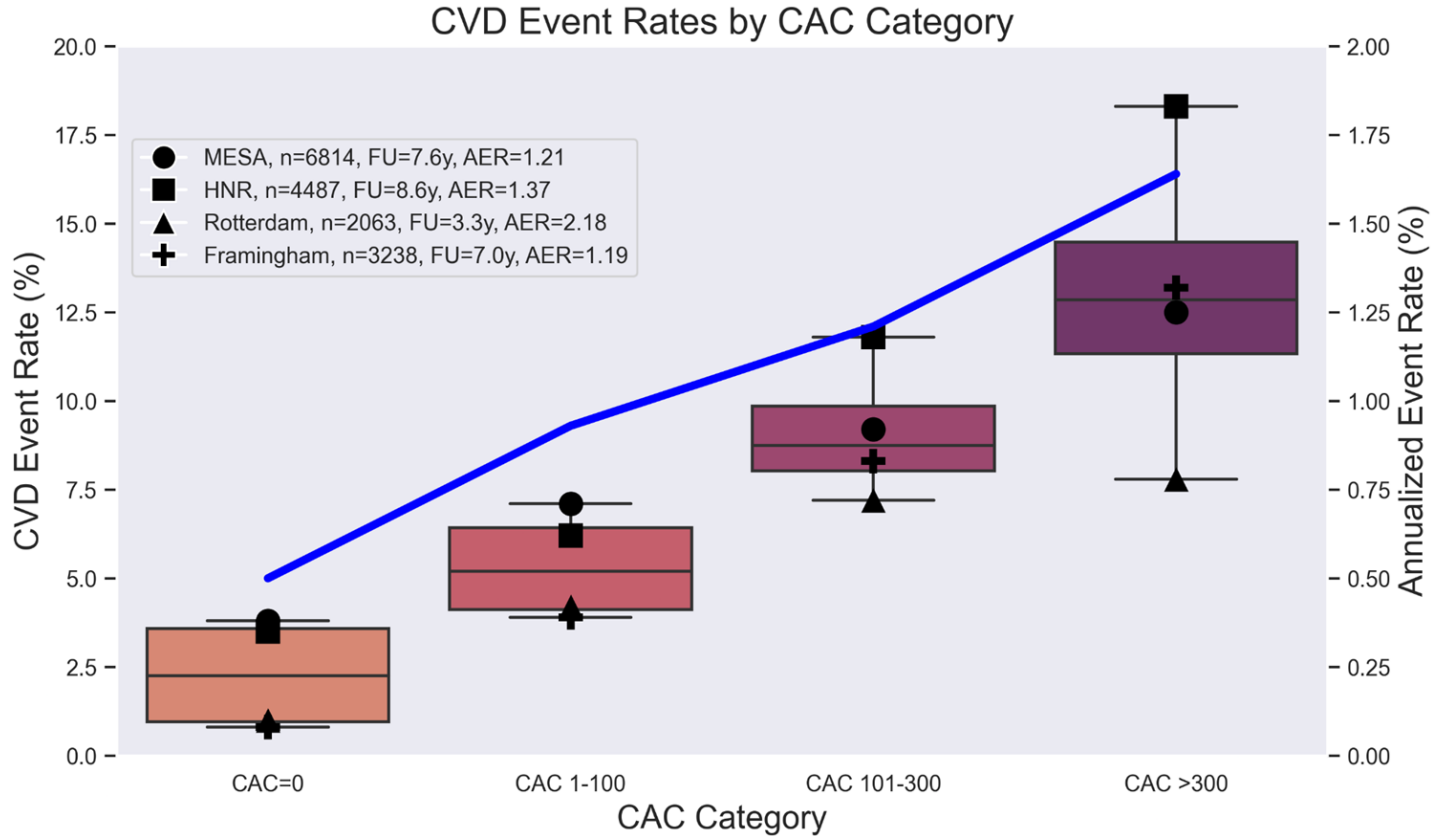
Heart Attack at 35

Millions of people are at risk without knowing it



**80% of CVD early deaths
worldwide could be prevented
with improved screening tools**

Presence of **coronary calcium** according to **risk score category**

