



Testing new models of research funding: One Brave Idea

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Disclosures

- Revenue from gene testing in cardiomyopathies
- Patents for cardiotoxicity testing in zebrafish
- Patents for drug discovery in zebrafish

- Novartis
- AtlasVentures
- ArrayBioPharma
- The Medicines Company
- Synthon
- Biogen Idec
- Sanofi
- Merck
- Pfizer
- Vertex
- AHA/Verily/Astra Zeneca

- Academic self-interest

One Brave Idea : AHA/Verily/Astra Zeneca

- \$75M for a single investigator
- 3 month/3 stage timeline
 - 250 words
 - 10 pages
 - Shark tank
- No constraints on use of funding
- Distinctive reporting structure

Executive Board
(Funders)

AHA

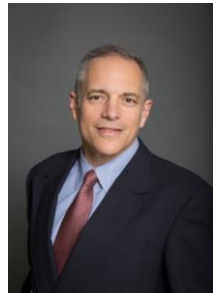
One Brave Idea™ Central Organization/CEO/CSO/CIO

Overall scientific strategy and direction

- › Core data science
- › Day to day operations/Project management
- › Scientific, reporting, financial and legal accountability
- › Incubator and other partnership development and maintenance
- › Cross-functional collaborative teams to maximize project velocity/efficiency
- › Lean and nimble

Scientific thought leaders - ad hoc advisors

**ONE
BRAVE
IDEA™**



Idea(s)

- Redefining coronary heart disease: at the edge of wellness
 - Redefinition of CHD in dynamic and quantitative biological terms
 - Identify new and much earlier true endophenotypes
 - Establish empiric approaches to moving from **deep to broad**
 - **New disease genes, new environmental contributors**
 - **New therapeutic approaches or new therapies**
 - **New preventative strategies**
- Testing new approaches to research execution and funding
- Contributing to a new ecosystem for discovery and care

Personnel: Initial core team



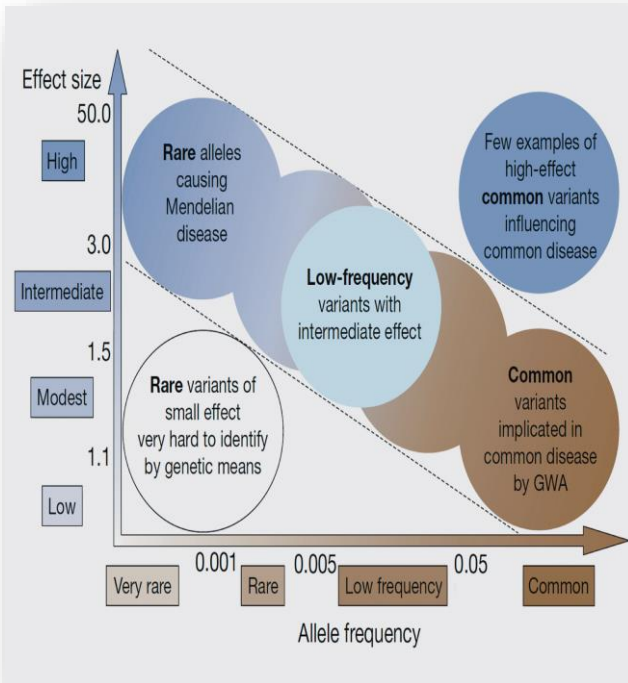
AHA/AZ/Google

Euan Ashley
Lazlo Barabasi
Elazer Edelman
Mike Gaziano
David Grayzel
Calum MacRae
Chris O'Donnell
Fritz Roth

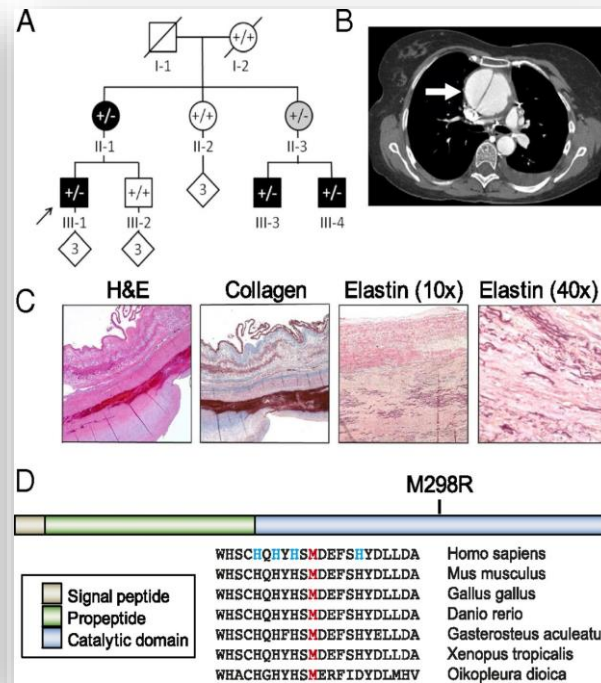
Ramachandran Vasan



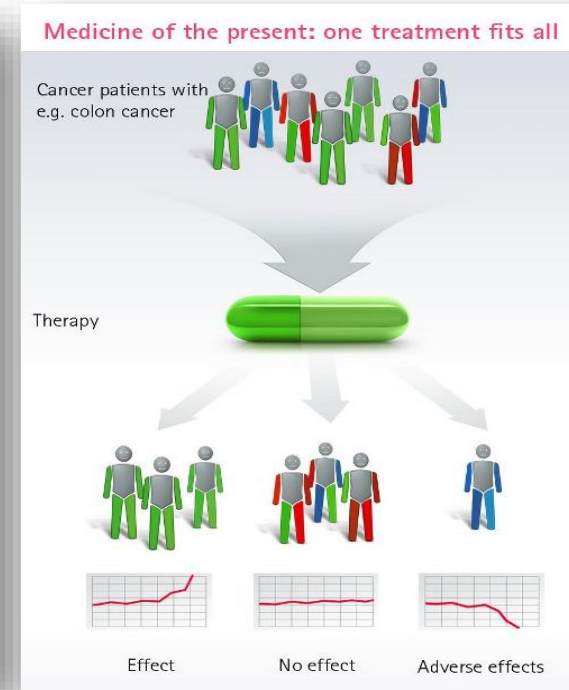
Project: Where is all the information?



