

Disclosure

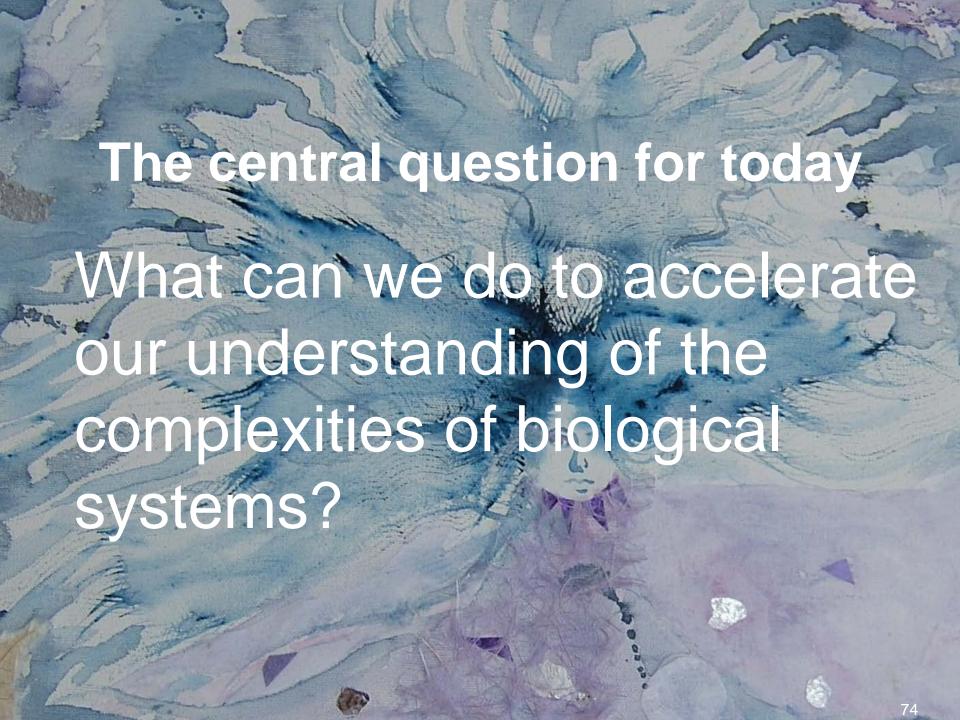


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- The authors of this research have <u>no</u> financial or other interests which pose <u>conflicts of interest</u>. Licenses to the Vanderbilt pump and valve technologies have been issued to KIYATEC, Inc. and CN Bio Innovations, which has also licensed the MicroFormulator. Our MicroClinical Analyzer patents have been licensed to Agilent. John Wikswo is an Inaugural Member of the Scientific Advisory Board of BiOasis Technologies, Inc.
- The views expressed in this document are solely those of the authors and do not necessarily reflect those of any of the funding agencies or companies. The EPA does not endorse any products or commercial services mentioned.

Abstract



Deconvolving the multiscale, spatiotemporal complexity of biology requires not only understanding the governing laws of physics and chemistry, but also decoding billions of years of genetically encoded history. Simple, passive observations cannot expose the nested, redundant levels of regulation in the historical instruction set, and active interventions are needed to disable specific biological functions to expose others. The concept of the hermeneutic circle applies to biology - one cannot understand the whole until one understands the parts, and one cannot understand the parts without understanding the whole. In this context, coupled microphysiological systems meet the criteria for a successful toy model: complicated enough to recapitulate key regulatory processes but simple enough to understand. Revealing the functions of such coupled in vitro systems will require untargeted analysis of the genome, transcriptome, proteome, lipidome, interactome, and metabolome at the level of cells and tissues, which places demands on the accessibility and interconnection of each micro-organ and establishes lower limits on their sizes. The grand challenge is to devise and integrate the requisite cells, microfluidic bioreactors, sensors, analytical techniques, closed-loop controls, and mathematical models (which may be underspecified). Machine learning and automated design of experiments may be critical to closing the hermeneutic circle.

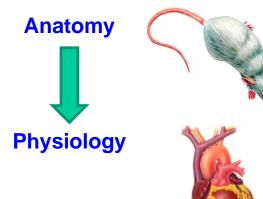


Seven Themes

- 1. A brief history of biology
- 2. Just how complex is biology?
- 3. What is the role of physics in understanding the complexity of biology?
- 4. Why do we need to emphasize external control?
- 5. How might Organoids and Organs on Chips change the way we study biology?
- 6. What does Multi-Omics offer?
- 7. Closing the circle

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The history of biology...

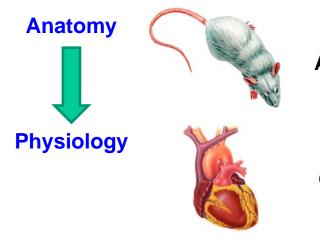




The birth of physiology



Galen of Pergamon AD 129-216

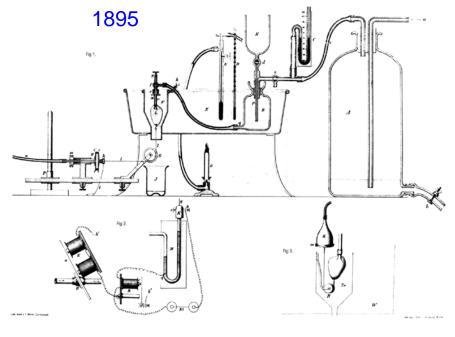


Animal

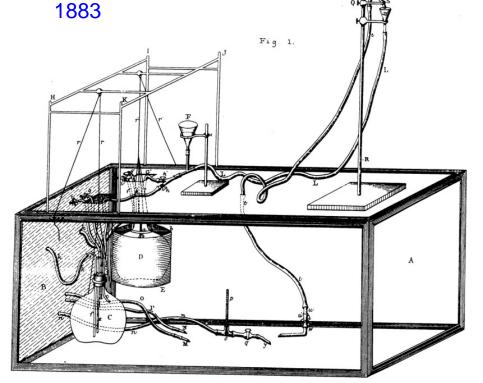
Isolated Organs





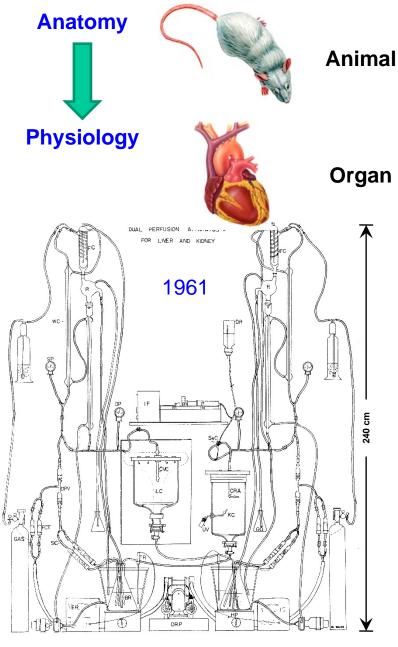


O. Langendorff. Untersuchungen am Überlebenden Säugethierherzen. Pflug.Arch.Eur.J.Phy. 61 (6):291-332, 1895, as shown in H.G. Zimmer. The isolated perfused heart and its pioneers. News Physiol.Sci. 13:203-210, 1998.



Martin HN. The direct influence of gradual variations of temperature upon the rate of beat of the dog's heart. Philos Trans R Soc London 1883 Jan 1:174:663-88.

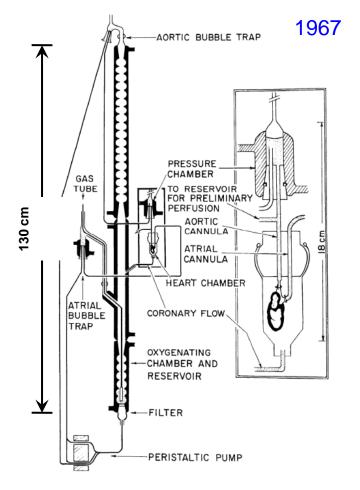
Conclusion: Body temperature affects heart rate. Missing from the image: Dog with cannulated heart & lungs, cast iron, water-filled pan, and Bunsen burners...



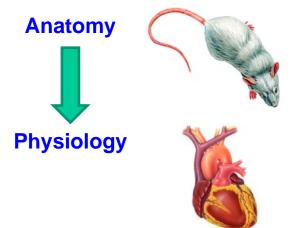
Avis FR. Investigations of liver and kidney. Design of a dual apparatus for research study. Science Teacher 1961;28(1):14-8.

Isolated Organs





- Neely JR, Liebermeister H, Battersby EJ, Morgan HE. Effect of pressure development on oxygen consumption by isolated rat heart. Am J Physiol 1967 Apr 1;212(4):804-14.
- Neely JR, Liebermeister H, Morgan HE. Effect of pressure development on membrane transport of glucose in isolated rat heart. Am J Physiol 1967;212(4):815-22.



Animal

Isolated Organs

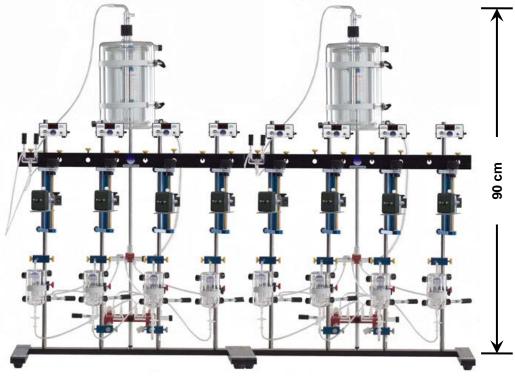


Organ

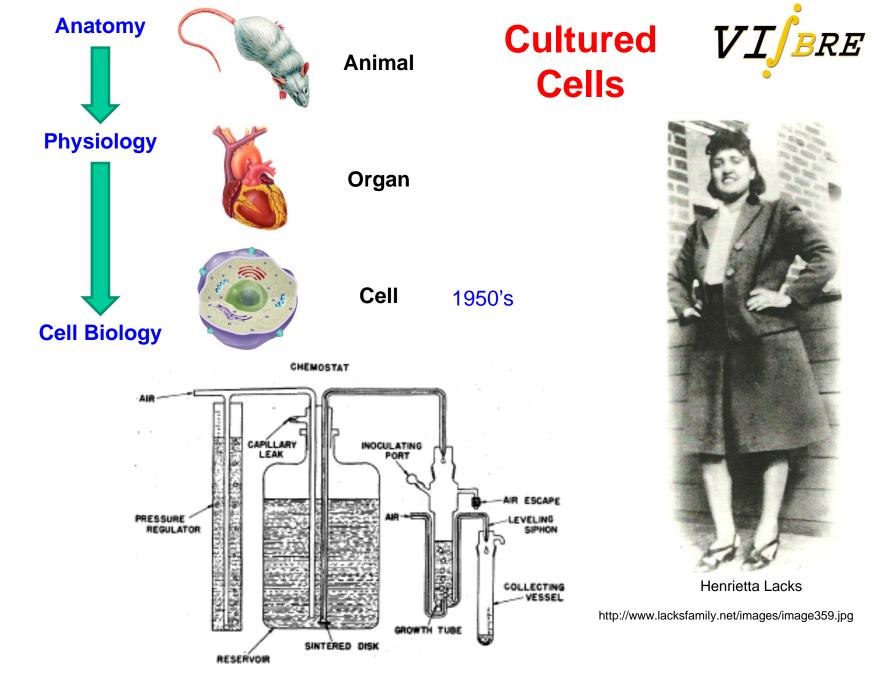
Today



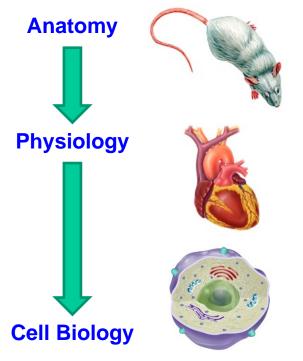
Holzer JR, Fong LE, Sidorov VY, Wikswo JP and Baudenbacher FJ. High resolution magnetic images of planar wave fronts reveal bidomain properties of cardiac tissue. Biophys.J. 87 (6):4326-4332, 2004.