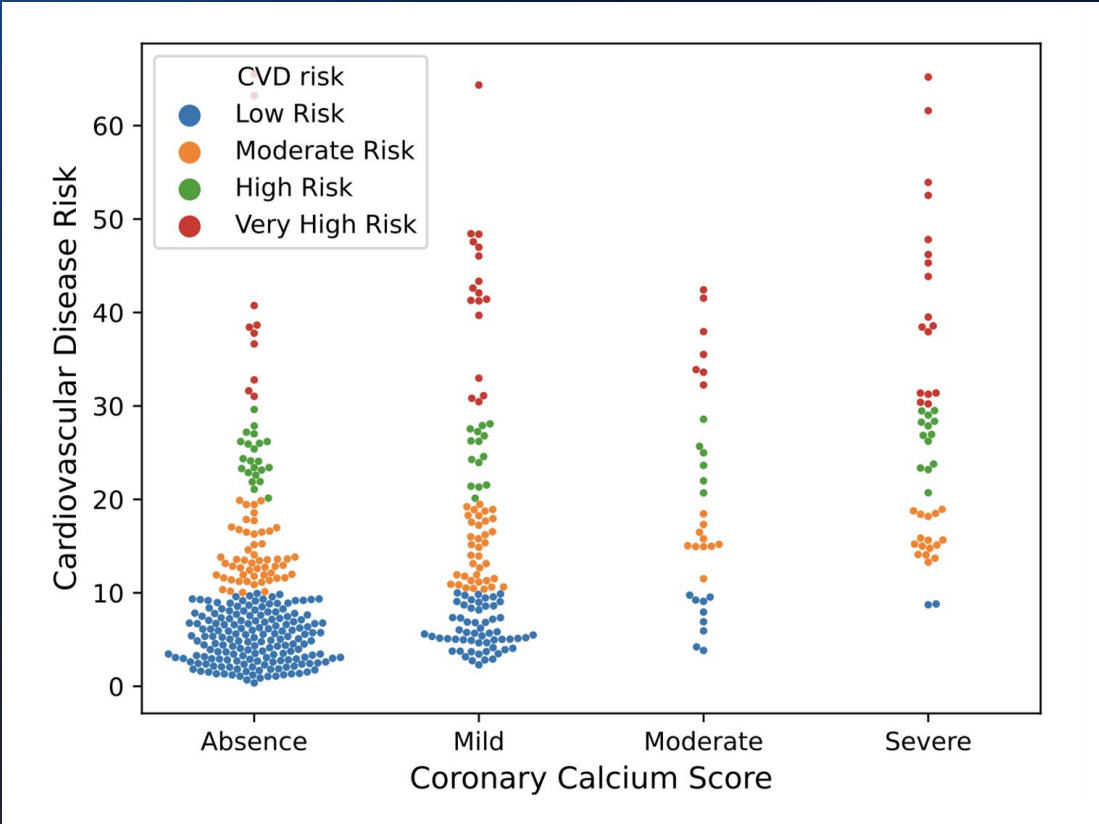


Gold standard of diagnosis has its challenges

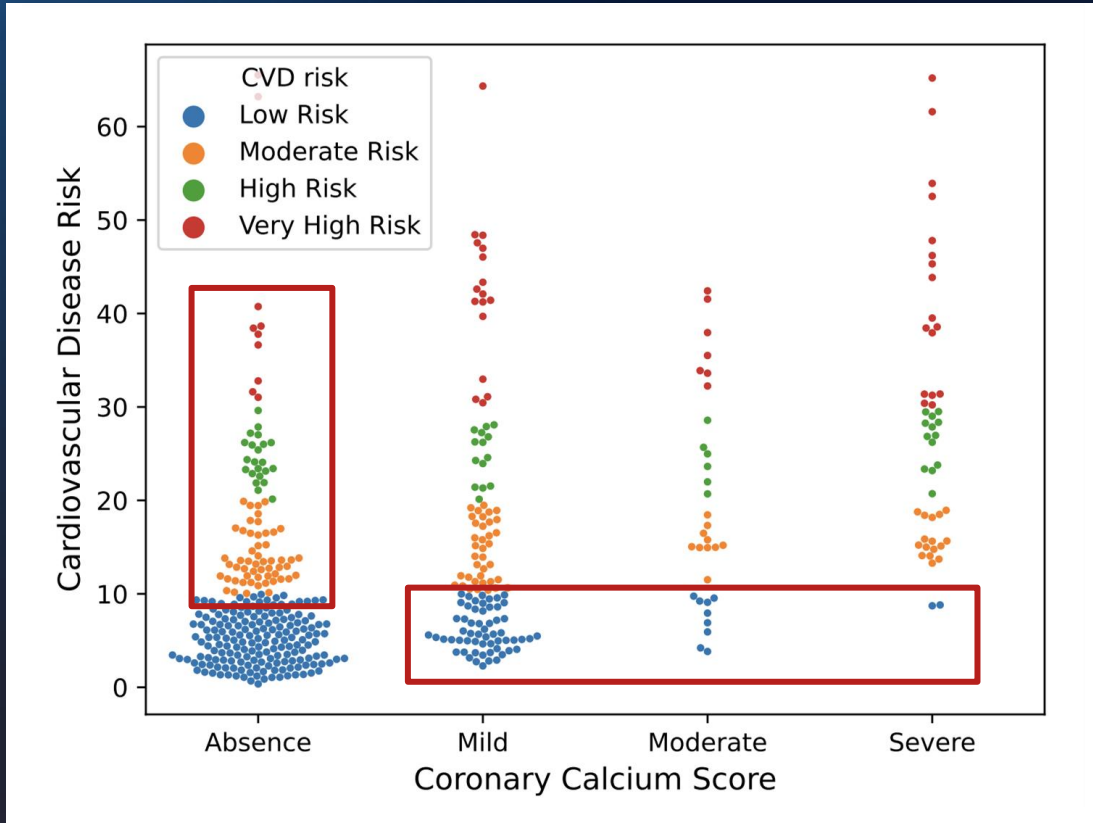
CT Scans not recommended
in low risk classification

Only 30% of countries have at least
one CT scanner per million people

CAC by Risk score (Framingham)

