

*Complex Adaptive Systems Forum:
Transformative CAS Initiatives in Biomedicine*

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Creating CAS-Based Knowledge Networks to Enable Molecularly-Based Medicine

- State of Precision(Personalized Medicine)
- The National Biomarker Development Alliance(NBDA)
- A 21st Century *In Silico* Medicine Consortium

Increasing Chronic Disease Burden: Looming Healthcare Crisis

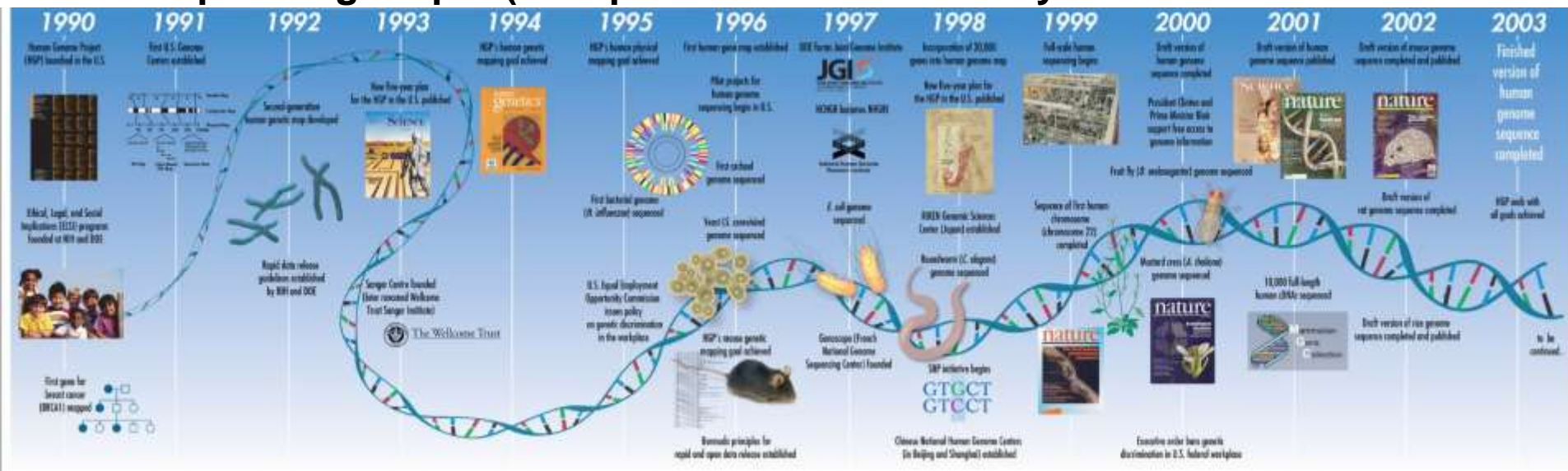
<u>Disease</u>	<u>Current Burden</u>	<u>Projected Cost</u>	<u>Future Estimates</u>
<i>Alzheimer's</i>	<i>*5.4 million survivors (6th leading cause of death)</i>	<i>\$130 billion/year</i>	<i>~16 million (Mid-century)</i>
<i>Cancer ** cases/year</i>	<i>12 million survivors (1.5 million new cases/year) (2nd leading cause of death)</i>	<i>\$263 billion/year</i>	<i>~ 2.0 million new (30% in 20 Years)</i>
<i>Diabetes ***</i>	<i>25.8 million (18.8 M Diagnosed/ 7 M Undiagnosed) (7th leading cause of death)</i>		

** Alzheimer's Association; ** ACS; *** Diabetes Association*

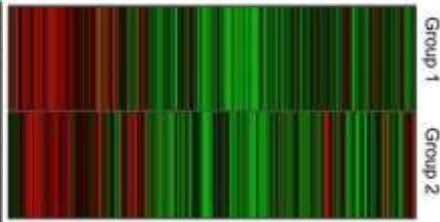
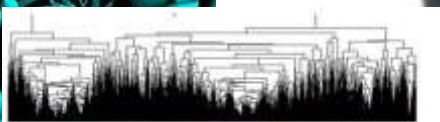
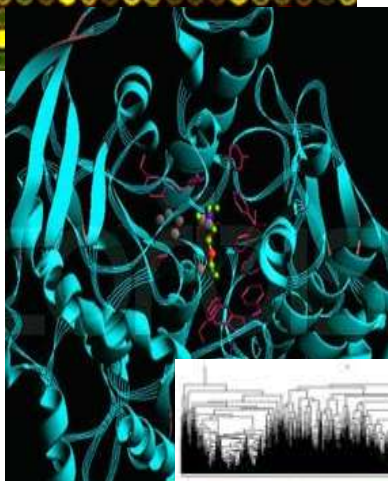
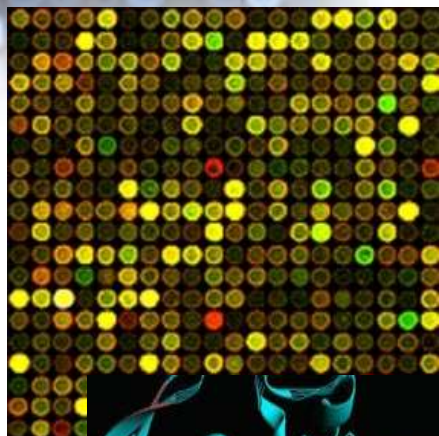
Genomics: Establishing a Foundation for Precision (Personalized) Medicine