Courtesy of Desmond Radnoti of Radnoti LLC
In regular use by academics and pharma



- Novick A and Szilard L. Experiments with the Chemostat on Spontaneous Mutations of Bacteria. PNAS 36 (12):708-719, 1950.
- Aaron. Novick and Leo Szilard. Description of the Chemostat. Science 112 (2920):715-716, 1950.



Animal

Cultured **Cells**

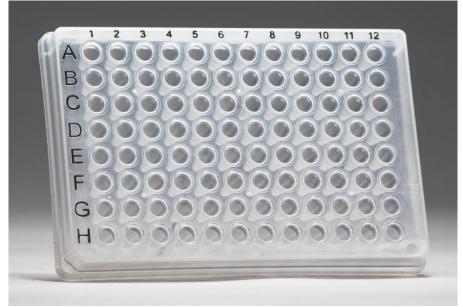


Organ

glasslaboratory.com/files/2245127/uploaded/GL-P100%20Petri%20Dish.jpg

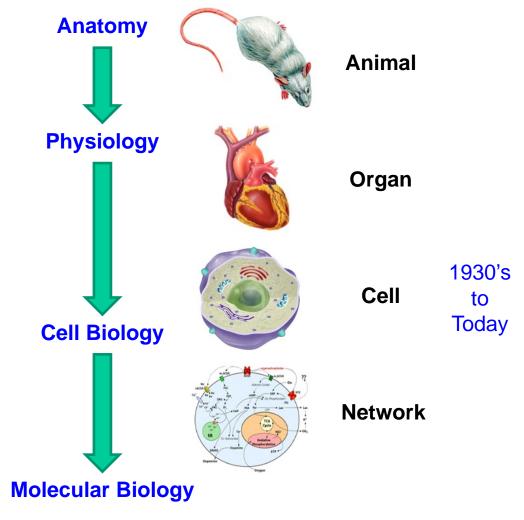
Cell

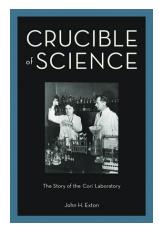
Today





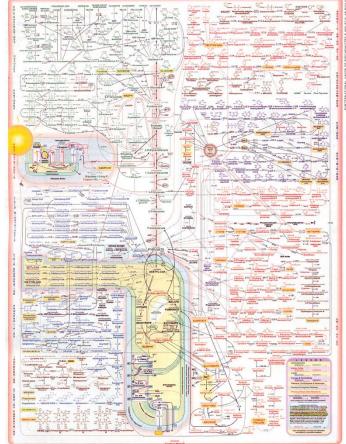


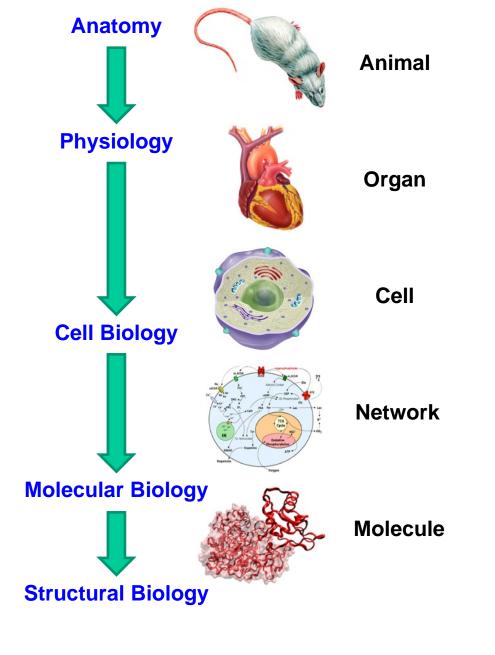






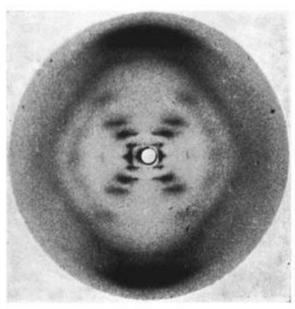
Crucible of Science: The Story of the Cori Laboratory, John H. Exton, Oxford, 2013





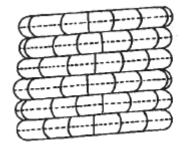


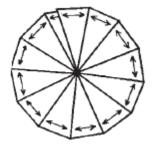
1952



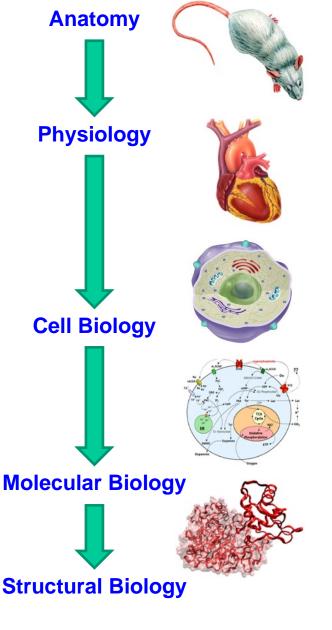
http://www.insight.mrc.ac.uk/2013/04/25/behind-the-picture-photo-51/

1955





Rosalind E. Franklin. Structure of Tobacco Mosaic Virus. Nature 175 (4452):379-381, 1955.



Animal

Automated high throughput 1 screening

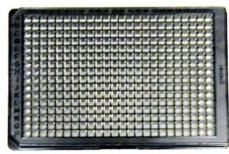
Organ

Today

1536 Well ~8 μl

Cell





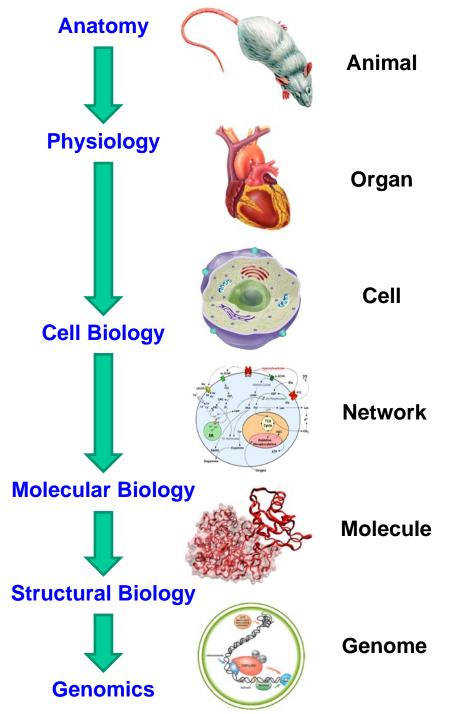
384 Well ~40 μl

Molecule





Images courtesy of Dr. David Weaver, Vanderbilt VICB High-Throughput Screening Facility



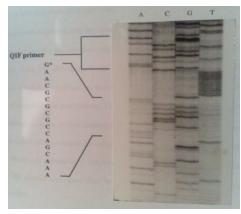
The Human VI BRE Genome Sequenced



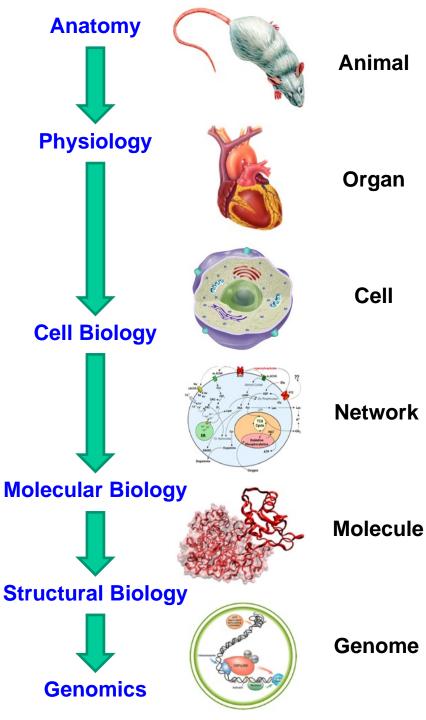
2003



http://www.wired.com/2008/07/british-institu/



http://nsaunders.wordpress.com/2011/12/22/sequencingfor-relics-from-the-sanger-era-part-1-getting-the-raw-data/



Cell

The \$1,000 Genome



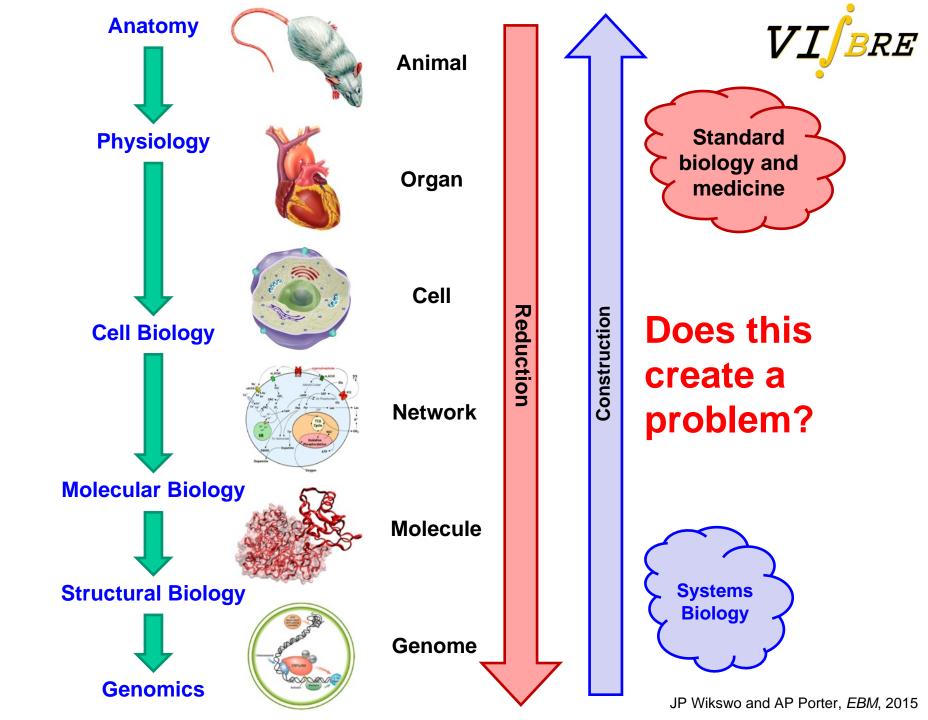
Today



1000 Gb max output, 4000M max read number, 2x125bp max read length, 600 gigabases (Gb) per day per system http://systems.illumina.com/systems/hiseq_2500_1500.ilmn



Conclusion: Technology drives major advances in biology!



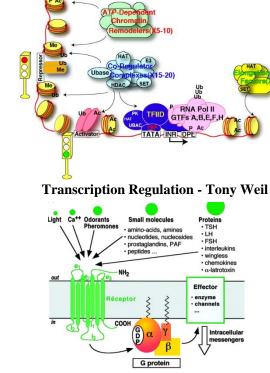
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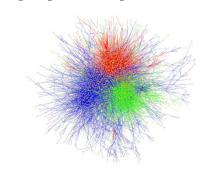
Why is biology so complex?

- The human genome codes for ~20,000 unique proteins
- RNA splicing and proteolysis may result in 50,000 to 500,000 unique proteins.
- Post-translational modifications may increase this to a million
- An individual cell expresses between 10,000 to 15,000 different proteins at any one time.
- Proteins interact with each other.





G-coupled protein receptors – Heidi Hamm

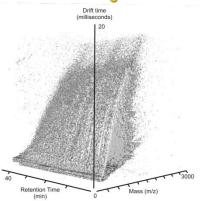


Human proteome, and its binding interactions - Simonis and Vidal.

Why is biology so complex, con't?

- Today, one can easily detect 100,000 chemical species in 100 µL of rat serum.
- Cells are NOT well-stirred bioreactors but have anomalous diffusion and active transport.
- 10⁹ 10¹¹ interacting cells in some organs.
- We must consider the microbiome.
- Cell signaling is dynamic, non-linear, multiscale, redundant, and has positive and negative feedback.