Genetics



Clinical genomics



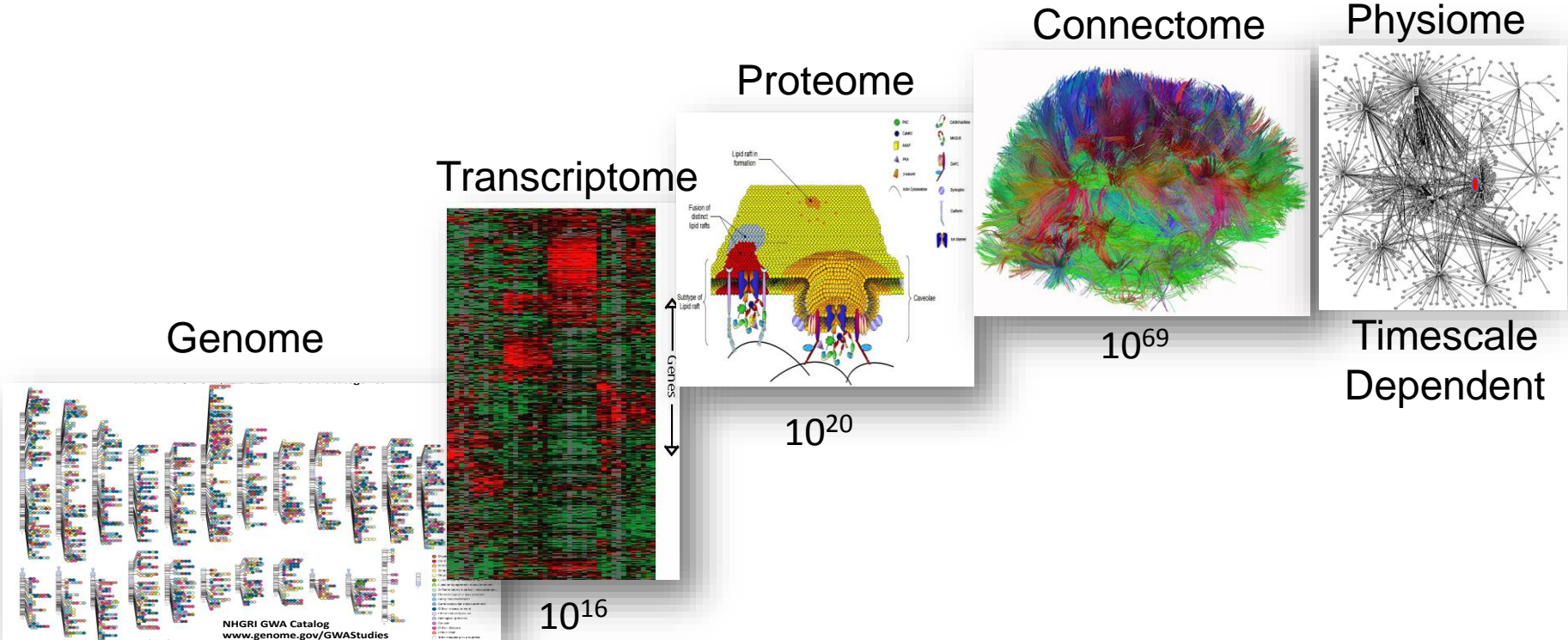
Clinical trials



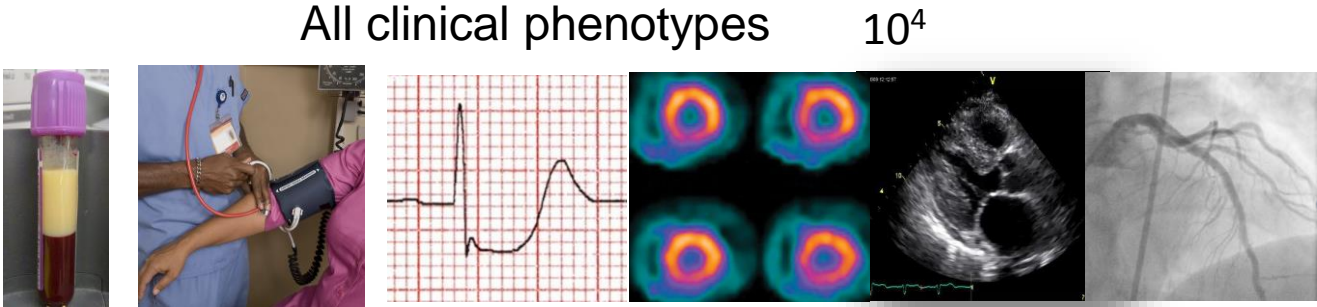
Care redesign

- **Phenotype is limiting in multiple areas of biomedical science**
 - Static or limited dynamic range
 - Almost all aggregates
 - Unidimensional with no organizing metadata
- **Few if any conditioning variables ever measured**
- **Most medical data**

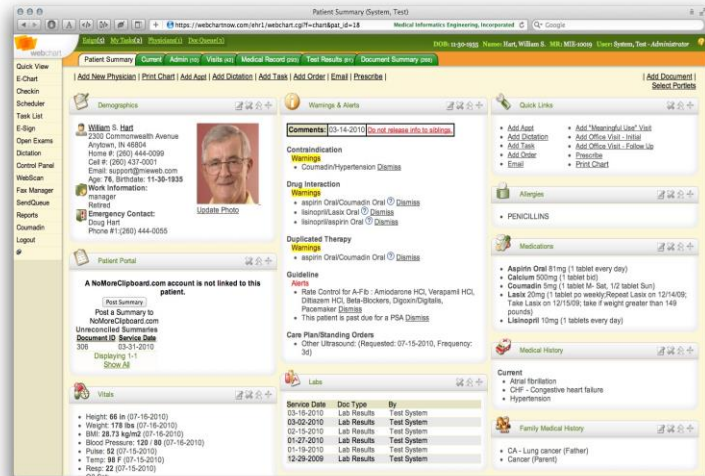
The phenotype gap



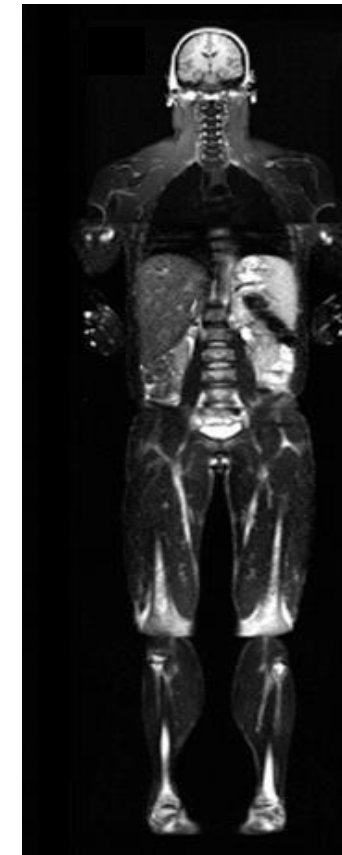
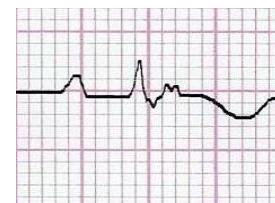
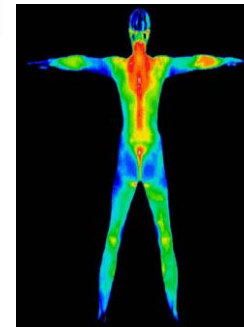
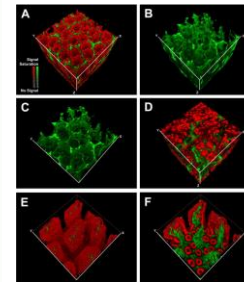
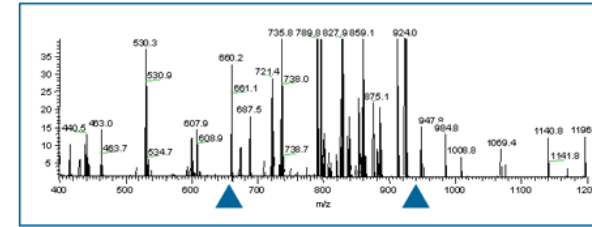
Exposome