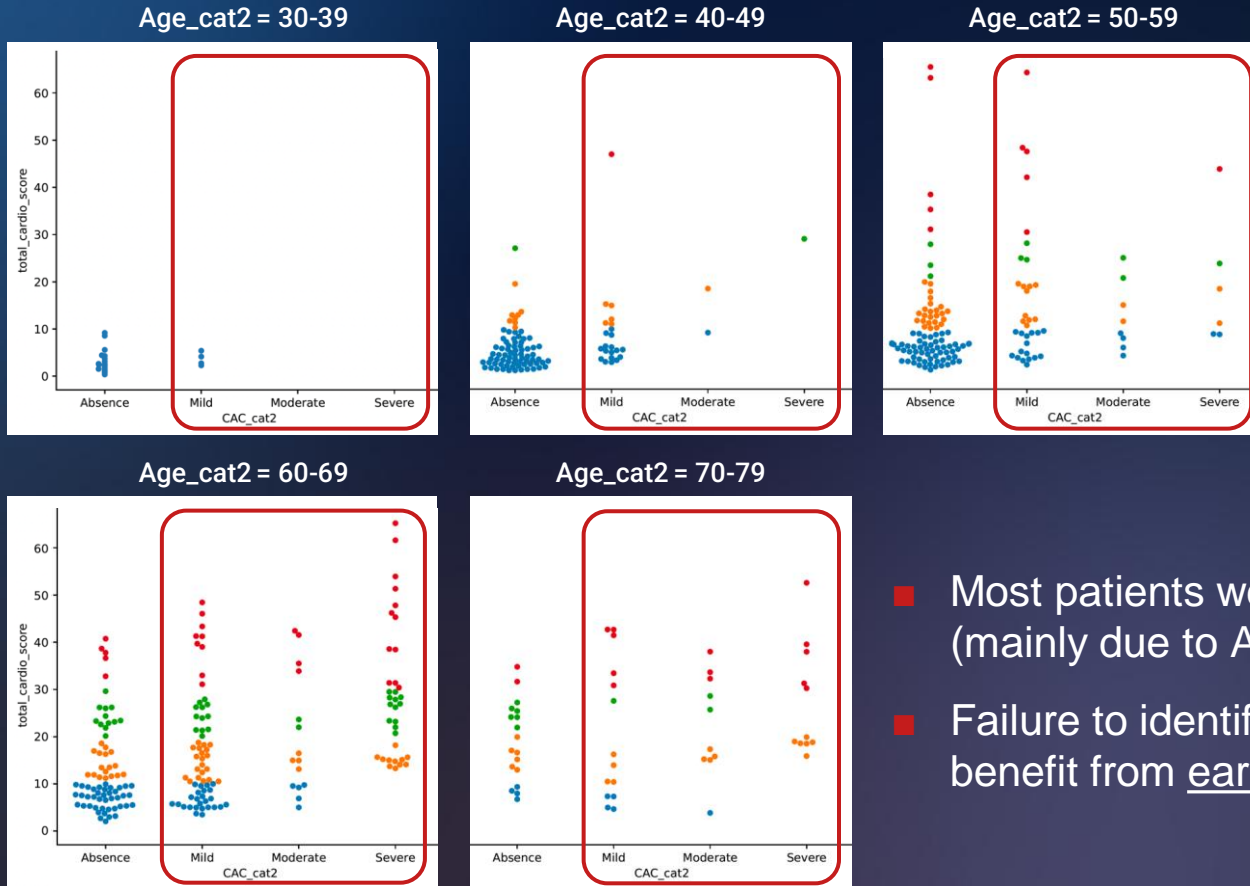


CAC by Risk score (Framingham)



- Prevalence of CAC>0 → 45 %
- Risk equations are not reliable for determining the presence or absence of coronary artery calcification (CAC)
- Misclassification rate = 33%

CAC by Risk score (Framingham) - By Age



- Most patients were classified as low risk (mainly due to Age)
- Failure to identify patients who would benefit from early prevention

Determination of the presence and extent of coronary calcifications by transcriptomic analysis of whole blood assisted by artificial intelligence: pilot study

The CCT-1 study