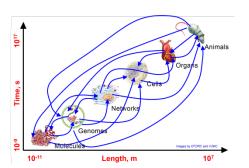




UPLC-nESI-IM-MS John McLean



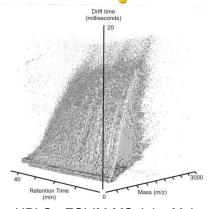
3.1 x 3.2 x 1.2 µm³ beta cell Brad Marsh, PNAS, 2001



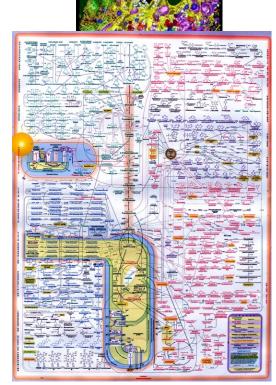
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- We must consider the microbiome.
- Cell signaling is dynamic, non-linear, multiscale, redundant, and has positive and negative feedback.
- Metabolism has 8,000 reactions.
- Models might need Avogadro's number of PDEs, *i.e.*, a Leibniz of PDEs (1 L = N_a).
- We need new experimental approaches.





UPLC-nESI-IM-MS John McLean



Biology spans lots of space and time

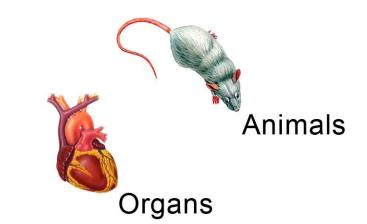


1017

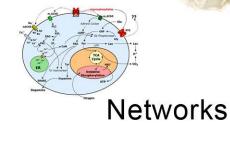
Time, s

6-O





Cells



Genomes

Molecules

Images by CFDRC and VUMC

10-11

Length, m

10⁷

Part of the problem is that human biology is COMPLEX. Organs talk to each other, but we seldom hear what they are saying.

Organs, Organs, Organs

Cardiovascular

Heart **Blood Blood vessels** Digestive Salivary glands **Esophagus** Stomach Liver Gallbladder **Pancreas**

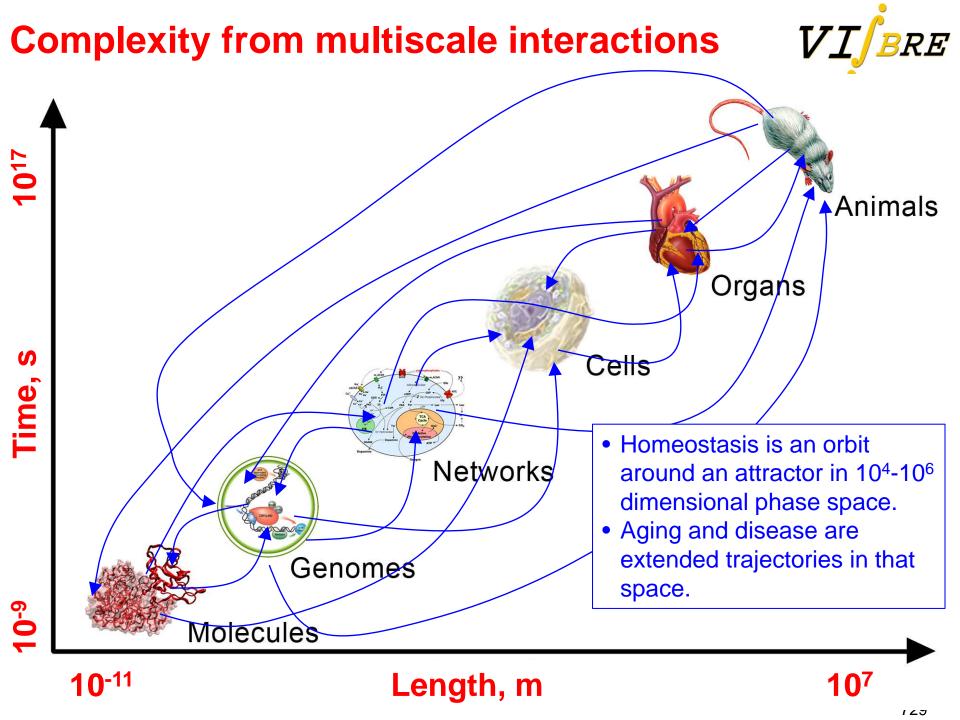
Excretory **Kidneys Ureters** Bladder Urethra Immune Leukocytes **Tonsils** Adenoids **Thymus**

Reproductive **Ovaries** Fallopian tubes Uterus Vagina Mammary glands **Testes** Vas deferens Seminal vesicles **Prostate**

Intestines Colon Rectum Anus **Endocrine** Hypothalamus Pituitary gland Pineal gland **Thyroid Parathyroids** Adrenals

Spleen **Appendix** Integumentary Skin Hair **Nails** Muscular Muscles Golgi tendon organ Nervous Brain Spinal cord Nerves Eyes

Penis Respiratory Pharynx larynx **Trachea Bronchi** Lungs Diaphragm Skeletal Bones Cartilage Ligaments **Tendons**



A possible failure mode



Ontological failure: The phenomenon you are interested in requires elements or laws outside of the set you have been given.

D. Bray. Reductionism for biochemists: how to survive the protein jungle. *Trends Biochem.Sci.* 22 (9):325-326, 1997.



The solution to ontological failure

Get more data...



It's the numbers....

Where do we get a mole of numbers?

The Catch



- Modeling of a <u>single</u> mammalian cell may require >100,000 <u>dynamic</u> variables and equations, maybe > 1,000,000
- Cell-cell interactions are critical to system function
- 10⁹ 10¹¹ interacting cells in some organs
- Cell signaling involves highly DYNAMIC biochemical cascades with positive and negative feedback
- Multiple, overlapping regulatory mechanisms
- Many of the interactions are nonlinear
- Models might have a Leibnitz (1 L = N_a) of PDEs
- The data don't yet exist to drive the models
- Hence we need to experiment...

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Helmholtz on Cells



"The behavior of living cells should be accountable in terms of motions of molecules acting under certain fixed force laws."

Herman von Helmholtz, 1870

Quoted in Max Delbruck, "A Physicist looks at biology" Trans. Conn. Acad. Arts Sci. 38:173-190, 1949.

Bohr on Complementarity



"... we should doubtless kill an animal if we tried to carry the investigation of its organs so far that we could describe the role played by single atoms in vital functions. In every experiment on living organisms, there must remain an uncertainty as regards the physical conditions to which they are subjected, and the idea suggests itself that the minimal freedom we must allow the organism in this respect is just large enough to permit it, so to say, to hide its ultimate secrets from us."

N. Bohr. "Light and Life," *Nature* 131 (3309):457-459, 1933.

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Schrödinger on Life



"Present day physics and chemistry could not possibly account for what happens in space and time within a living organism."

"... a living organism ... can only keep ... alive ... by continually drawing from its environment negative entropy."

Schrödinger, What is Life, 1943



"When I was invited to review the influence of *What is Life?* I accepted with the intention of doing honor to Schrödinger's memory. To my disappointment, a close study of his book and of the related literature has shown me that what was true in his book was not original, and most of what was original was known not to be true even when it was written."

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What was original was from Max Delbruck.



What makes biology so different from physics, chemistry, and engineering?

What makes biology different from physics or chemistry?



Physics and chemistry describe dynamic interactions in terms of fundamental or phenomenological laws that govern the state of the matter being studied.

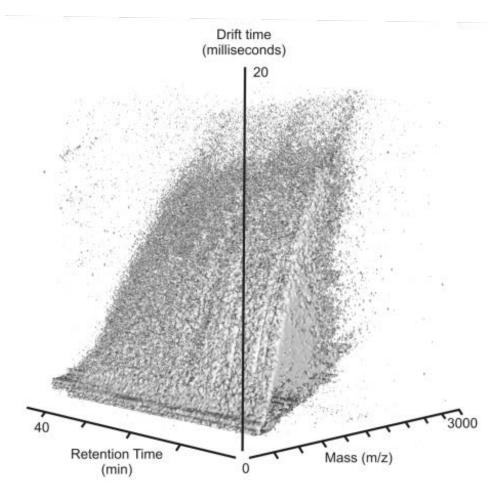
*Ohm's law, Hooke's law, the Standard Model, ... conservation of mass, Dalton's law, quantum mechanics ...

Biology has laws, but the operation of every living organism is determined not only by the laws of biology, physics and chemistry, but also by <u>historic instructions</u> that may be specific to each individual organism.

"... any living cell carries with it the experiences of a billion years of experimentation by its ancestors. You cannot expect to explain so wise an old bird in a few simple words."

Max Delbrück, "A Physicist Looks at Biology," 1949





Build untargeted mass spectrometers, and you will have enough data.

How do you deal with a Leibnitz of non-sparse PDEs involving 100,000 nonlinear variables?

Carefully, very carefully

A possible failure mode



Ontological failure: The phenomenon you are interested in requires elements or laws outside of the set you have been given.

There is a second possible failure mode

Epistemological failure: You have enough elements and the laws do apply, but you yourself cannot understand the explanation that they provide.

D. Bray. Reductionism for biochemists: how to survive the protein jungle. *Trends Biochem.Sci.* 22 (9):325-326, 1997.

Houston, we have a problem.



 The human brain can process only seven pieces of data at a time.

"...the seven-point rating scale, the seven categories for absolute judgment, the seven objects in the span of attention, and the seven digits in the span of immediate memory..."

G.A. Miller, "The Magical Number Seven, Plus or Minus Two: Some Limits on our Capacity for Processing Information," Psychological Review, 63, 81-97 (1956).

A Really Hard Problem



If the human brain were so simple that we could understand it, we would be so simple that we couldn't.

Emerson M. Pugh, 1938

Yet one more Really Hard Problem



Pugh's observation applies to biology:

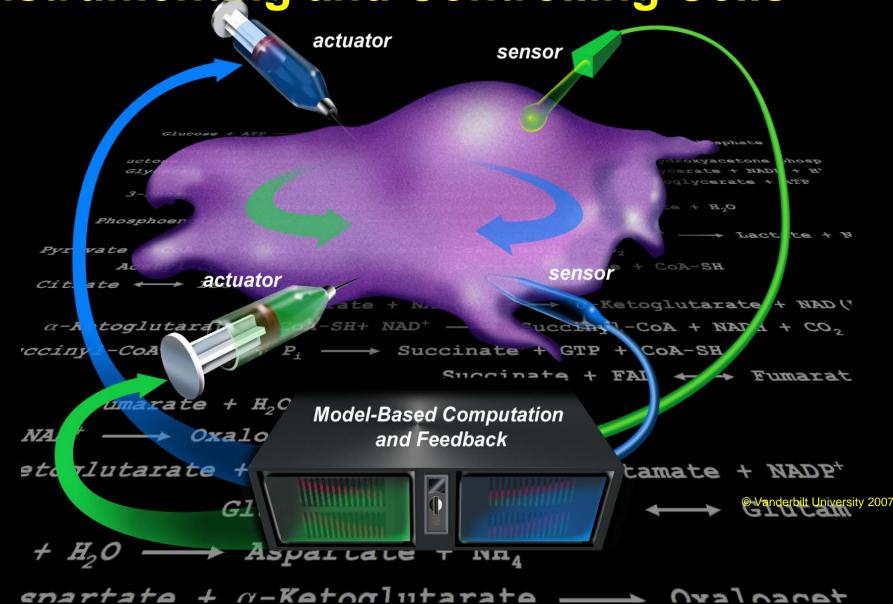
Human biology may be too complicated for humans to <u>fully</u> comprehend.

John Wikswo

Seven Themes 1. A brief history of b 2. Just how nderstanding the . Can sensors, actuators, controllers and robot scientists help address biological complexity? 5. How might Organoids and Organs on Chips change the way we study biology? What does Multi-Omics offer? '. Closing the circle

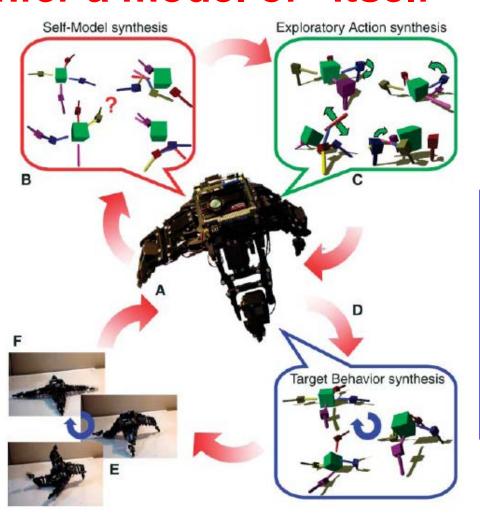
VIIBRE 2001

Instrumenting and Controlling Cells



Machine Learning: A robot that can infer a model of "itself"





A robot can be programmed to conduct experiments to derive a model of itself.