Moving beyond legacy phenotypes



Glycosuria



EKG

Specific metabolites

Microbiome

Microcirculation confocal imaging

Adipose tissue mapping

Thermography

Everything else

Unstructured

18th century

Semi-subjective and duplicative

Lack of standardization

Cross sectional and static

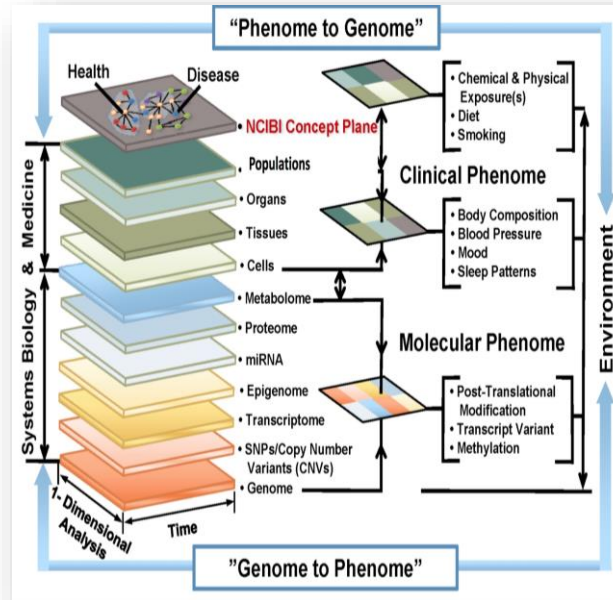
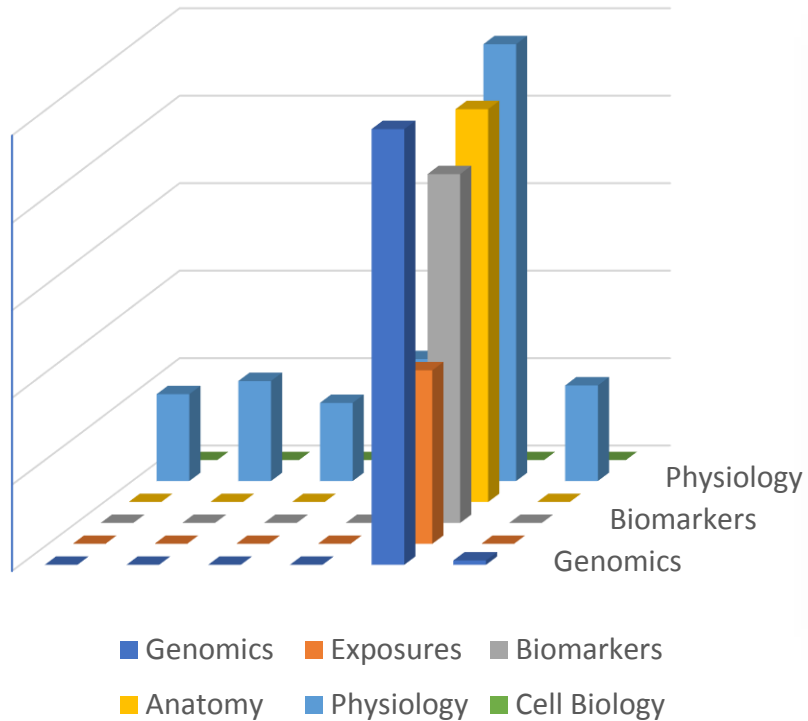
No metadata

High threshold for innovation

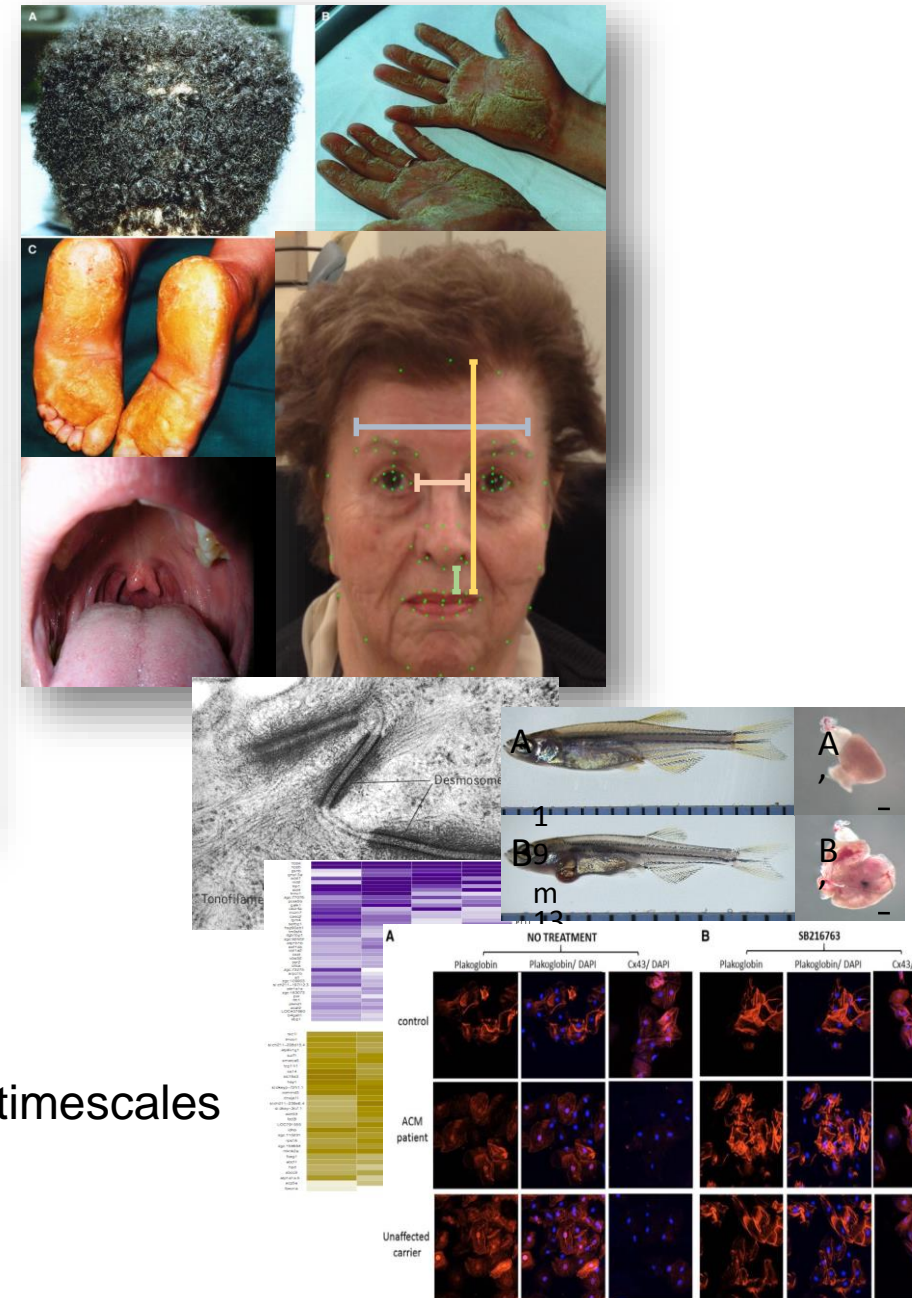
Tied directly to implementation evidence base