The CCT-1 Study: Study design & Data collection

Baseline

5 Year follow up

Participants



- Non prior CVD
- Men 40-70 years
- Women 50-70 years

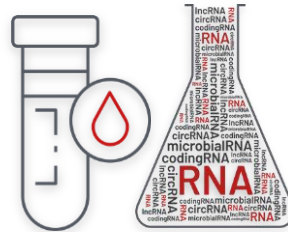
Chest CT Comparator



CT-Baseline/Outcomes

- Coronary artery calcium (CAC)
- Aortic calcium
- Steatosis
- Epicardial fat

Whole blood RNA



Clinical data



- Sex-Age
- Blood Pressure
- Body weight/ Height
- Drug treatments

Incident events

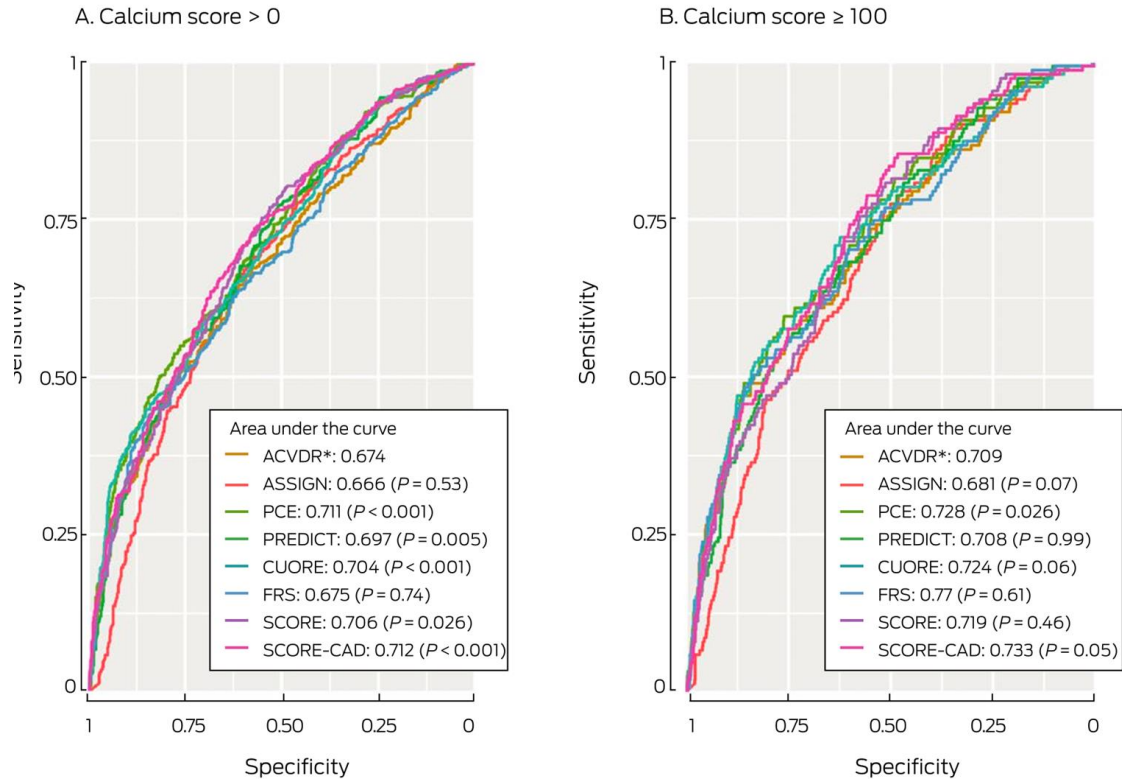
- Death from any cause
- AMI, angina
- Stroke or transient ischemic attack
- Hospitalized Heart failure
- Vascular revascularization
- Cancer