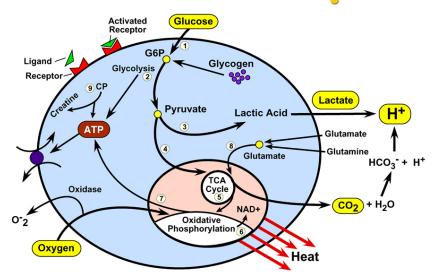
My hypothesis: Machine learning and model inference with automated experimentation can be extended from robots to bioreactors.

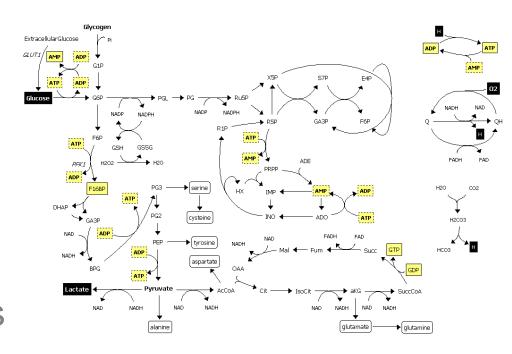
J. Bongard, V. Zykov, and H. Lipson, Resilient Machines Through Continuous Self-Modeling, Science, 314, 1118-1121, 2006

Note to self: 2006

VIJBRE

- Can I design and build a hybrid silicon/biological system that proposes and generates models and conducts experiments on itself to identify the underlying equations that govern cellular dynamics?
- Extracellular: \$3 4
 million and 3 5 years
- Intracellular: \$15 20
 million and 5 10 years



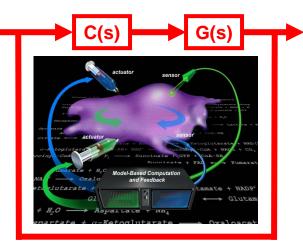




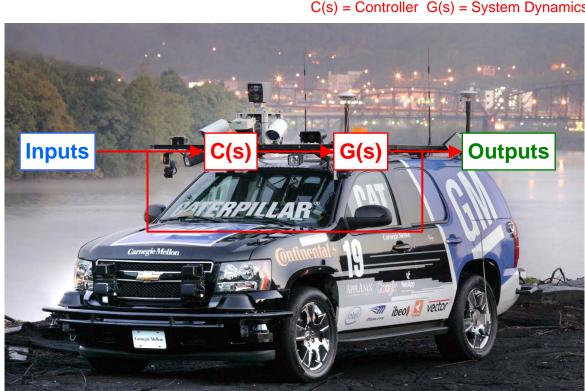
First step – control biology!

INPUT ACTUATORS

- Chemical
- Electrical
- Genetic
- Mechanical
 - Optical
 - Thermal
- Scaffolding



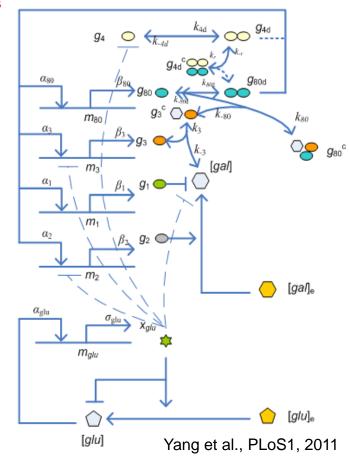
C(s) = Controller G(s) = System Dynamics

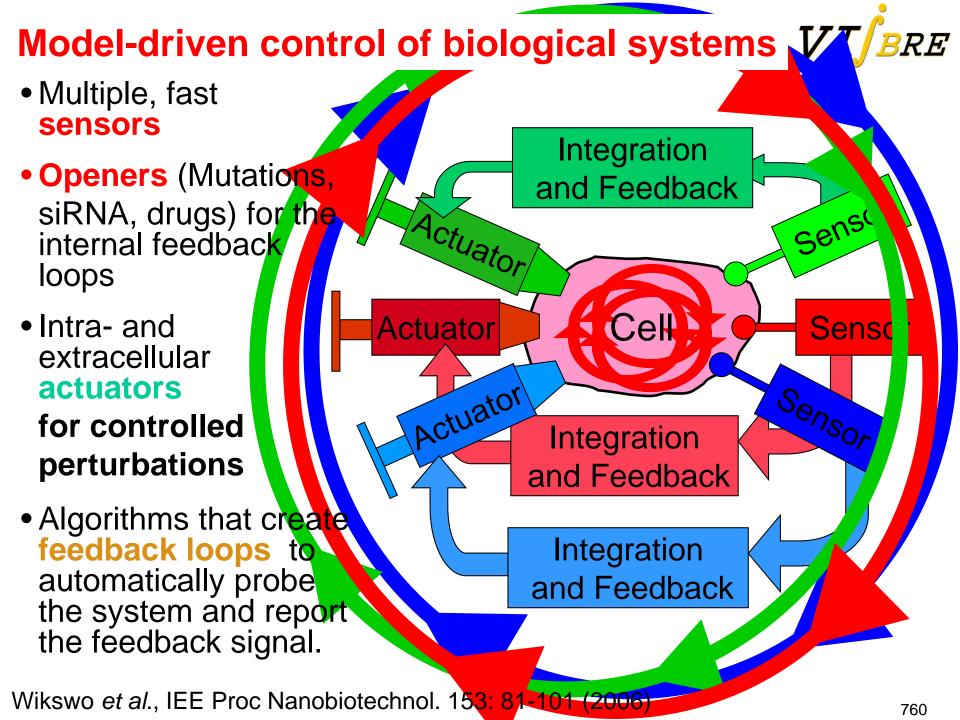


From: LeDuc, Messner, Wikswo. How do controls approaches enter into biology? Annual Reviews of BME, 2011.

OUTPUT SENSORS

- Apoptosis
- Differentiation
- Gene / Protein Expression
 - Growth
 - Metabolism
 - Motility
 - Signal Transduction





Artificial Intelligence, per Wikipedia



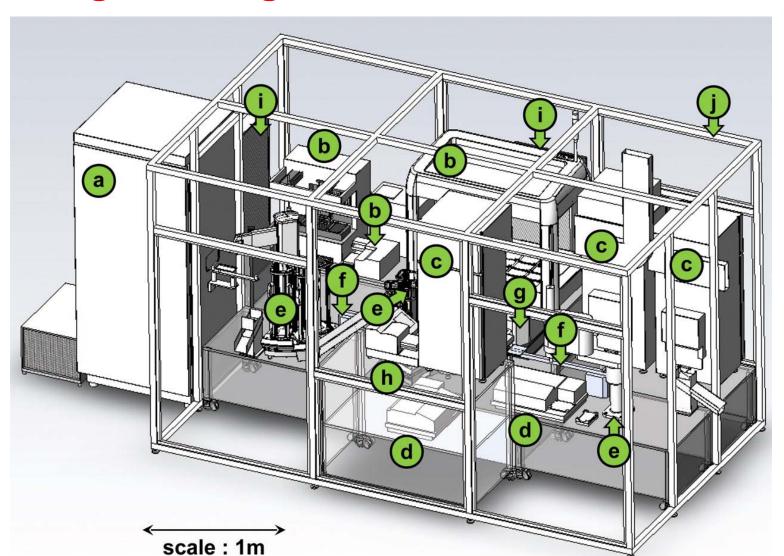
- Narrow AI (or "weak AI") is the use of software to study or accomplish specific problem solving or reasoning tasks. Weak AI, in contrast to strong AI, does not attempt to simulate the full range of human cognitive abilities. (Siri, Alexa...)
- Strong AI is artificial intelligence that matches or exceeds human intelligence — the intelligence of a machine that can successfully perform any intellectual task that a human being can. ... Strong AI is associated with traits such as consciousness, sentience, sapience and self-awareness observed in living beings. (Turing test...)

Narrow versus Strong Automated Biology



- Narrow AB: the use of software to study or accomplish specific laboratory biology tasks
 - High throughput screening (HTS)
 - Automated microscopy (ImageJ)
 - Biomedical data mining (KEGG, MAGINE, SIMONE)
 - Computer-aided medical diagnosis
- Strong AB: automated biological experiments driven by autonomous, machine-learning systems
 - Identification of orphan gene function (King et al.)
 - Inference of metabolic networks (Schmidt et al.)
 - Optimal design of experiments
 - Development of iPSC differentiation control paradigms

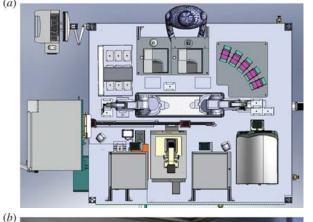
Strong AB: King's Robot Scientist "Adam"



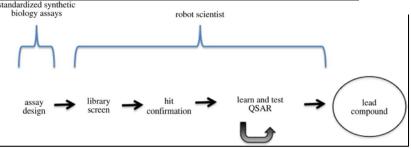
- •6,657,024 optical density measurements@595nm to form 26,495 growth curves
- Formulated and tested 20 hypotheses concerning genes encoding 13 orphan enzymes

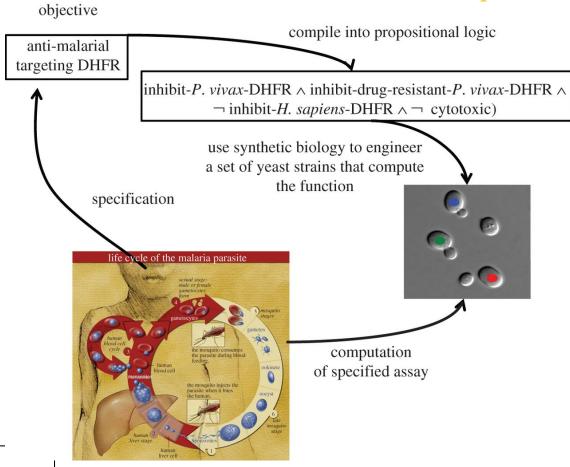
Strong AB: King's Robot Scientist "Eve"











Chimeric yeast can inform drug mechanism of action in mammalian cells.

What do we need for automated inference of biological models?