Vision: understanding genomic changes would enable targeted intervention

- **Since 2003:** Sequencing costs fell from ~\$ 1.0 million in early 2001 to less than \$10,000 per genome (\$1000/genome possible); Numbers of genomes sequenced 2011 --~ 30,000(NHGRI); Number of genomes projected end of 2014 - 1.0 million (NGRI); total cumulative global sequencing output (on a path to reach ~ 28PB by late 2014)



Genome Sequencing ++ =,Data “Tsunami” = Paralysis



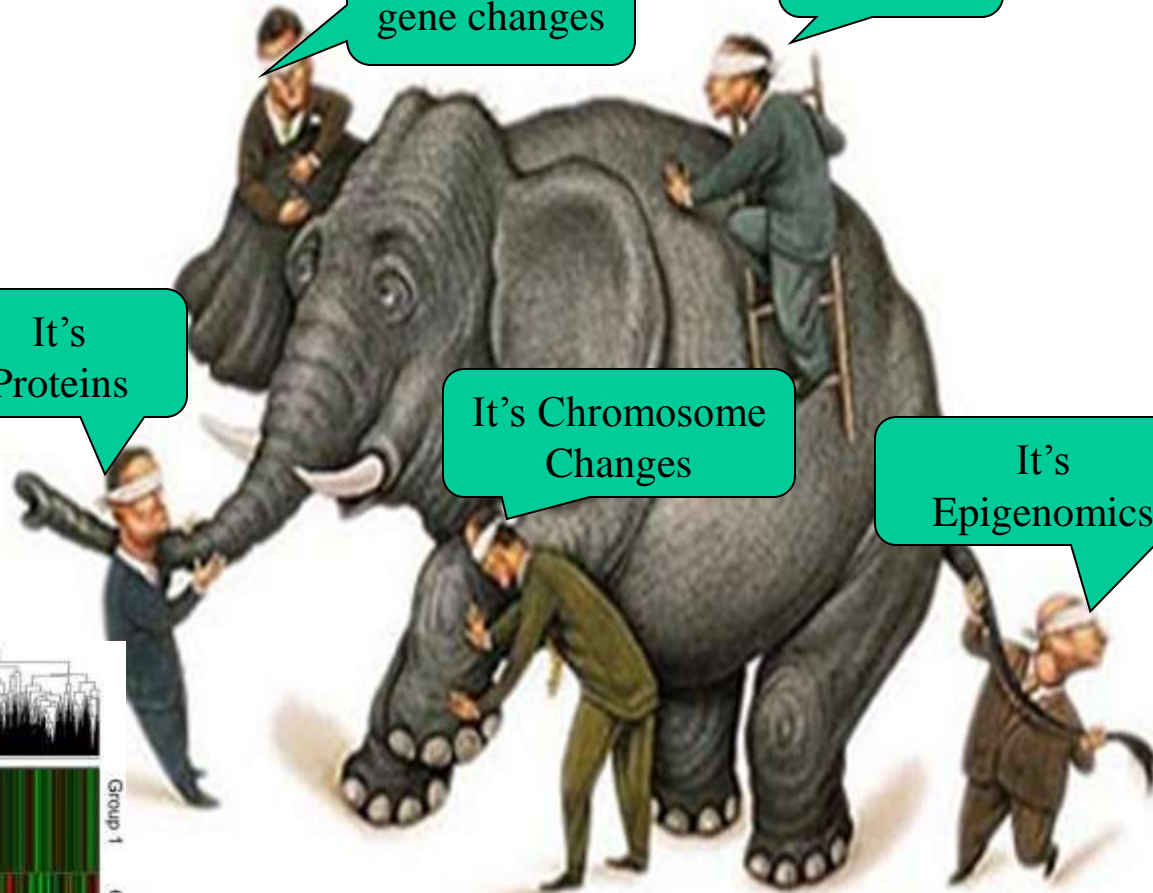
It's Proteins

It's single gene changes

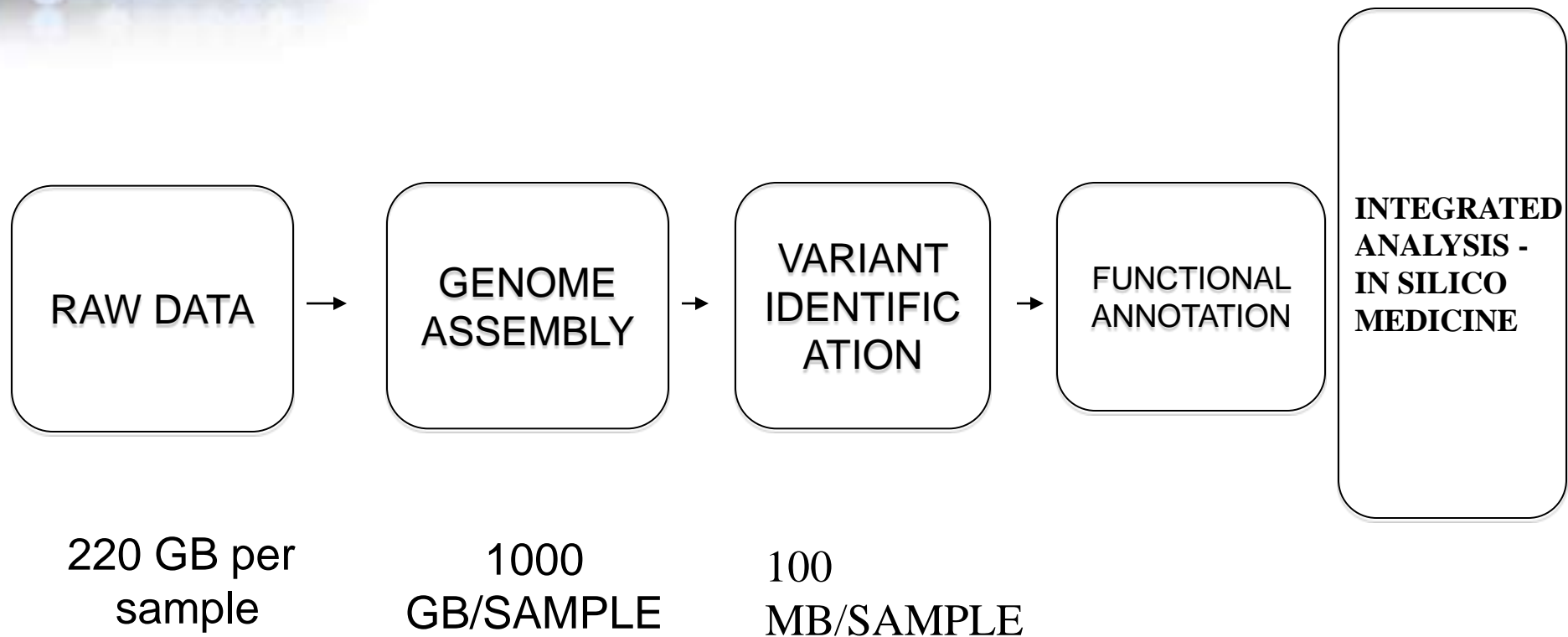
It's RNA

It's Chromosome Changes

It's Epigenomics



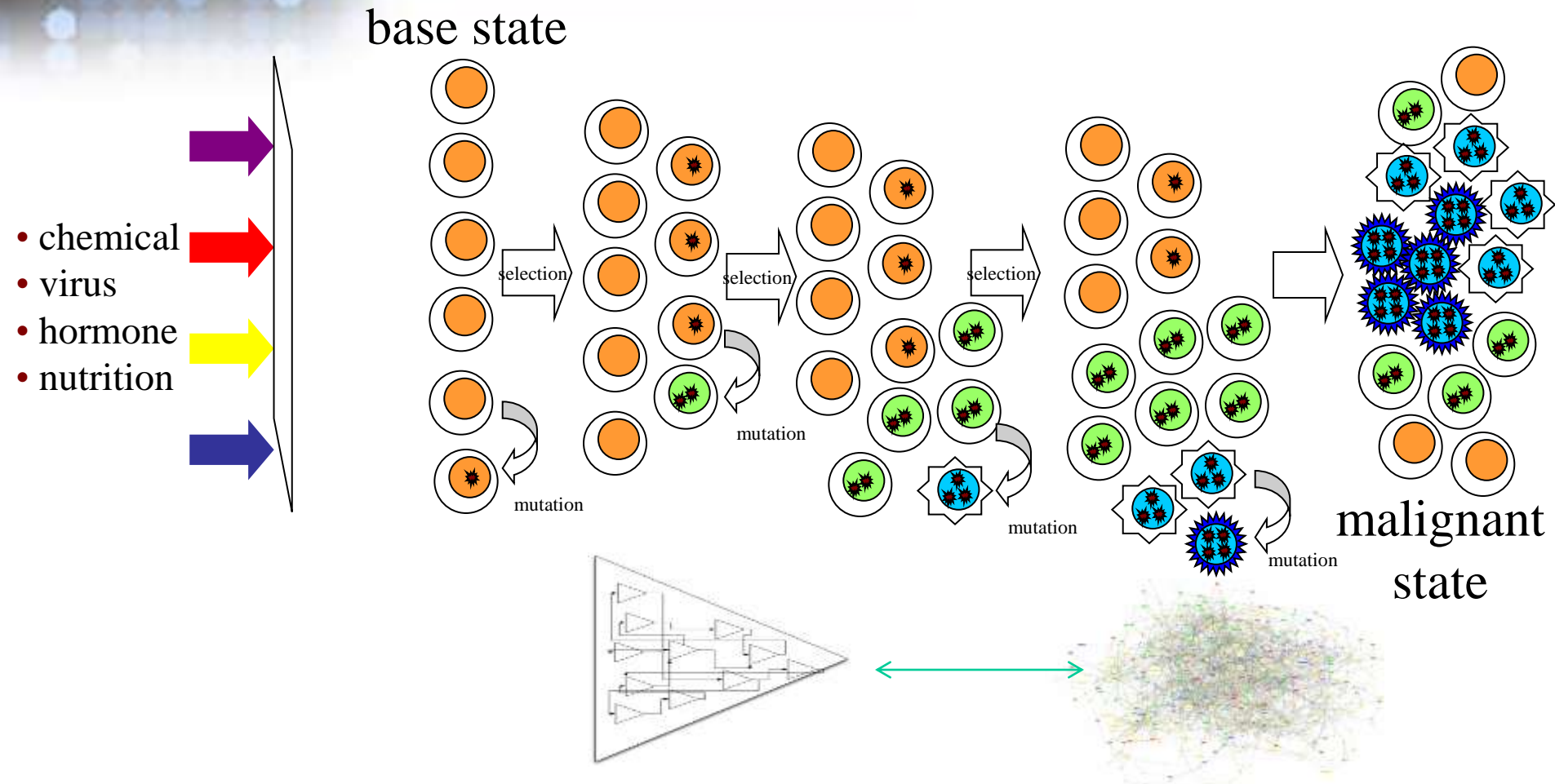
The Genomics Pipeline Today – from Raw Data to 1st Order Function