Diagnostic results
Precision for CAC detection

Prognostic results
Precision for predicting
incident events

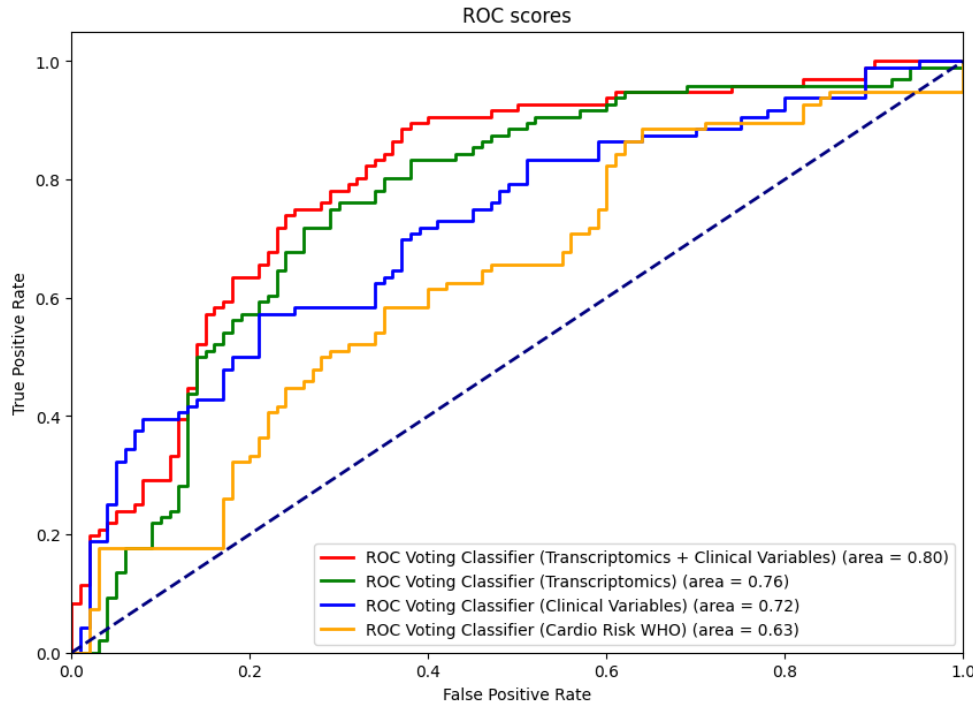
CCT1 study results. Area Under the Curve (AUC)

Prediction of coronary calcium compared to the traditional approach



CCT1 study results. Area Under the Curve (AUC)