Broad vs deep

Information Space



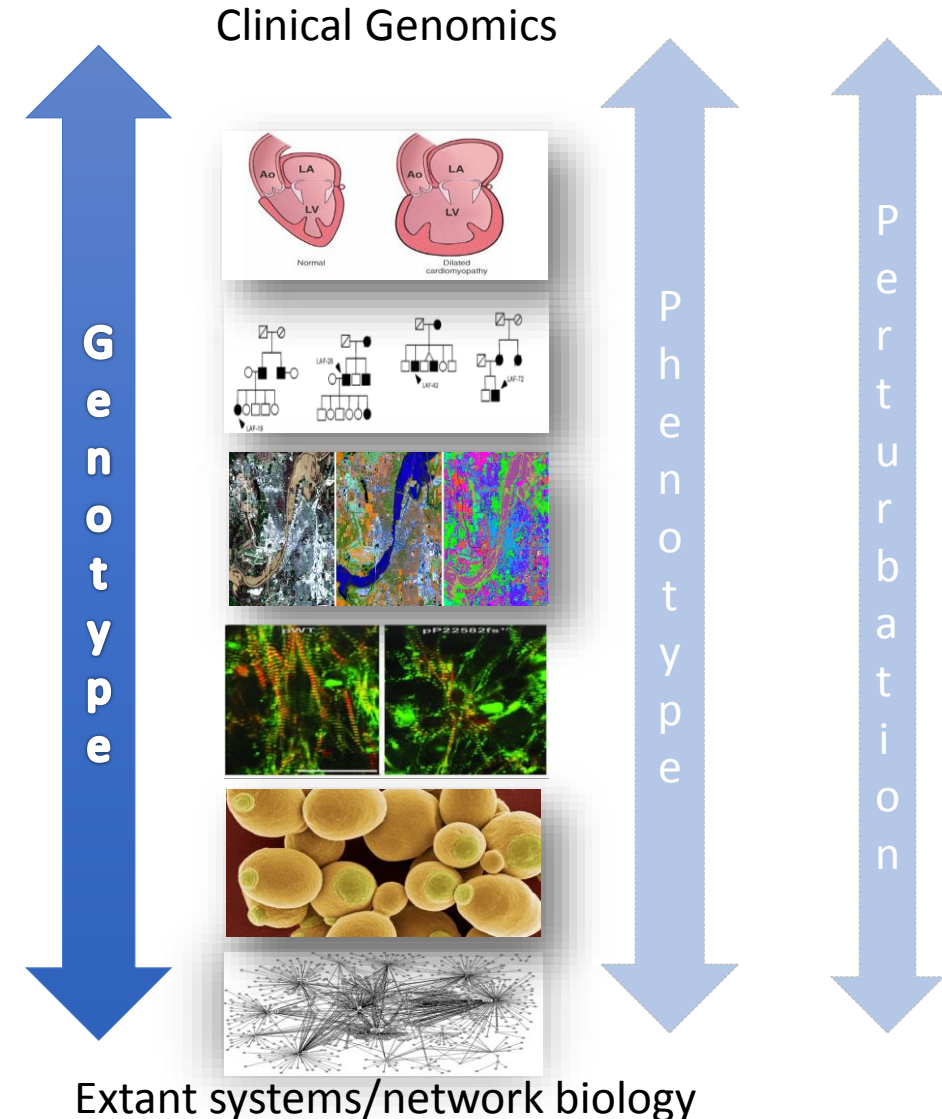
Kohane et al.



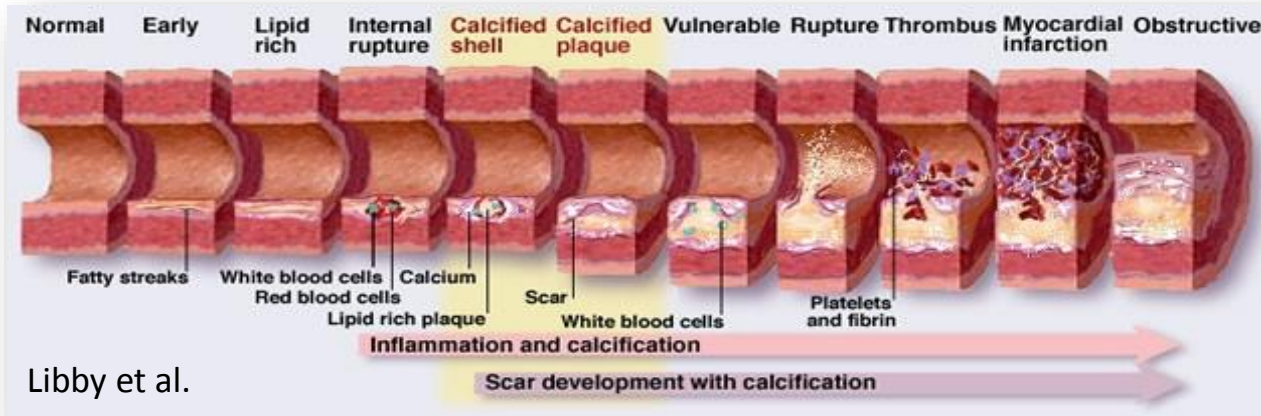
- Comprehensiveness
- Organizing metadata across datasets and models: perturbations and timescales
- New datasets-shared across biological models
- **A computable molecular/cellular/physiologic ‘physical exam’**

Linking clinical and basic science

- Mapping relationships across species
 - Genome
 - Phenome
 - Perturbations
- Multiple 'omics
- Cell biology, physiology
- Environmental and drug responses
- Multiple dimensions improve specificity
- Shared phenotypic lexicon
- 'Mechanistic' phenotypes in all species
- 'Co-clinical' modeling: real time application