Storage, Management and Transfer

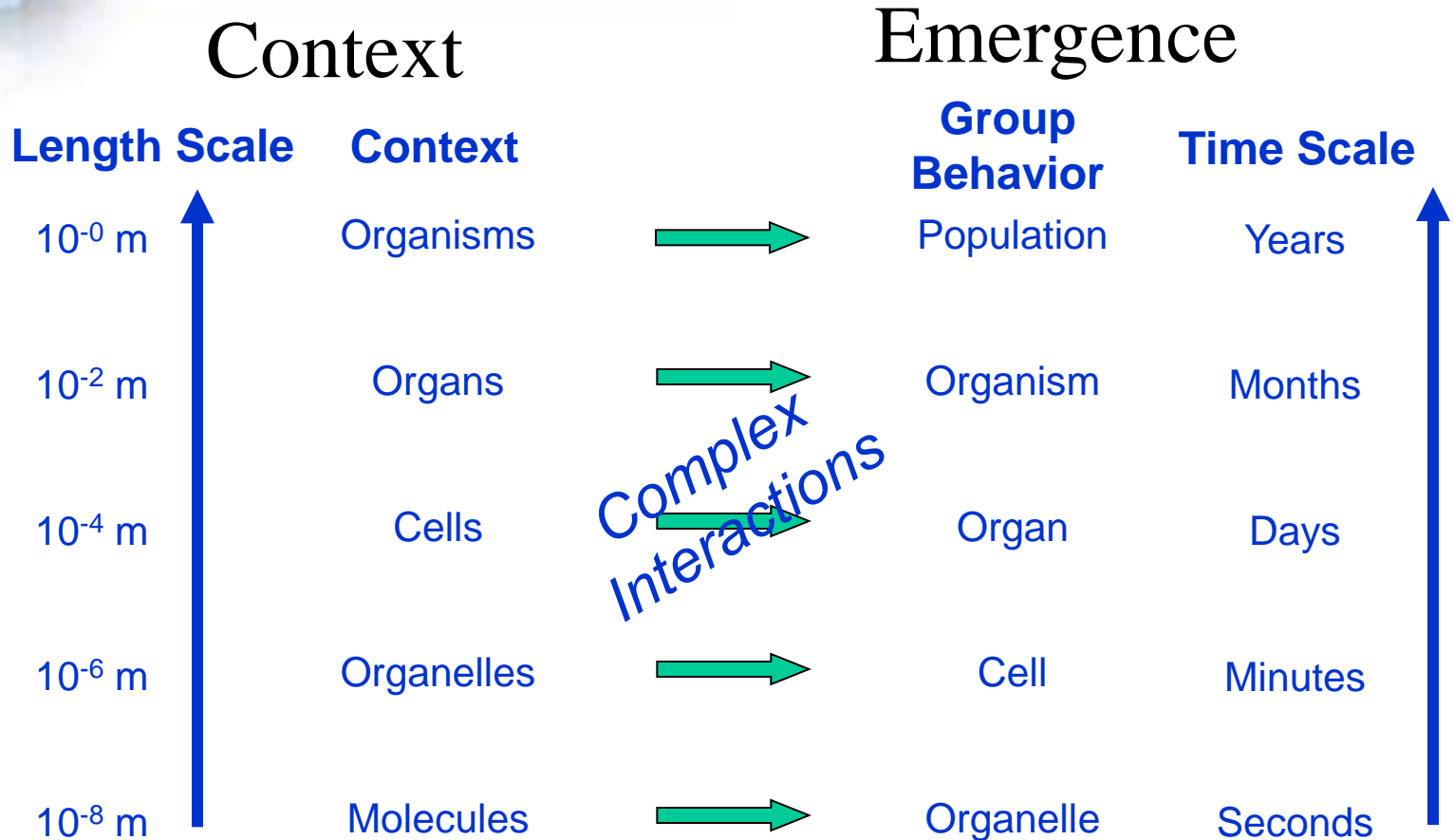
Data Compression

Cancer is a Complex Evolving System (composed of multiple subsystems)