- Closed-loop control of biology
- Automated design of experiments
 - Cells
 - Matrix chemistry and architecture
 - Metabolic and signaling molecules and other clues
- Quantitative measurements
 - Fluorescent imaging
 - Electrochemical measurements of bioenergetics
 - Untargeted ion mobility-mass spectrometry
 - New types of data from new sensing modalities
- Automated data mining and interpretation
- Organotypic 3D tissue culture, including dynamic control of the extracellular matrix

Narrow versus Strong Automated Biology

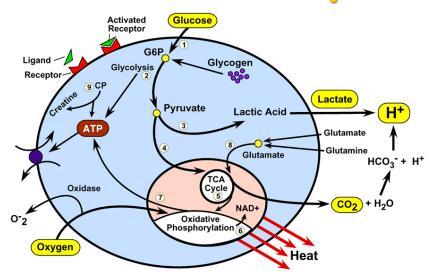


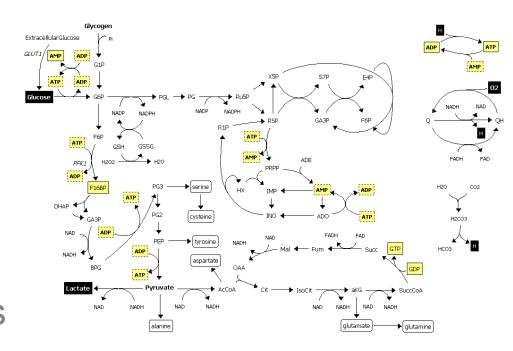
- Narrow AB: the use of software to study or accomplish specific laboratory biology tasks
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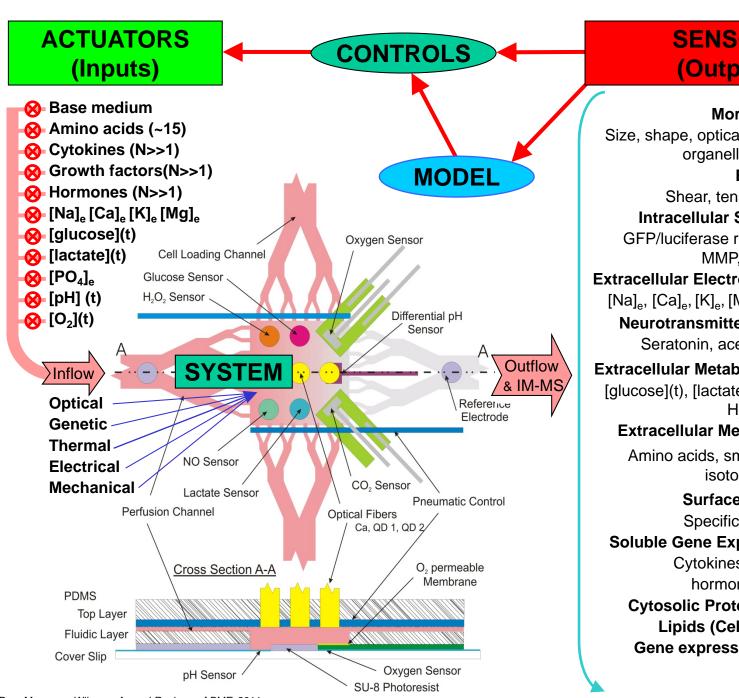
Grand Challenge: 2006

VIJBRE

- Can we design and build a hybrid silicon/biological system that proposes and generates models and conducts experiments on itself to identify the underlying equations that govern cellular dynamics?
- Extracellular: \$3 4
 million and 3 5 years
- Intracellular: \$15 20
 million and 5 10 years







SENSORS (Outputs)

Morphology

Size, shape, optical density, motility, division, organelle configuration

Force

Shear, tension, deformation

Intracellular Signaling (Optical)

GFP/luciferase reporters, [Ca], pH, Vm, MMP, GFP FRET

Extracellular Electrolytes (Electrochemical)

[Na]_e, [Ca]_e, [K]_e, [Mg]_e, [PO₄]_e, [Cl]_e, [HCO₃]_e

Neurotransmitters (Electrochemical)

Seratonin, acetocholine, GABA, ...

Extracellular Metabolites (Electrochemical)

[glucose](t), [lactate](t), [pH](t), $[O_2](t)$, NO(t), $H_2O_2(t)$...

Extracellular Metabolites (GC IM-MS)

Amino acids, small metabolites, stable isotopic markers

Surface Expression

Specific affinity probes

Soluble Gene Expression (nESI IM-MS)

Cytokines, growth factors hormones, enzymes

Cytosolic Proteins (MALDI IM-MS) **Lipids (Cell Lysate IM-MS) Gene expression (mRNA Arrays)**

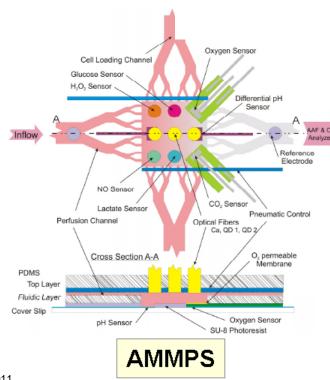


Symbolic-Regression and Estimation-Exploration Algorithms 7 can design experiments to select best symbolic model



Estimation / **Exploration**









Inferring Metabolic Models using the SRA and EEA



Target Model placed in black box with 10% noise

$$\frac{dS_{1}}{dt} = 2.5 - \frac{100 * A_{3}S_{1}}{1 + 13.68 * A_{3}^{4}}$$

$$\frac{dS_{2}}{dt} = \frac{200 * A_{3}S_{1}}{1 + 13.68 * A_{3}^{4}} - 6 * S_{2} - 6 * S_{2}N_{2}$$

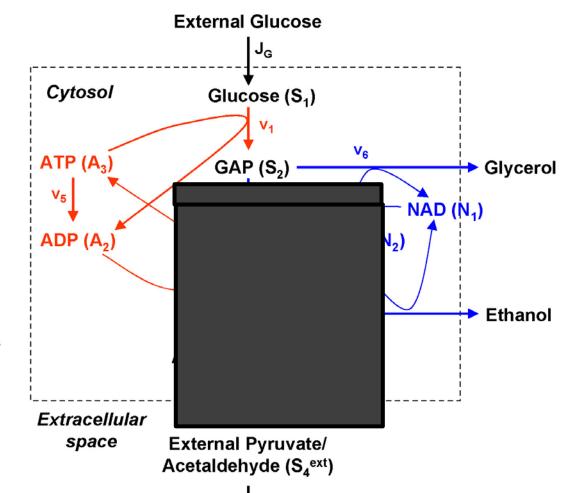
$$\frac{dS_{3}}{dt} = 6 * S_{2} - 6 * N_{2}S_{2} - 64 * S_{3} + 16 * A_{3}S_{3}$$

$$\frac{dS_{4}}{dt} = 64 * S_{3} - 16 * A_{3}S_{3} - 13 * S_{4} - 100 * N_{2}S_{4} + 13 * S_{5}$$

$$\frac{dN_{2}}{dt} = 6 * S_{2} - 18 * N_{2}S_{2} - 100 * N_{2}S_{4}$$

$$\frac{dA_{3}}{dt} = -1.28 * A_{3} - \frac{200 * A_{3}S_{1}}{1 + 13.68 * A_{3}^{4}} + 128 * S_{3} + 32 * A_{3}S_{3}$$

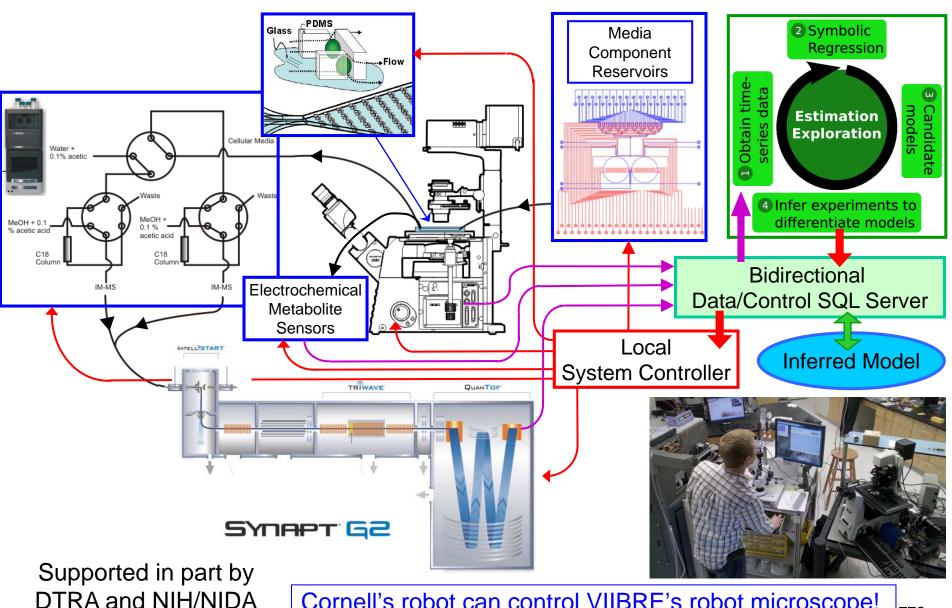
 $\frac{dS_5}{dt} = 1.3 * S_4 - 3.1 * S_5$



Schmidt, Vallabhajosyula, Jenkins, Hood, Soni, Wikswo, and Lipson. Automated probing and inference of analytical models for metabolic networks. Phys.Biol. 8 (5):055011, 2011.

2014: VIIBRE's Robot Scientist for Automated Omni-Omics





Cornell's robot can control VIIBRE's robot microscope!



So how do we implement the biology for our Robot Scientist?

How have we been studying biology?

People

We are severely limited in isogenetic controls, interventions, and data when studying normal subjects and patients.

Animals

Animals, including non-human primates, are not people and have significant genetic and physiological differences.

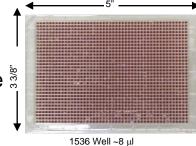
Cells in vitro

2D biology on plastic: Many biological experiments are conducted on cells that

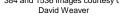
- have cancer.
- are inbred.
- are diabetic.
- are potatoes on a stiff plastic couch without exercise
- enjoy neither gender nor sex,
- live almost entirely in the dark,
- gorge themselves on sugar once a day,
- may be slowly suffocating in an increasingly acidic environment.
- live in their own excrement,
- never bury their dead,
- may take a complete or only partial bath every day or two,
- and talk only to cells of like mind.







384 Well ~40 ul



384 and 1536 images courtesy of

One might get reproducible, statistically significant results, but are they relevant to human biology and disease?

Seven Themes 1. A brief history 2. Just ho understanding the complexity Can sensors, actuators, controllers nd robot scientists help address. delogical complexity 5. How might Organoids and Organs on Chips change the way we study biology? 6. What does Multi-Omics offer? 7. Closing the circle



Circa 2011, organoids and organs-on-chips break into the limelight

- Schmeichel and Bissell "Modeling tissuespecific signaling and organ function in three dimensions." J Cell Science (2003)
- https://www.ted.com/speakers/mina_bissell

A hot, new in vitro model for biology

Complex 3D biology is a better model than 2D biology.

3D Organoids

Are self-organizing models with tissue-level functions and disease phenotypes.

Demonstrate development

Can be transplanted

Can be a medium-to-high throughput assay

Hard to replicate an individual organoid

May benefit from engineered hydrogels

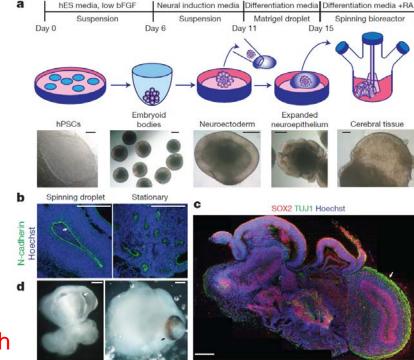
Hard to perfuse or apply uniform shear stress

Hard to quantify barrier functions

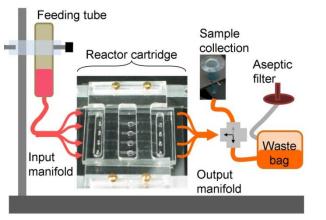
Hard to visualize when living

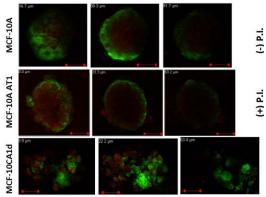
Hard to integrate with other organ systems with proper volumes

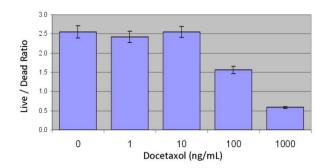
Contributions from Kapil Bharti (NIH/NEI)



Lancaster, ..., Knoblich. Cerebral organoids model human brain development and microcephaly. Nature, 2013.







Markov, ..., McCawley. Thick-tissue bioreactor as a platform for long-term organotypic culture and drug delivery. Lab Chip 12:4560-4568. 2012.

The "Media Volume" problem

Relative sphere sizes:

nL media vs pL cell

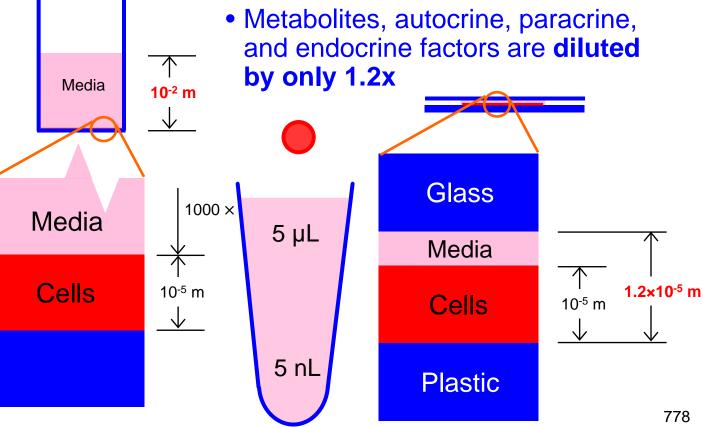


Conventional culture

- A typical picoliter cell requires a nanoliter of media per day.
- A 10 μm layer of cells covered by a 10,000 μm layer of media.
- A 5 nL spheroid in 5 μL of media
- 1 or 2 days between fluid changes
- Metabolites, endocrine, autocrine, and paracrine factors are diluted 1000fold.