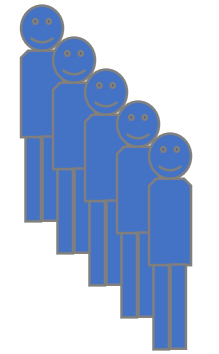


OBI: Detecting the earliest phases of disease

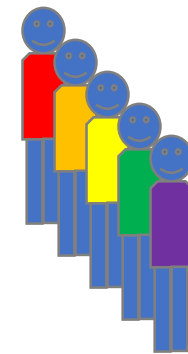


20-40 years

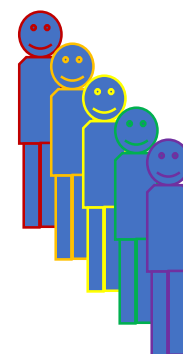
15-20 years



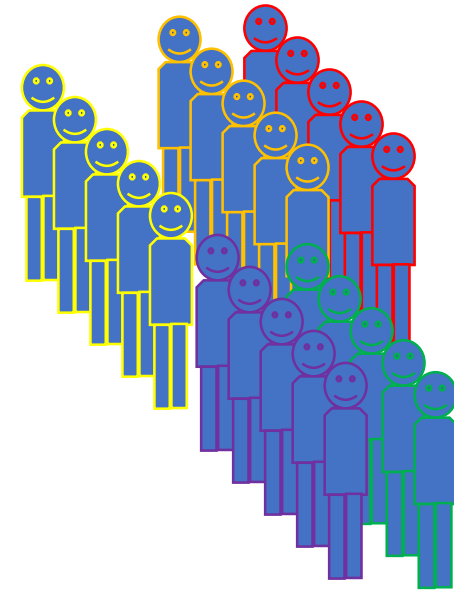
Generic pre-CHD



Different underlying forms of CHD



New ways to detect different forms of CHD



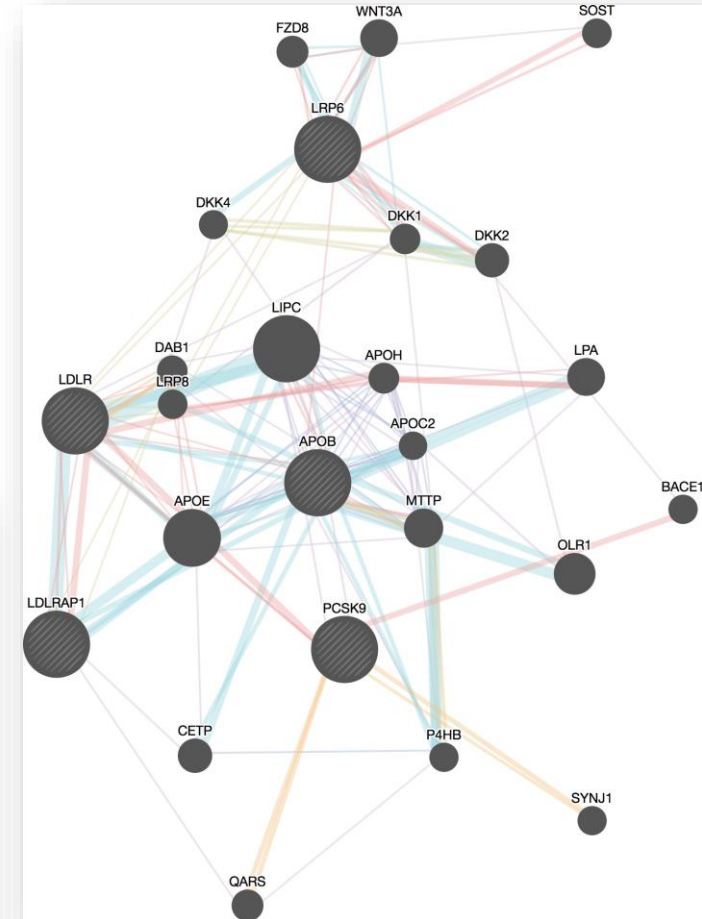
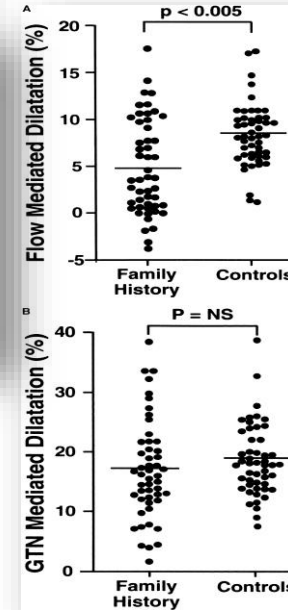
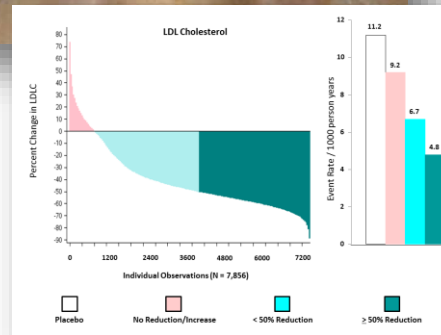
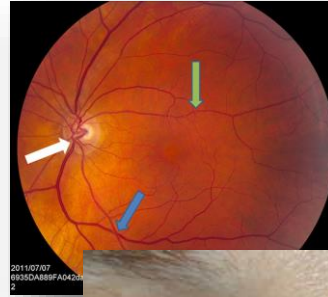
Scaling to general population for screening and disease Rx

- Most of what we know about CHD has emerged from a focus on the **latter 15-20 years** of the disease
- CHD represents many different disorders which resemble each other most in their later stages
 - Identifying new translatable markers of the very earliest stages of CHD
 - Define new underlying causal factors for CHD,
 - Develop technologies for population detection,
 - Move towards new therapies and preventative strategies

New pathways in atherosclerosis?

- Preclinical phenotypes
 - Discrete genetics
- Core genes
 - LDLR, ApoB, PCSK9
- Extant biology predictions
 - HTN
 - T2DM
 - Cognitive decline

Preclinical phenotypes



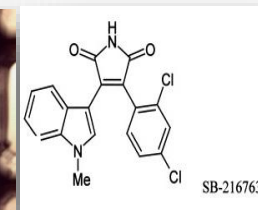
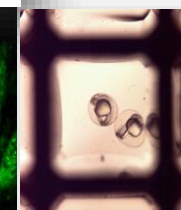
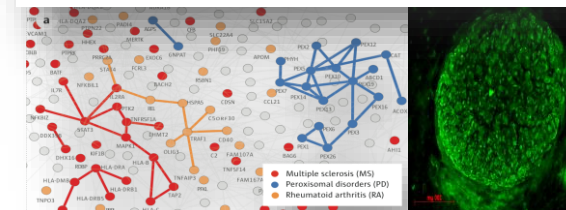
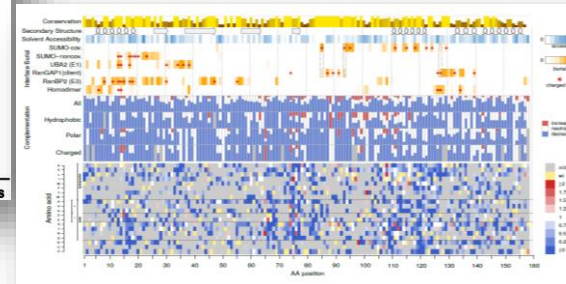
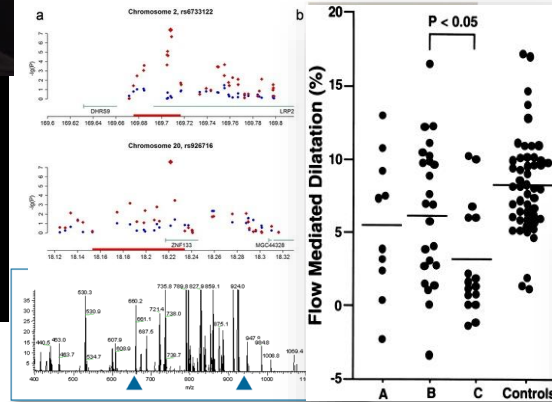
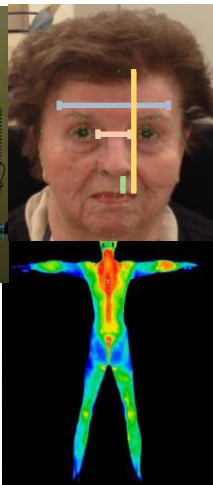
A generalizable approach for phenotype discovery