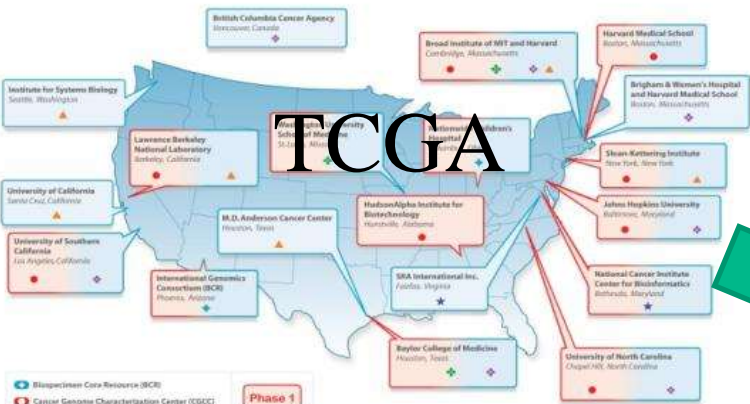
*Co-Evolution of Information-Driven Communication Between
Cancer Cells/networks and their Environment (In Context) Across Scales*

Complex Diseases (e.g. Cancer) are CAS – All Share 2 Features



Genomic alterations (information) drive the development of new information that enables the emergence of many diseases – e.g., cancer is an emergent property of the interactions of elements/subsystems across scales

Big DATA (Cancer is the Prime Example)



TCGA

Data - Data
Data - Data
Data - Data
(Issues: Public and Private - Quality, Access, Consents, Etc.)

Pharma/Bio

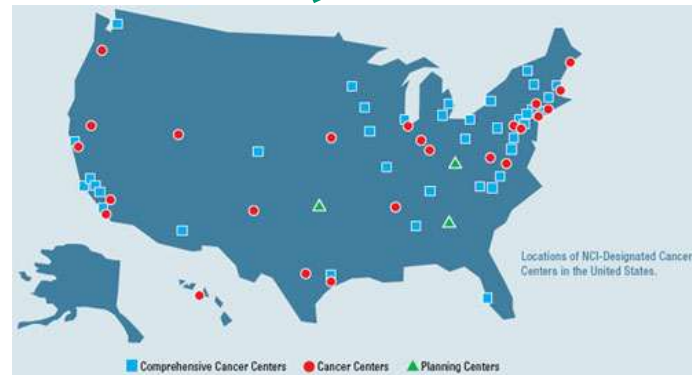


- Sanger Institute
- ICGC
- International Sequencing Projects

Individual investigators

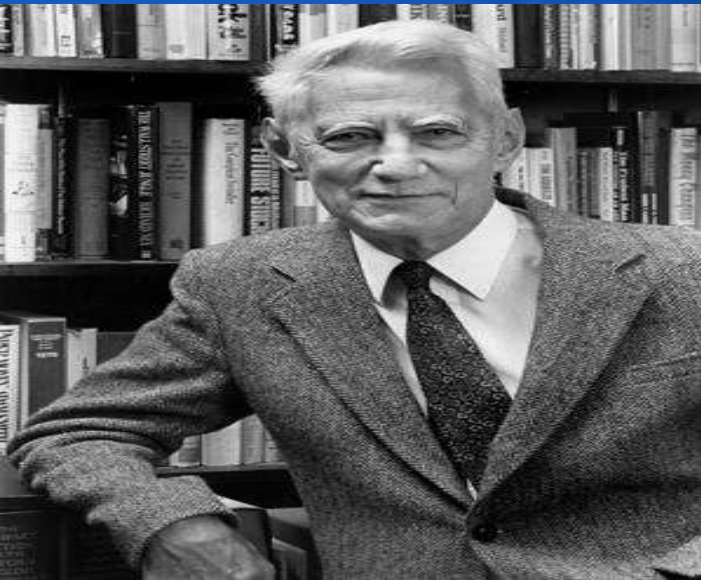


Hospitals



NCI-Designated Cancer Centers – Academic Medical Centers

The CAS-Based Diseases will Require Theoretical Constructs that Utilize Information and Evolutionary Theory to Interrogate Multi-Dimensional Data