Prediction of ANY coronary calcium compared to the traditional approach

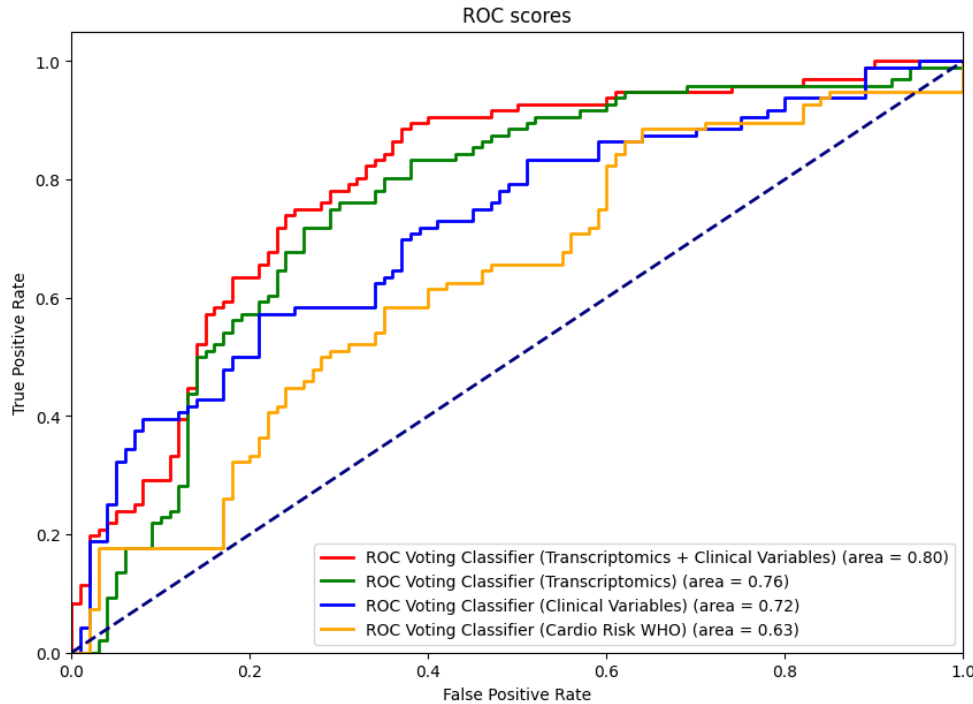


- CardioRisk WHO 0.63
- Clinical Variable 0.72
- Transcriptomics 0.76
- **Clinical Variables + Transcriptomics 0.80**

Outcome: Presence of coronary atherosclerosis (CT- Coronary Agatston units > 0)
Covariates: sex, age, smoker status, BMI

CCT1 study results. Area Under the Curve (AUC)

Prediction of ANY coronary calcium compared to the traditional approach



Variables Included in the model:

- Age and Sex
- Linear RNA
- transcripts
- circRNA
- Blood microbiome.

Outcome: Presence of coronary atherosclerosis (CT- Coronary Agatston units > 0)
Covariates: sex, age, smoker status, BMI