- A typical picoliter cell requires a nanoliter of media per day.
- A 10 μm layer of cells is covered by a 2 μm layer of media.
- 5000 fluid changes/day



Another hot new in vitro model for biology Complex 3D biology is a better model than 2D biology.



Organ Chips

Better than 2D biology Ideal for barrier functions Can reproduce physiological flows Provide a thick ECM for scaffolding and drug/factor binding Support organ-organ interactions Sufficient tissue for multi-omics of

Can use minimal media volumes Will be vascularized soon May ultimately reduce drug costs Possible to build a single-patient homunculus

10's to 1000's of variables

Could build animals-on-chips Can require microfluidics and control Not yet high throughput Are expensive today (hardware,

effort, human cells, real estate) Not fully validated vs in vivo, e.g., no

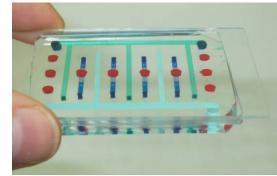
WGCNA yet Can't be transplanted Mammary gland on a chip

Heart on a chip

Prabhakar Pandian and

Kapil Pant,

SynVivo/CFDRC

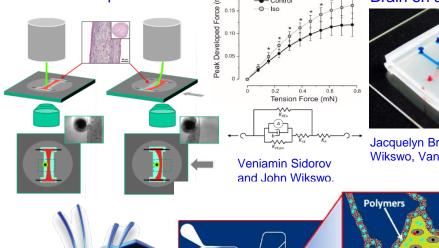


Lisa McCawley and Dmitry Markov, Vanderbilt

T cells in a lymph node on a chip 高高高高高高高高高 多有线 多黄色的 化黄色 Shannon Faley, Kevin Seale and John Wikswo,

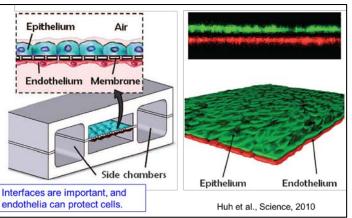
Brain on a chip

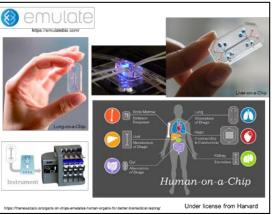
Jacquelyn Brown and John Wikswo, Vanderbilt

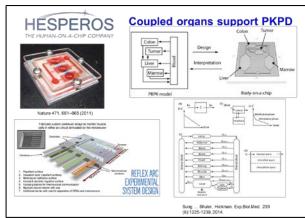


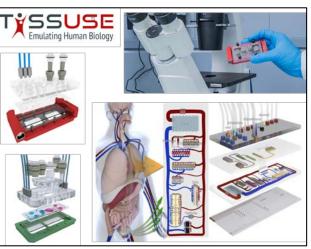
http://www.synvivobio.com

Organ-Chips, Organ Chips, Organ Chips...

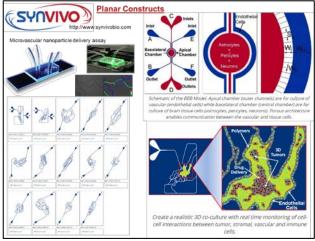


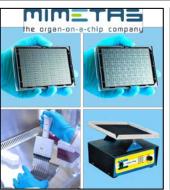










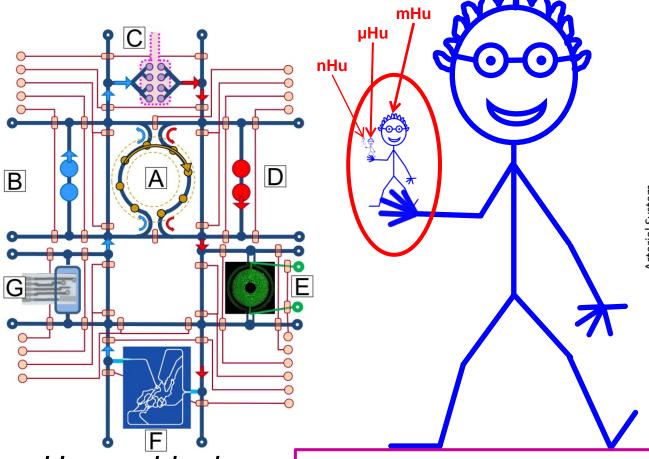




Others, and there's room to grow!

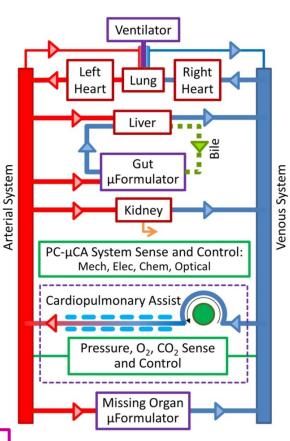
John Wikswo's goal – Determine how best to fit two new *Homunculi* species into the biomedical research ecosystem.





Homo chippiens NanoHuman (nHu)

What can organs-on-chips do for basic research and tox-safety? Single organs and/or coupled-organ homunculi?



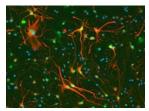
Homo minutus MicroHuman (µHu)

Tissue Chips at Vanderbilt





 Human iPSC-derived neuronal cells



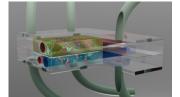
hiPSC glutamatergic neurons



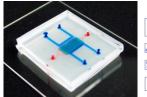


TSC-patient hiPSCs are being used to create brain microvascular endothelial cells, astrocytes, pericytes, and both excitatory and inhibitory neurons 2016

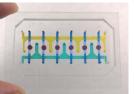
Bioreactors



VIIBRE NVU concept 2012



VIIBRE NVU as built 2014



Mammary gland-on-a-chip 2016

 Control hardware



VIIBRE NVU Perfusion Controller 2014



VIIBRE 24-port valve 2015



MicroFormulator 1.0, 2015



MicroFormulator 2.0, 2016



SmartMotor 2.0, 2016

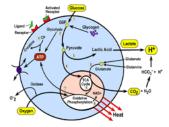
 Analytical chemistry & metabolomics



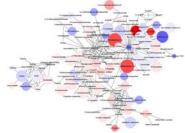
In-line MS of organ chip 2013



VIIBRE MicroClincial Analyzer



Core Carbon Metabolism



MS metabolomics 2016

Translation



CFD3G...



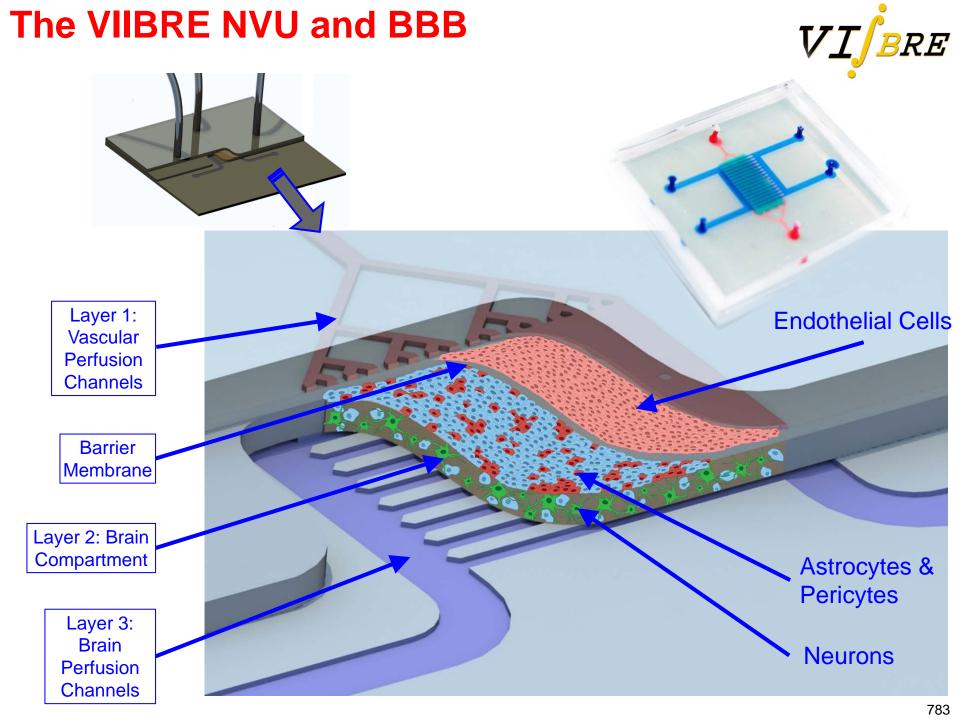
National Center
for Advancing
Translational Science



Issued U.S. patents: 7,435,578;

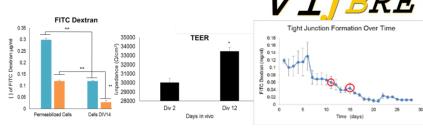
7,534,601; 7,704,745; 7,713,733; 7,790,443;

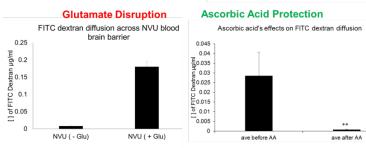
7,974,003; 7,977,089; 7,981,649; 8,129,179; 8,339,704

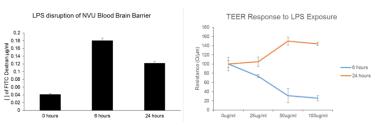


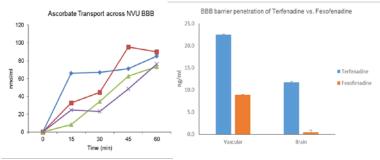
NVU/BBB measurements

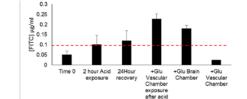
- Tightening of the BBB with time after assembly
- Disruption by glutamate in the brain compartment
- Tightening by ascorbic acid in the vasculature
- Differential responses over time to inflammatory agents (LPS and cytokine cocktails)
- Differential transport across the BBB: ascorbic acid (Y), Terfenadine (Y), Fexofenadine (N)
- Response to combined insults (brain glutamate + acidification)











784

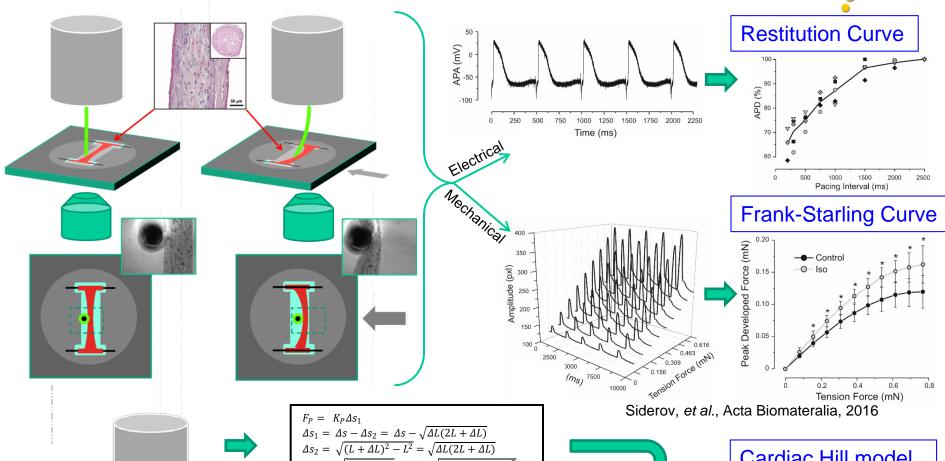
Cardiac I-Wire

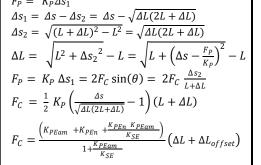
F, $\Delta F(t)$

Stim(t)