Claude Shannon

Information
Theory



Evolutionary
Theory



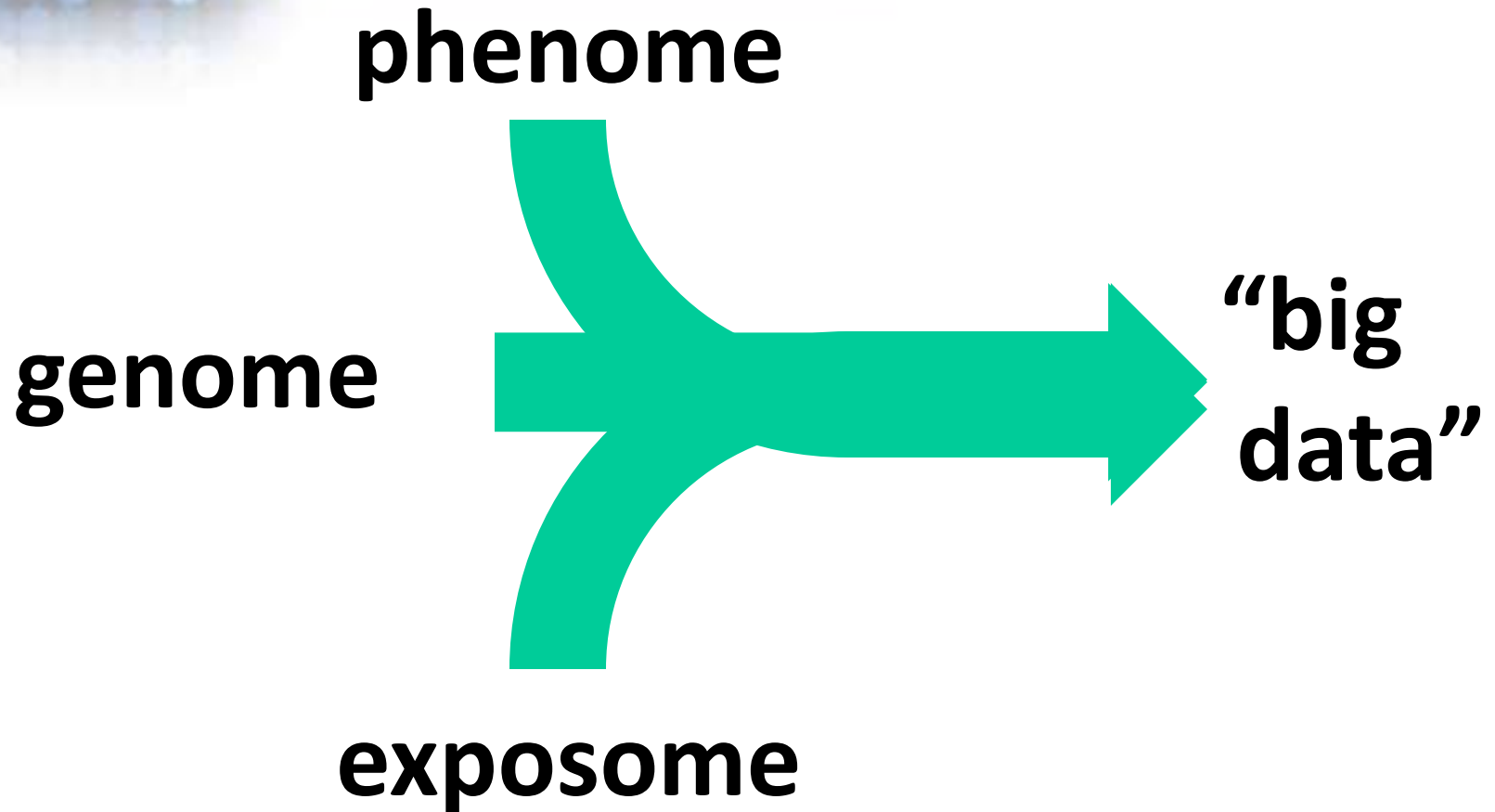
Charles Darwin

Information Theory -mathematical basis for describing/measuring information Management/decision making in cancer (e.g., can defining changes in fitness(implies changes in information content) be used to predict future disease states ?

Evolutionary Theory - Perhaps our best hope to predict evolution of resistant states in cancer (Prediction of most probable states)

“the Mind can not possibly grasp ...the full effects of many slight variations, accumulated during an almost infinite number of generations – C. Darwin