Phenotype and perturbation discovery
Cells to organisms

Validation in kindreds and genotyped cohorts

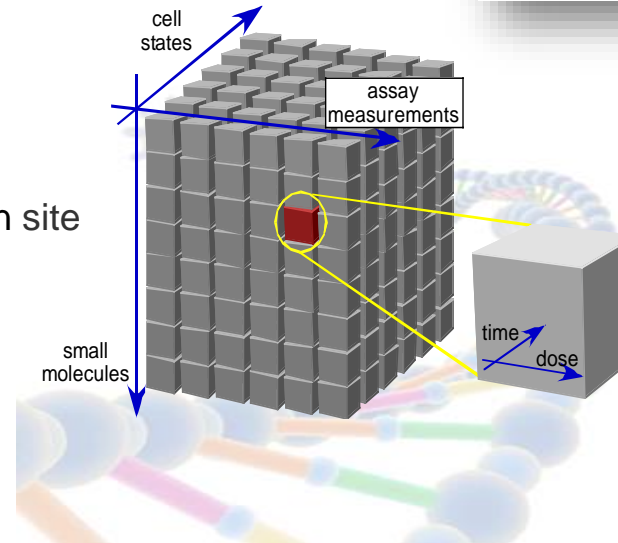
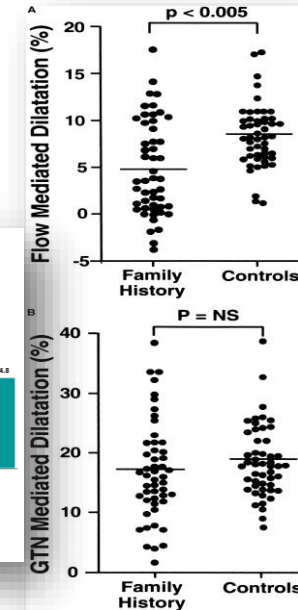
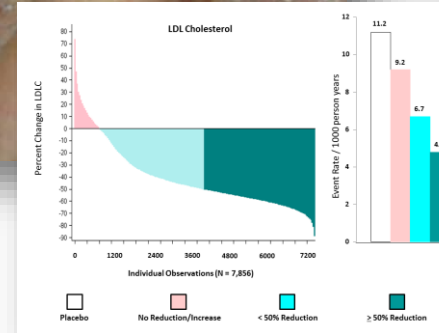
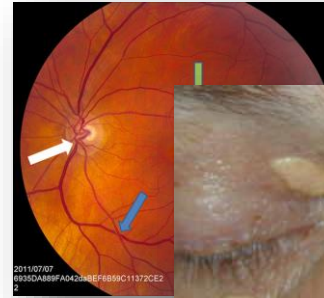
New causal pathways
Perturbation/phenotype discovery

Democratization of phenotyping
Large simple trials



Orthogonal phenotypes: scalability by design

- Rapid cycle development
 - Plug and play
 - Flexible bioengineering and computational infrastructure
 - Benchmarking technologies
 - Developing our own technologies
 - Rigorous biologic insight for positioning
 - Stimulus identification
- Disease-enriched populations
 - Large numbers of subjects
 - Large proportion already genetically defined
 - Active Precision Medicine efforts
- Based in outpatient clinic
 - Physician encounter occupies <20% time spent on site
 - Integrating genomics and care reinvention
 - Mapping onto existing disease framework
 - Controlled environment: stimulus-response pairs
- Efficient scaling to population cohorts
 - FHS
 - Million Veterans Program
 - Verily, Microsoft, AHA My Research Legacy



Funding mechanism considerations

- PPG format
- Venture fund: 'for profit' vs 'not for profit'
- Discrete commercial entity
- Closed end vs sustainable
- Additional partners and fundraising
 - Industrial/Foundations/Philanthropy
 - Focused on alignment: Pharma/Tech/Biotech/Device/Retail/other
 - In-kind resources
 - Governance
- Partnerships with traditional funders
 - Joint investments: shared returns
 - Federal and international cohorts-fee for service
 - Training mechanisms
 - Infrastructure development

Structural features of program

- Administrative
 - Central core with fiduciary, legal and reporting responsibility
 - Renewable engagement of scientific team members and SAB
 - Flexibility to continue to engage/disengage based on science
 - Executive board with oversight
- Highly goal directed
- Objective go/no go metrics for each funding component
 - Scientific rigor
 - Alignment with goals of program
- Efficient funding cycles-<6 weeks
- Indirect costs
- Intellectual property

Executive Board
(Funders)