 $F_c(t)$, $\Delta L(t)$





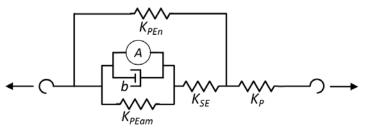


 $\frac{\left(1 + \frac{K_{PEam}}{K_{SE}}\right) f_1(\Delta L) - K_M(\Delta L + \Delta L_{offset}) - F_A}{b\left(1 - \frac{f_2(\Delta L) - K_{PEn}}{K_{CF}}\right)}$

Nashville single-string guitar equation

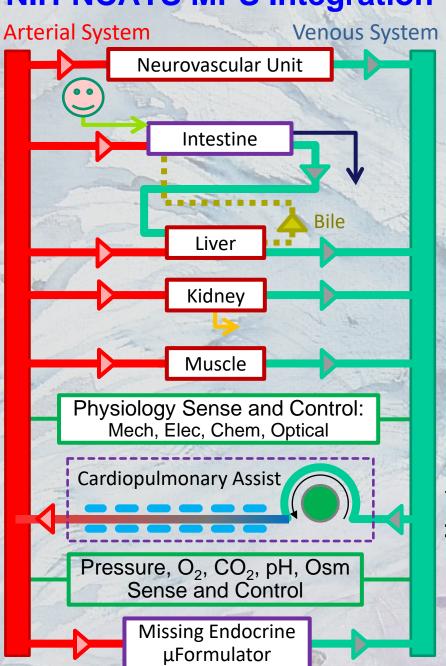


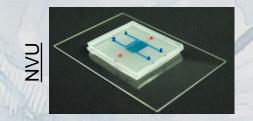
Cardiac Hill model



Schroer, et al., Acta Biomateralia, 2016

NIH-NCATS MPS Integration





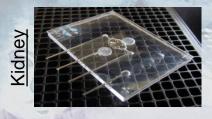




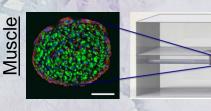












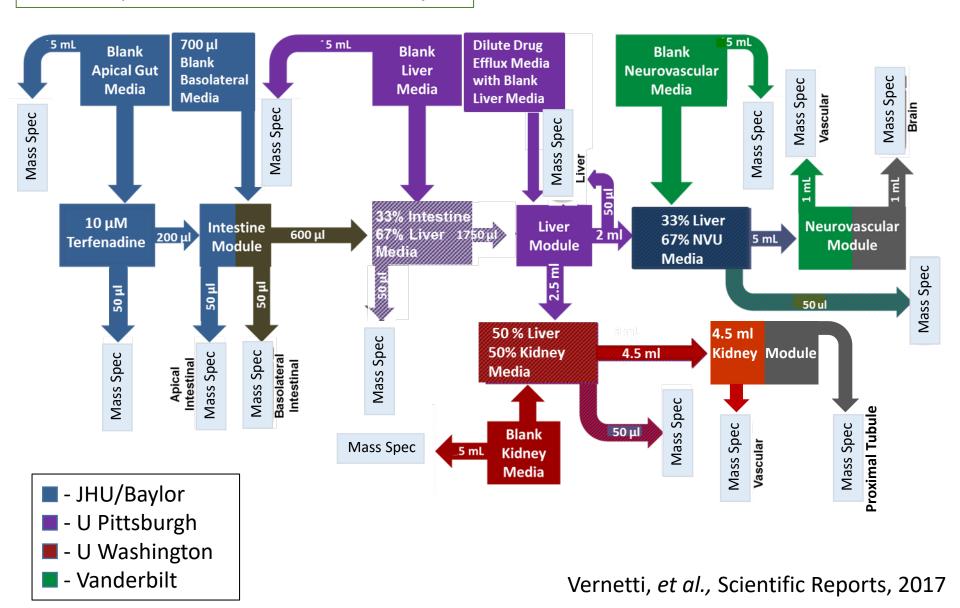




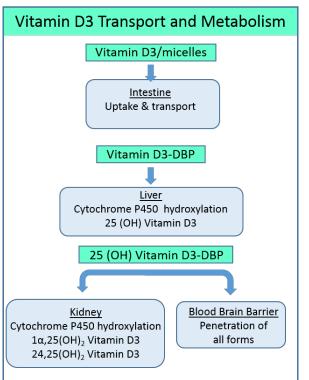


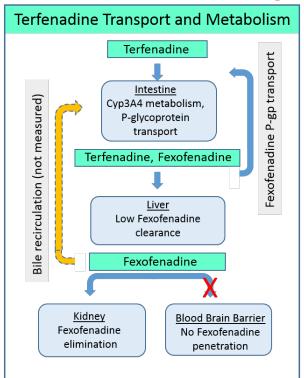
Work Flow for Functional Coupling Experiment

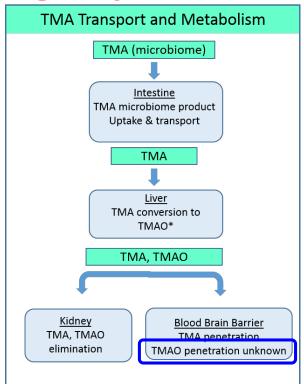
Goal: Couple Gut, Liver, Brain, and Kidney



Work Flow for Functional Coupling Experiment







TMAO penetration into human CSF confirmed the NVU observation: Del Rio, et al., Nutrients, 2017

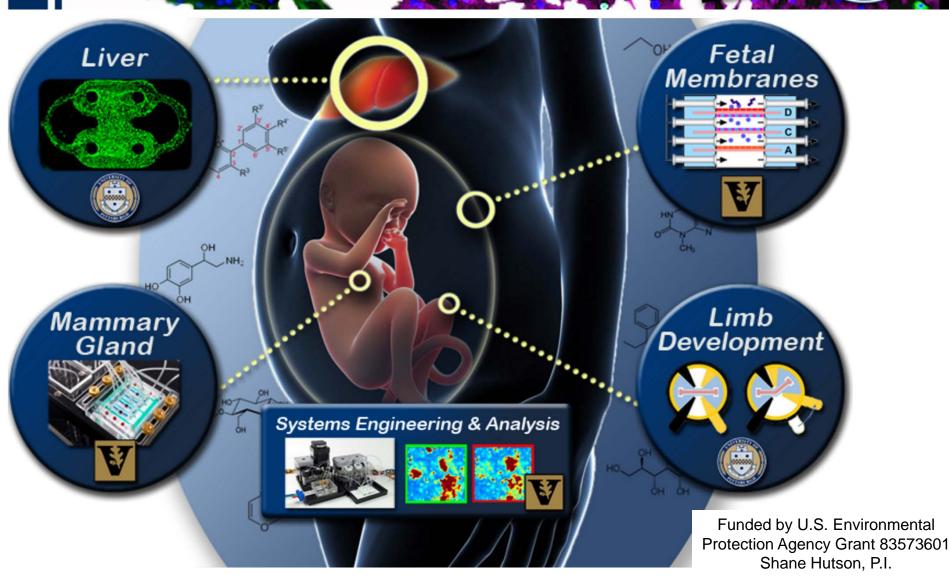
We found 26% TMAO penetration into the NVU brain chamber!

Test Agent/Metabolites	Clinical MPS Model	Intestine	Liver	Kidney	ввв
TMA TMAO	Clinical	Uptake & Transport	TMA → TMAO < 5% TMA Clearance	> 95% TMAO Excreted	TMAO Penetration: Unknown
	MPS	Uptake & Transport	TMA → TMAO < 1% TMA Clearance	~46% TMAO Excreted	26% TMAO Penetration
Terfenadine (Ter) Fexofenadine (Fex)	Clinical	Ter → <i>Fex</i> ; <i>Fex</i> CounterTrans	< 1% Bio T < 95% Fex Clearance	11% Fex Excreted	~0% Fex Penetration
	MPS	Ter → Fex; Fex CounterTrans	< 1.4% Bio T (est.) < 80% Fex Clearance	~ 1% Fex Excreted	~ 0% Fex Penetration
Vitamin D3 (VD3) 25(OH)VD3; 1α,25(OH) ₂ VD3; 24,25(OH) ₂ VD3	Clinical	Uptake & Transport No metabolism	VD3 → 25(OH)VD3	25(OH) $VD3 \rightarrow 1\alpha,25(OH)_2VD3 \& 24,25(OH)_2VD3$	VD3 & 25(OH)VD3 Penetration
	MPS	Uptake & Transport No metabolism	VD3 → 25(OH)VD3 & 24,25(OH) ₂ VD3	$1\alpha,25(OH)_2VD3$ & $24,25(OH)_2VD3$ below LOQ	0.4% VD3 & 6% 25(OH) VD3 Penetration

Key Concordances Between MPS and Clinical Fate for Three Test Agents. Key: Uptake - by jejunum endothelial cells; Transport - from apical to basolateral media; → = Metabolism; CounterTrans = Transport from basolateral to apical media; est. = estimated. Excreted - into proximal tubule lumen; LOQ = limit of quantitation; Penetration - through blood-brain barrier.

Vernetti, et al., Sci. Reports, 2017

Vanderbilt-Pittsburgh Resource for Organotypic Models for Predictive Toxicology



Problems that are at the MPS cutting edge



How accurately can

we recreate micro-

vasculature and the

basement membrane?

The full metastatic cascade

- Localized formation of the primary tumor
- Intravasation into vascular and lymph systems
- Dissemination through <u>vascular and lymph systems</u>
- Extravasation into a competent organ
- Colonization and proliferation with seed-soil interactions

Testing immuno-oncology drugs

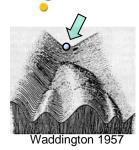
How accurately must we recreate adaptive immunity?

- Requires isogenetic innate and adaptive immune system, tumor, and metastatic niche to avoid host-versus-graft reactions and MHC-HLA incompatibilities.
- May require organ-specific lymph nodes, immune-active spleen and bone marrow for proper programming of multiple types of immune cells.
- CD34+ progenitor cells and B cells have yet to be derived from iPSCs.

790

VIIBRE's Tissue Chips Challenges

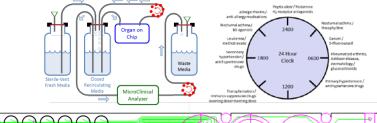
- Human iPSC-derived neuronal cells
- Reduce costs
- Shorten time from patient to iPSC to mature phenotypes
- Develop genotype libraries
- Learn how to control iPSC differentiation

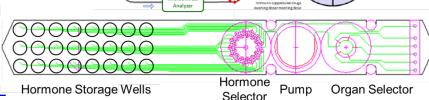


Bioreactors

- Reduce volumes
- Vascularize
- Eliminate PDMS
- Add electrodes to the NVU
 - Reduce volumes
 - Reduce size and cost
- Control hardware Recirculate

 - Add diurnal hormone and nutrient variations





Analytical chemistry & metabolomics

- Reduce volumes
- Detect more analytes on-line at lower cost
- Infer metabolic and signaling networks



- **Translation**
- Make it cost-effective and easy for conventional biologists, toxicologists, and pharmacologists to use organs on chips without a gigantic capital investment or an engineering degree
- Start answering medical questions and solving medical problems

How good a model do we need?



• It depends upon the question you are asking

The best material model for a cat is another, or preferably the same cat.

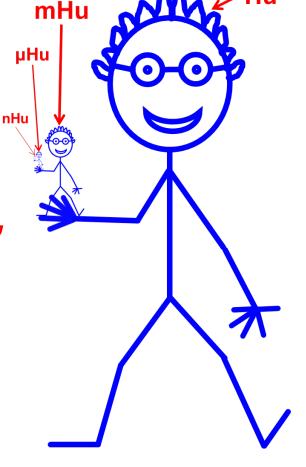
Arturo Rosenblueth and Norbert Wiener. The

Arturo Rosenblueth and Norbert Wiener. The Role of Models in Science. Philosophy of Science 12 (4):316-321, 1945.