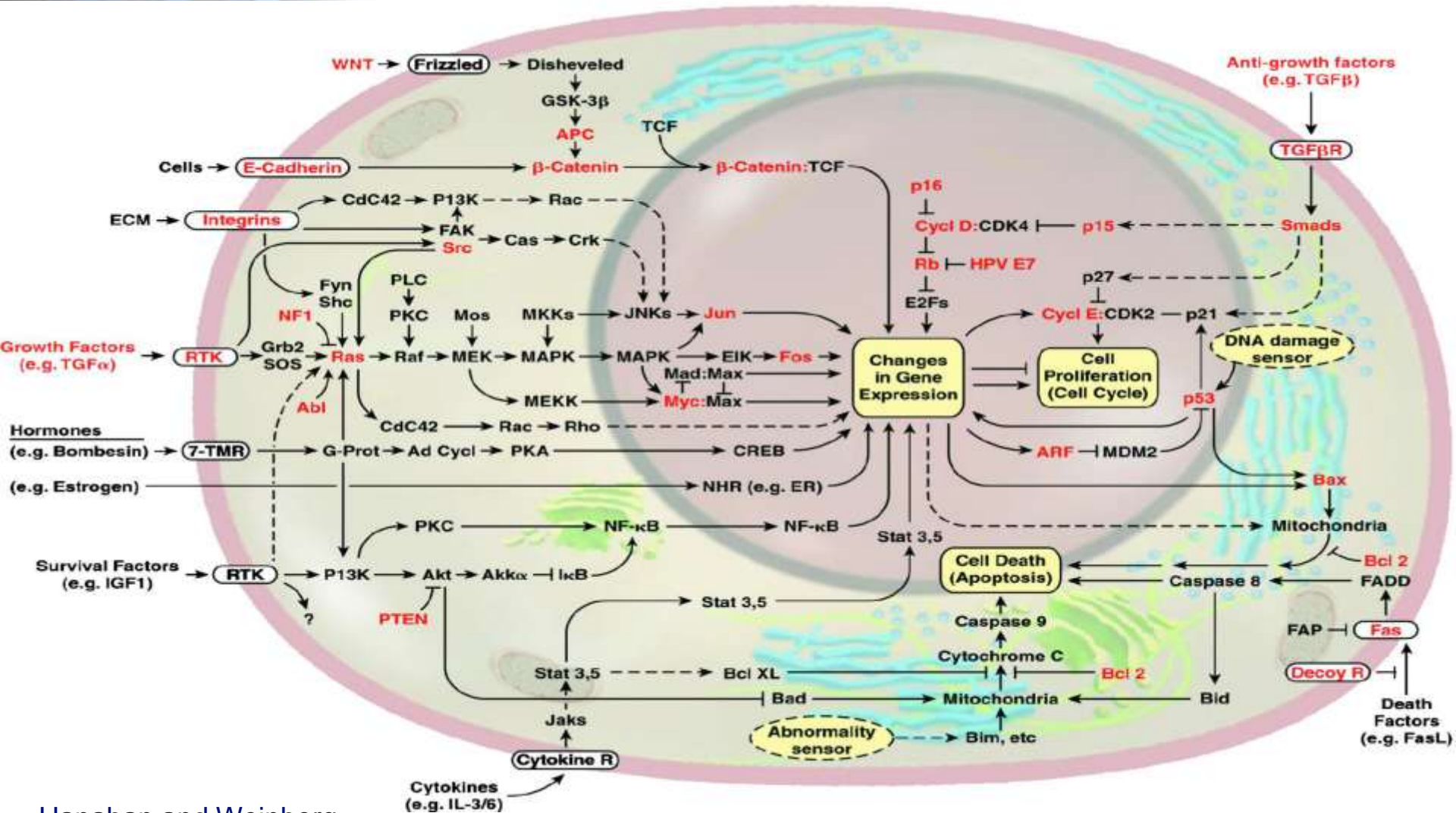
The *In Silico* Medicine: Integration – and Analysis of Multi-Dimensional – “BIG DATA”



***In silico* Medicine Consortium Status**

- **Discussions of concept and construct began ~ 1 year ago with staff from Biodesign, Mayo Clinic, Intel and Others (“Big Data Project”)**
- Goal 1 is an *in silico* center that creates a high quality continuum in the genome sequencing space that maximizes functional information and minimizes noise.
- Goal 2 is to create an integrated in silico approach that aggregates and enables “big data” from the genome, phenome and exposome
- Goal 3 is a new generation of tools for physicians that maximize disease amelioration.
- Goal 4 is knowledge network that serves as a learning system to ensure that patients benefit from the system.
- **Currently – developing partner agreements – initiating demonstration projects**
- **Seeking computational/modeling scientists from ASU**