AHA

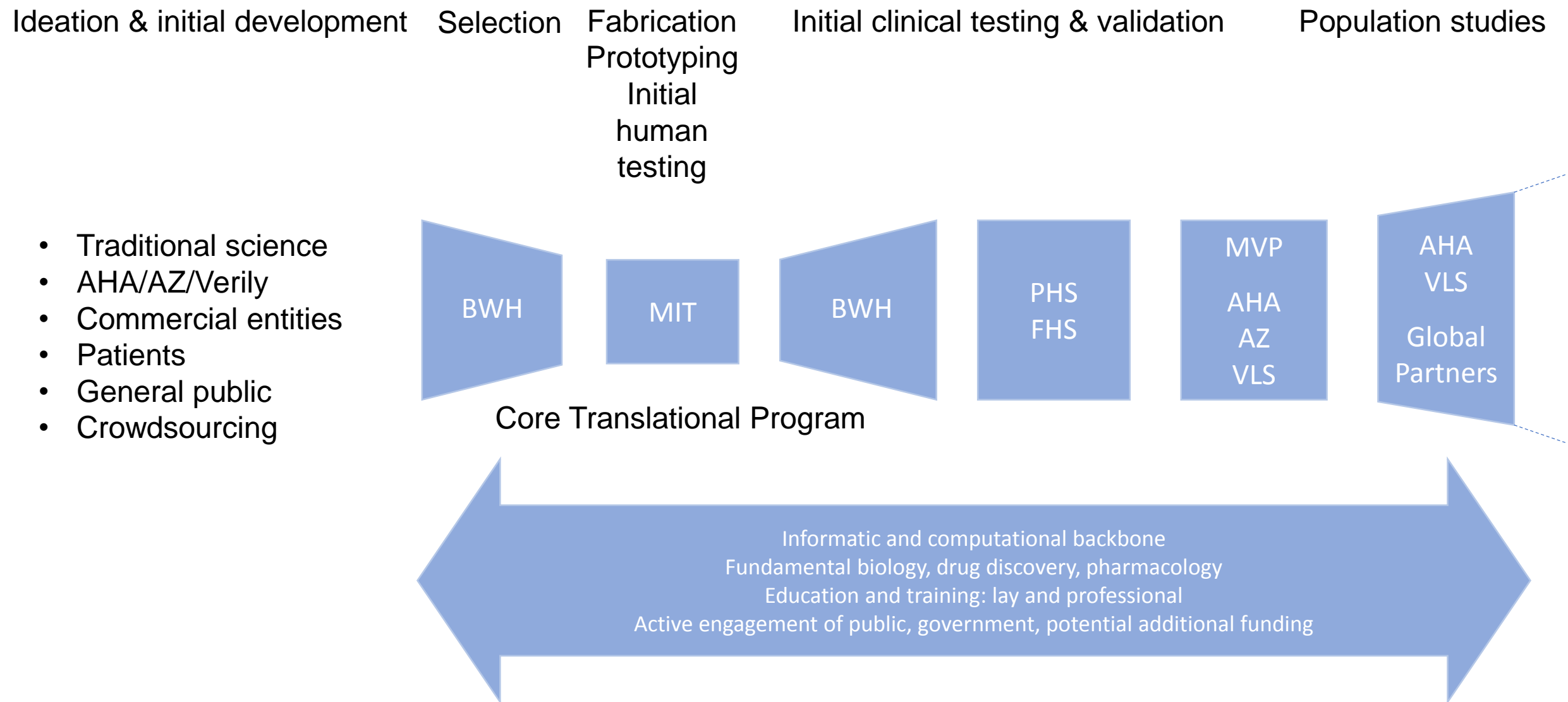
One Brave Idea™
Central Organization

CEO

CSO/CDO

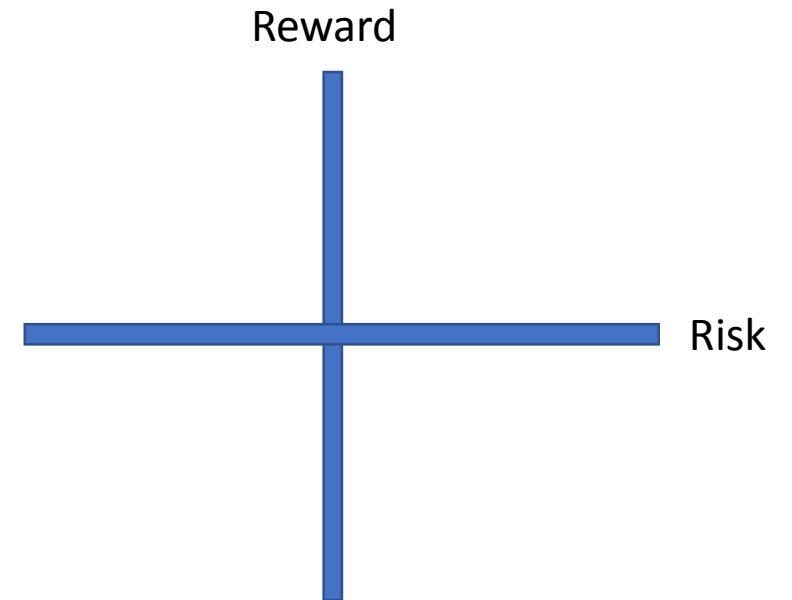
Active projects
Participating scientists

Core infrastructure



Initial science strategy

- Core data science
- Balanced portfolio
 - Early initiation of low risk/high yield
 - Exposure quantitation
 - Extant atherosclerosis cell biology
 - Population science with existing data
 - Prioritization of high risk/high return projects
 - Early basic science
 - Foundation for later implementation
 - Developing criteria for moving to scale
- Internal and external RFAs
 - 'Quality, price and performance'
 - Testing the structure of the program
- Optimization of teams for projects
 - Members
 - Locations
- Exploring partnership mechanisms



Agnostic vs Directed
Funding vs Engagement

Culture

- **Emphasis on alignment and engagement**
- **Diversity-personnel, ideas etc**
- **Building community engagement for the long haul**
- **Balancing comprehensiveness and utility-the academic conundrum**
- Rigorous metrics including effect size
- Active cross-fertilization between projects **within OBI**
 - 20% rule or equivalent: rewarding multi-disciplinarity/engagement
- Teaming across groups to advance goals
- Team member development
 - Maintenance of long term fundability for investigators
 - Development of orthogonal skills for all team members

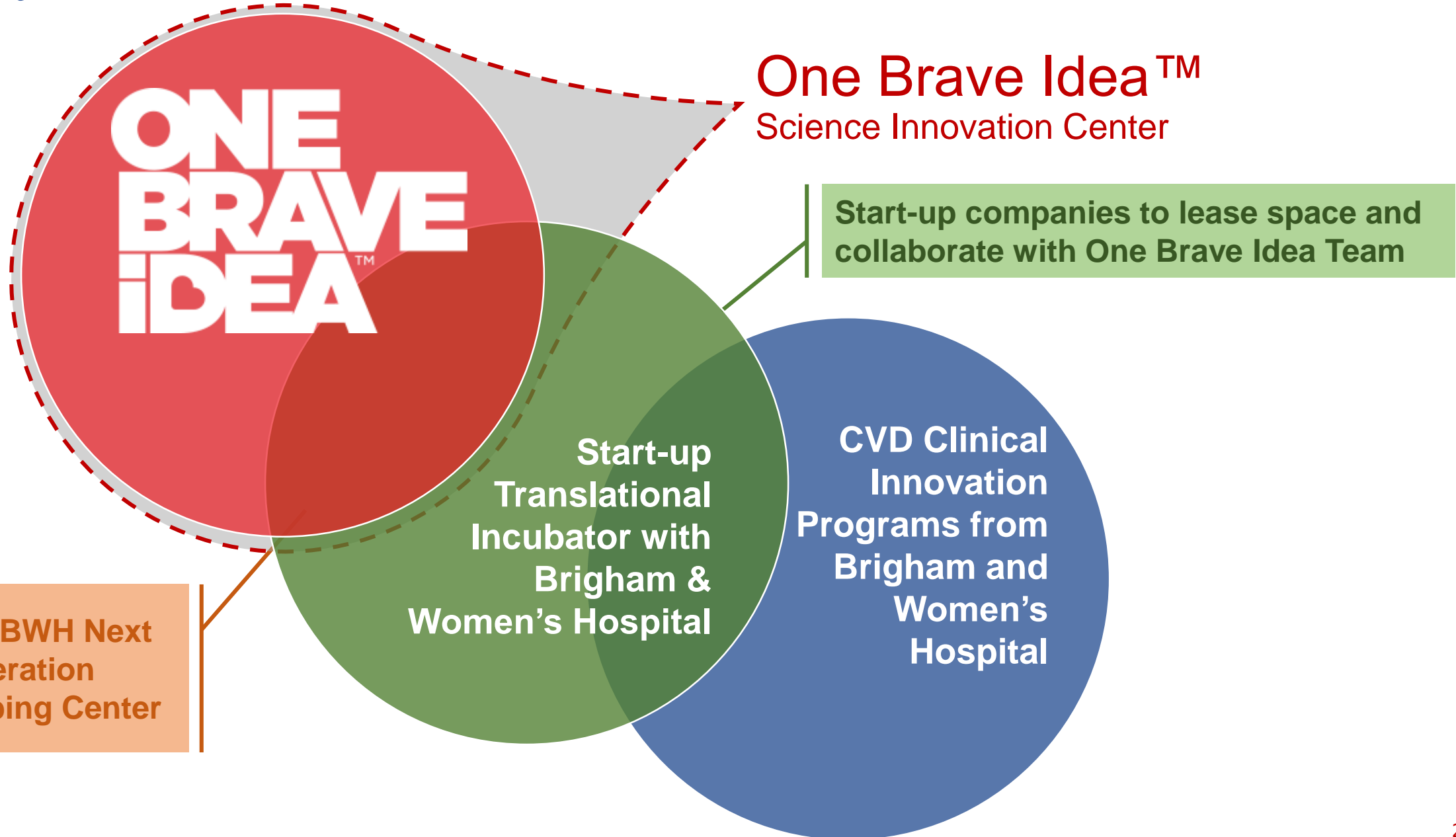
Reappraisal in new funding context

- Creating and testing approaches to:
 - Team science
 - Data accessibility
 - Science process-planning, budgeting, execution
 - Academic-private partnerships
 - Public-private partnerships
- Communication
 - Engagement across entire program: channels
 - Scientists/Funders
 - Project management
 - Regular videoconferencing: I2 based solutions
 - Shared dashboards-metrics
- Publication
- Sociology
- Economics
- Process
- Sustainability or otherwise-by design