Early detection of subclinical CAC in asymptomatic individuals

The CAC-TRAIT study

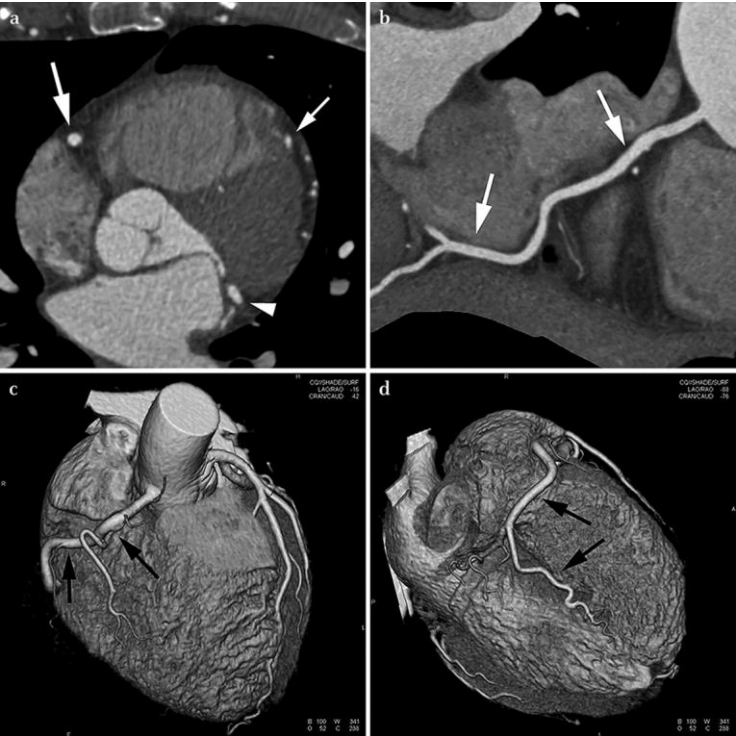
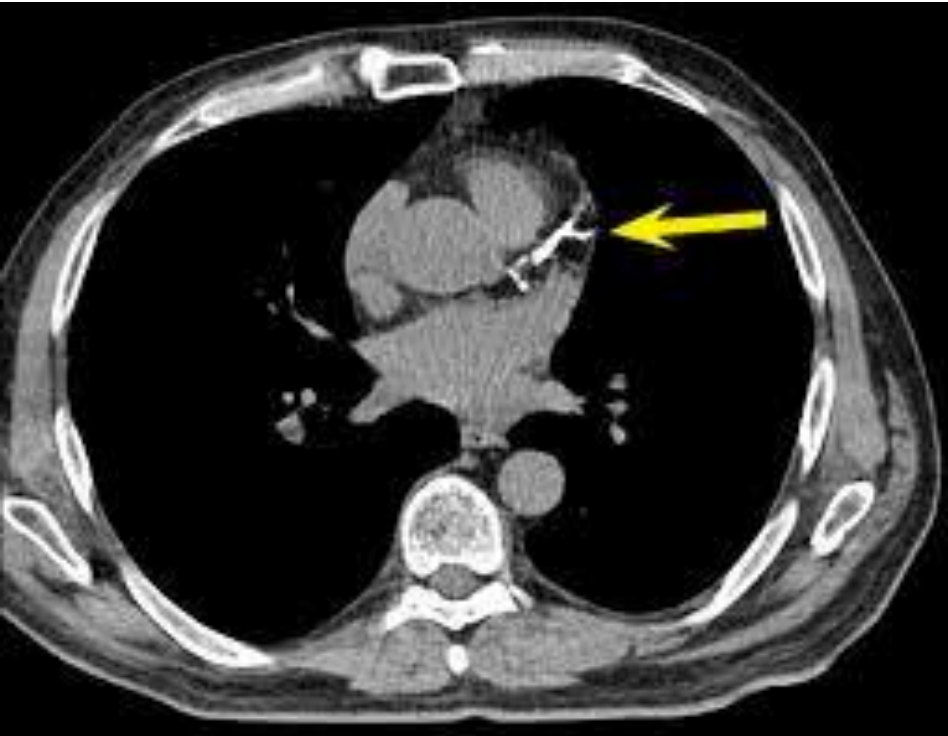
clinicaltrials.gov ID NCT05619042

Whole Blood Transcriptome Patterns according to the Coronary Atherosclerotic Plaque Burden determined by CT Angiography

The CORPLAQ-TRAIT pilot study

CT Scan

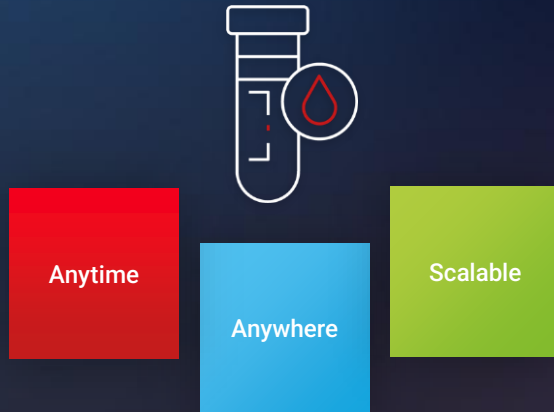
CT Scan versus angioCT Scan



Universal Genomic Screening for multiple pathologies

Artificial Intelligence and **Genomic Sequencing** allows us to face the challenges of complex diseases like CVD and make them available to anyone on the planet.

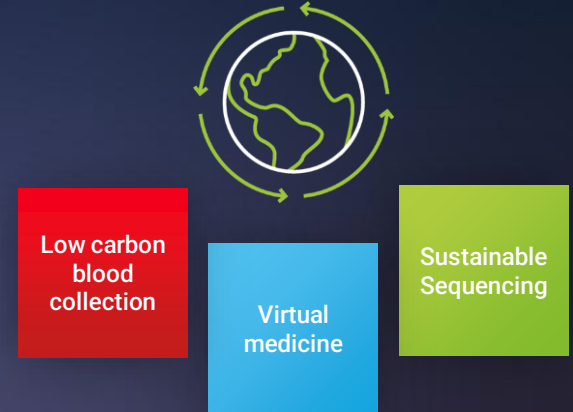
Universal **Accessibility**



Universal **Affordability**



Universal **Sustainability**



"The future of cardiology is personalized medicine, and AI is essential to making that happen."

– Eric Topol

Thank You

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