Genomic Changes, Pathways, Networks: Early in the Discovery Process = Biomarkers

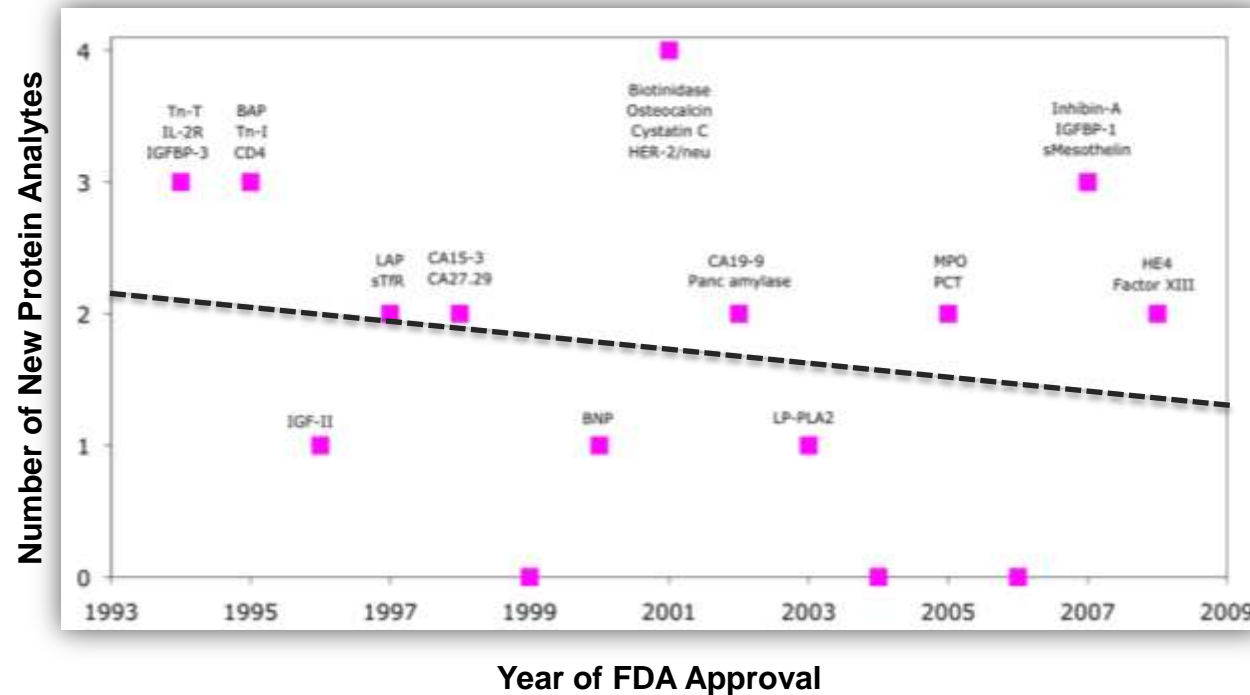


Potential of Biomarkers

- New **target discovery** (understand underlying biology)
- Drug **development** – markers of toxicity, metabolism, etc. - **pharmacogenomics**
- **Early detection** (broad or specific detection / corroboration of specific disease stage)
- Identify **molecular basis of disease phenotypes**
- **Rational** choice of **treatments (patient stratification)**
- **Assessment** of treatment **effectiveness**
- **Prognostic biomarkers**
- **Prevention markers**
- **Surrogate endpoints for clinical trials**

Status of Protein-Based Biomarkers

Few biomarker candidates translating into clinical utility and at a steadily declining rate

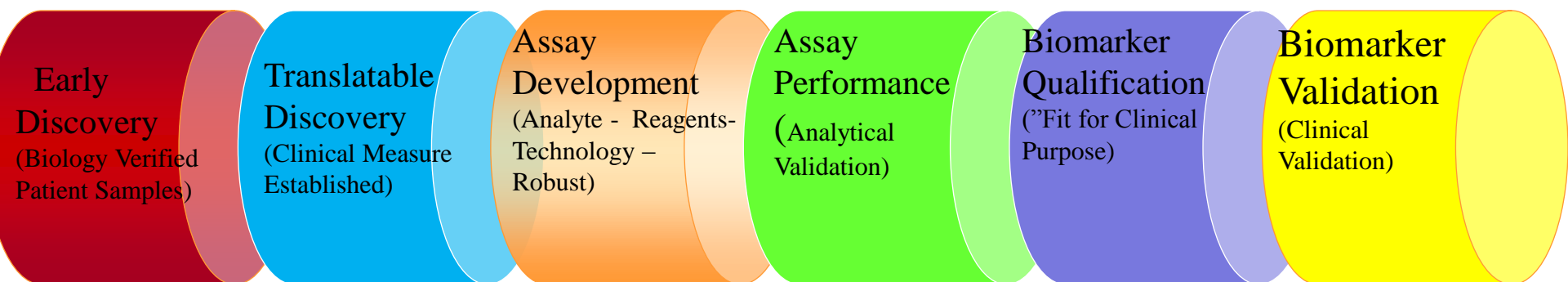


- Well over 1260 putative protein biomarkers described in the literature
- Very few approved by FDA for clinical use
- Large numbers of biomarker publications and claims – but bias and lack of standards, reproducibility across labs and SOPs make discoveries questionable
- Molecular diagnostics business remains undeveloped

Why is the Development of Biomarkers Slow and Difficult?

- Knowledge of the biological space is insufficient to understand how biomarkers reflect function – especially in populations (heterogeneity)
- Biomarkers are often developed without attention to biospecimen and technology standards
- Proteomics technologies (and others) are not standardized – often findings are not reproducible
- Biomarkers are not specifically developed for a specific function or test (i.e. FDA – fit for purpose)
- Biomarkers are not qualified before entering clinical trials
- Little understanding of what is required to develop a diagnostic test vs. using a biomarker for other uses
- Regulatory pathways not final – currently in development
- There is not end-to-end standards based system for biomarker development

NBDA: An end to End Standards Based System for Biomarker Development



NBDA: Description and Status

- **Overall Description:** Under the leadership of ASU over the last nine months, the respective strengths of ASU –especially Biodesign, C-Path, IGC (Founding Partners) and Collaborating Partners (Mayo Clinic) and (Several Partners in Discussion) have converged to create the NBDA
- **Goals:** The NBDA will transition to become a 501C3 with an overall goal of developing the evidentiary standards needed for molecularly based and other biomarkers (agnostic in terms of diseases and classes of biomarkers) at each stage of the biomarker pipeline to ensure a predictable prospective path to clinical development and regulatory filing
- **Status:** NBDA planning (and strategic plan completed; start up funds raised; workshop to begin standards identification and process held December 13-14, 2012; preliminary discussions with several pharmaceutical and diagnostic partners; discussions with FDA, NIH National Center for Advancing Translational Research, etc.)

NBDA Future Plans

- NBDA is underway – official launch with aggressive communications strategy in March 2013
- Case studies of approved biomarkers finalized to be submitted for publication 1st quarter 2013
- Overall development process- preliminary plan following Decembere workshop
- Initial laboratory study completed – mid 2013 (DNA, expression, epigenomic results in fresh tissue vs. paraffin fixed patient materials)
- Precompetitive collaborative consortia with industry established – 2013
- Additional content focused workshops in 2013
- Additional fund raising underway

NBDA is the “Poster-Child” for use-inspired Research to enable creation of a rational system for the development of precision medicine. Lots of opportunities for collaboration