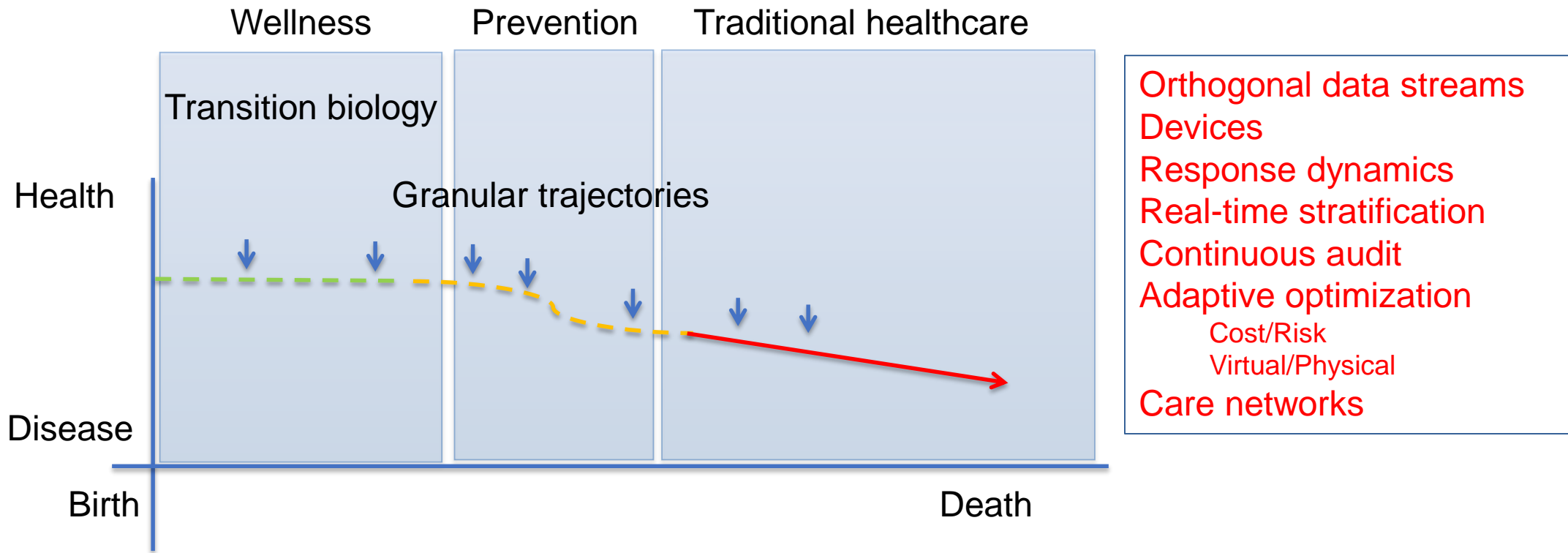
Partnership models

- Direct investment in the original project
- In-kind collaborations
- Joint investment in specific projects with shared risk/reward
 - e.g. a real world clinical trial
- Joint investment in communal projects
 - Data escrow or other strategies to overcome the long-term issues with 'de-identification'
- Joint investment in external RFAs
 - e.g. In specific population cohorts to test/validate new phenotypes
- Training and education

Physical location



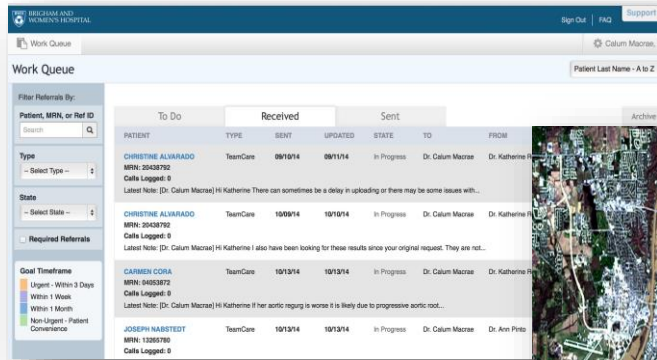
Information content drives integration and 'learning'



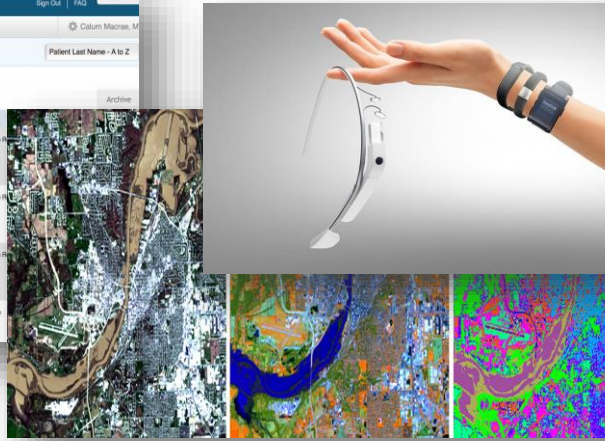
Areas of potential transformation

- Timing and resolution of care delivery, research and education
- True scalability
- Value as well as its measurement and attribution
- New partnerships across multiple areas: devices, delivery channels, etc,
- Biomedicine as learning platform: knowledge generation/ implementation
- Basic or Translational science/Hybrid trials/Real world randomization

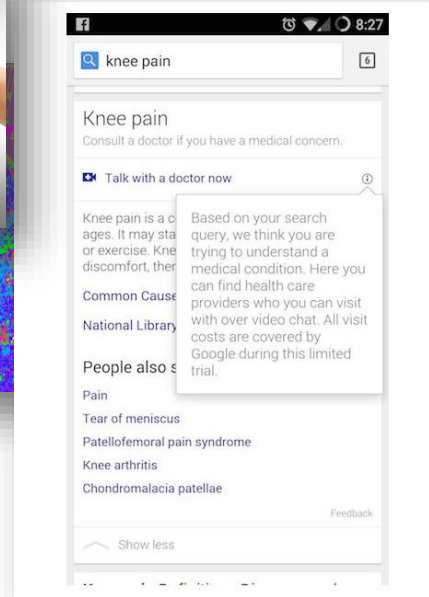
The real driver for new models of funding



New workflow



New data streams



New delivery systems

- Disruptive partners/competitors with deep pocket: PBMs, eHR companies, Global IT Players, VCs, Pharma, Banks, Real Estate, Supermarkets etc
- New revenue models
 - ACO, pay per click, subscription, added services
 - Renewed focus on knowledge generation, knowledge management and knowledge transmission
 - Different transaction types: direct to patient, educational, research premium
 - Lower cost larger markets
- Failure of many AMCs
- Emergence of small number of 'global' networks

Summary

- Information content is a core problem in biomedicine
 - Overcoming entrenched legacy phenotypes
 - Overcoming lack of comprehensiveness
 - Actively balancing broad (fewer metadata) vs deep (lower scale) phenotyping
- New models of funding are required that align all of the potential partners
 - Complement traditional biomedical science funding models
 - Directly associated with data sources
- Convergence of care and discovery
 - Data management, data science and decision support
 - Trajectories - across health and disease
 - **Funding**
- We need systematic approaches to acquiring the right information content
 - New earlier, orthogonal and more granular data and new data gathering tools
 - Eliminating non-biological silos
 - Structured perturbations-translatable by design to models-to allow integration
- Quantitative learning health systems-generalizable rules

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