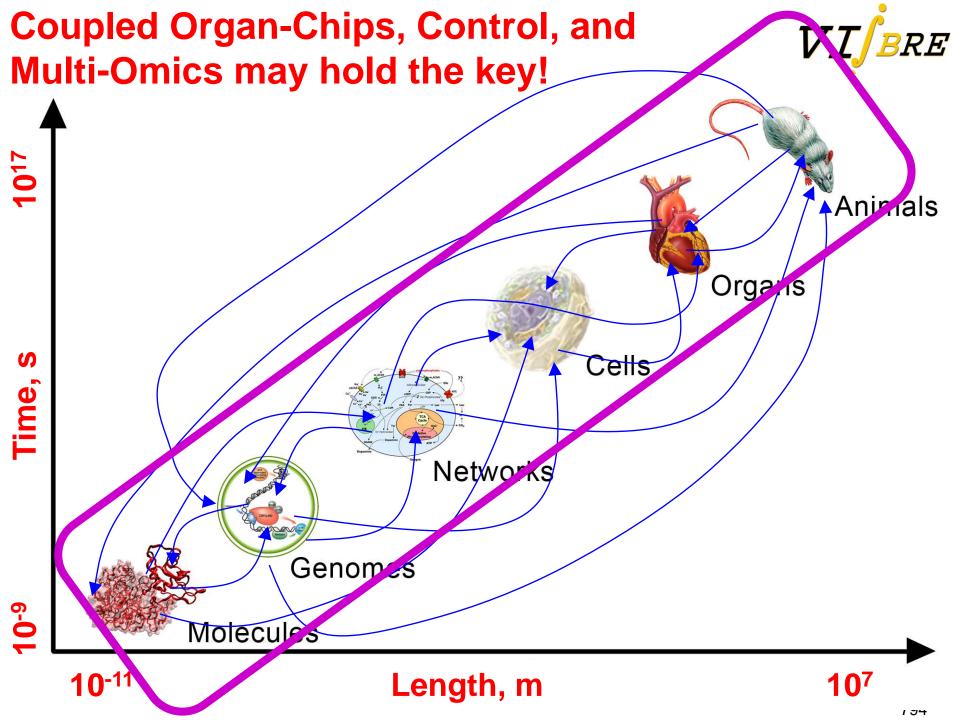
Make your theories simple enough, but not too simple.

~Albert Einstein

Make your organs-on-chips systems simple enough, but not too simple.



Seven Themes 1. A brief histor 2. Just he nderstanding the complexity an sensors, actuators, controllers nd robot scientists help address. jical complexity? 5. How might Organoids nips change the way we study 6. What does Multi-Omics offer? 7. Closing the circle



Monitoring Organ Health/Toxin Response



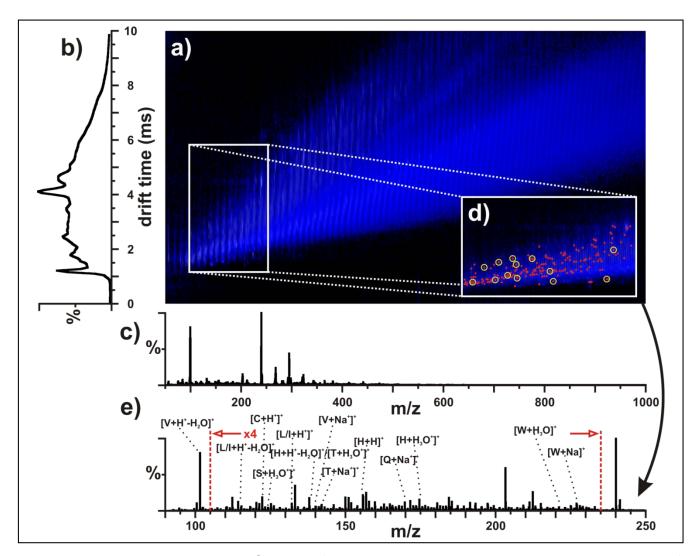
- Cellular morphology
 - Requires fluorescence microscopy
 - Organ-on-chip module should be HCS confocal compatible
- Genetically encoded fluorescent reporters
- TEER
- Bioenergetics
- Protein production
 - Albumin
 - Bile acids
 - Cytokines
 - Cyp activity
 - LDH release
- Transcriptomics
- Drug metabolism
- Untargeted metabolomics
- Metabolic activity (glucose, lactate, pH, oxygen)



The sensitivity of many assays is set by the ratio of cell volume to media volume!

Tyrode's solution perfused through a normal beating mouse heart Dynamic metabolomics

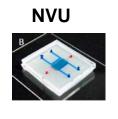


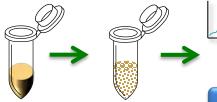


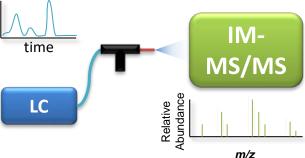
Sample from Franz Baudenbacher, slide courtesy of John McLean

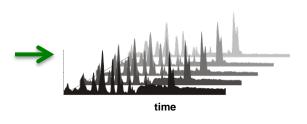
NVU/BBB UPLC-IM-MS workflow











Sample preparation

metabolites extracted using ice cooled methanol:H₂O (80:20), incubated -80°C overnight, spun down at 15,000 rpm, 15 min dried down *in vacuo*

Sample Acquisition

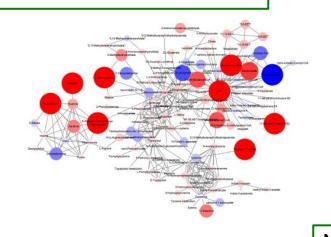
LC IM-MS/MS of metabolite extracts

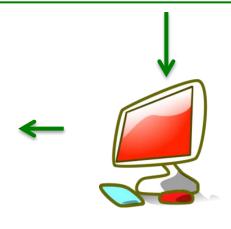
LPS or Cytokine treated samples

Data Alignment and Biostatistical Analysis

Progenesis QI

Pathways Analysis	p-value
Vitamin E metabolism	8.00E-05
Glutathione Metabolism	1.13E-03
Prostaglandin formation from arachidonate	6.48E-03
Aspartate and asparagine metabolism	9.95E-03
Drug metabolism - cytochrome P450	9.97E-03





Network & Pathway Module Output

Network and Pathway analysis *Mummichog*

Metabolomic pathway analysis with high mass-accuracy UPLC-IM-MS is accelerating the incorporation of untargeted metabolomics into mechanism of action studies.

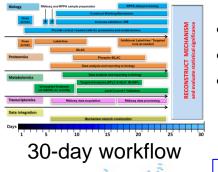
Brown et al., J. Neuroinflammation, 2016

Rapid Threat Assessment (RTA) of MoA

Richard Caprioli, PI, DARPA W911 NF-14-2-0022. **Objective**: Use multiomics to characterize drug and toxin

Mechanism of Action (MoA) in 30 days or less.

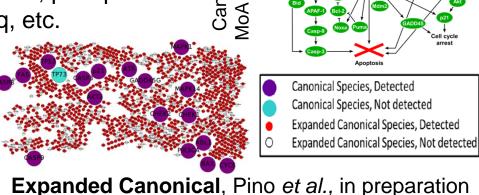
Challenge 1: A549 cells treated with 50 µM cisplatin for 1, 6, 24 and 48 h. MS proteomics (mudpit, SILAC, phosphoproteomics), IM-MS metabolomics, RNAseq, etc.



ECN - 2560 nodes

DDN - 6386 nodes

- 254,296 total features
- 55,898 unique species
- 13,483 (24%) species significantly changed



DDN: 40% of network measured (2,215 species) 86% of detected species changed significantly

Expanded Resistance vs
Data Driven Networks

ERN - 2557 nodes

Cisplatin

DNA damage

DNA dam

Time-resolved omni-omics has great potential!

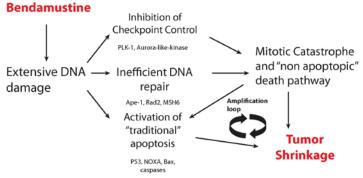
Expanded Canonical vs

Data Driven Network

New Canonical Cisplatin MoA Norris *et al.*, J. Proteome Res. 2017

RTA – Bendamustine MoA

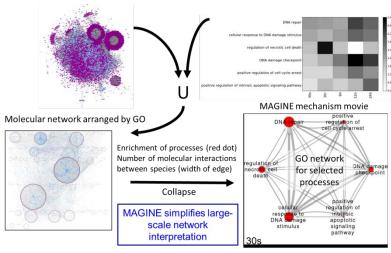
Known Mechanism of Action of Bendamustine



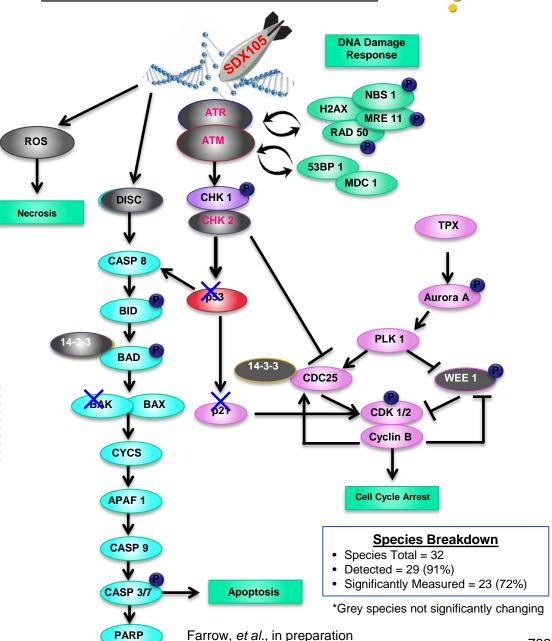
Leoni and Hartley, Seminars in Hematology (2011)

Vanderbilt RTA 2nd 30-day Challenge

- Acquire 781,072 data points spanning
 12 time points and 7 platforms
- How do we extract and integrate knowledge from these data?

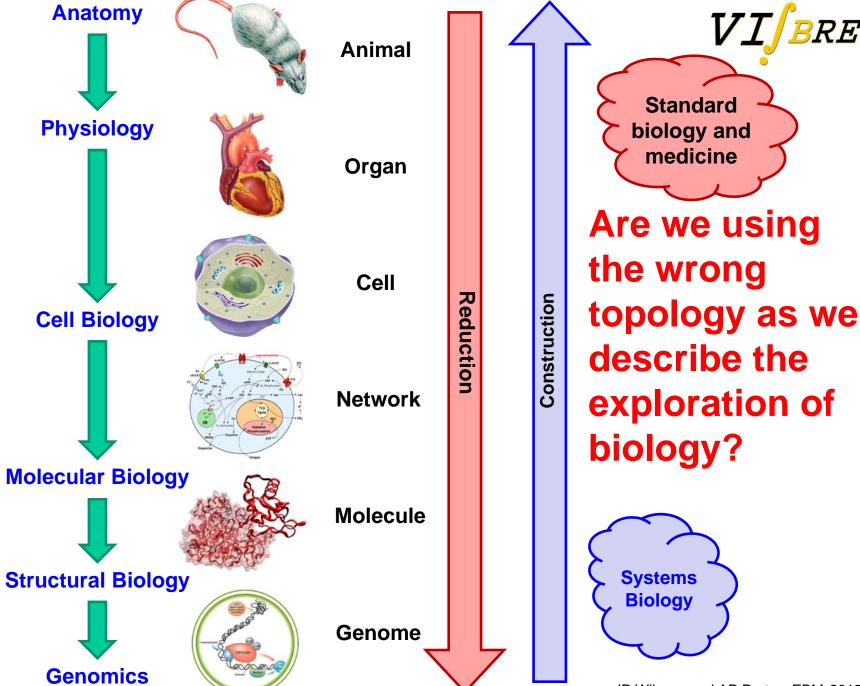


Vanderbilt-RTA Postulated Mechanism of Action



Pino, et al., in preparation

Seven Themes 1. A brief history 2. Just ho understanding the complexity Can sensors, actuators, controllers nd robot scientists help address. biological complexity? 5. How might Organoids nips change the way we study What does Multi-Omics of 7. Closing the circle Julia Wikswo



JP Wikswo and AP Porter, EBM, 2015

Hermeneutics, noun

[hərmə n(y)oodiks/]

The study of the methodological principles of interpretation (as of the Bible).

http://www.merriam-webster.com/dictionary/hermeneutic

The first order art and the second order theory of understanding and interpretation of linguistic and non-linguistic expressions.

http://plato.stanford.edu/entries/hermeneutics/

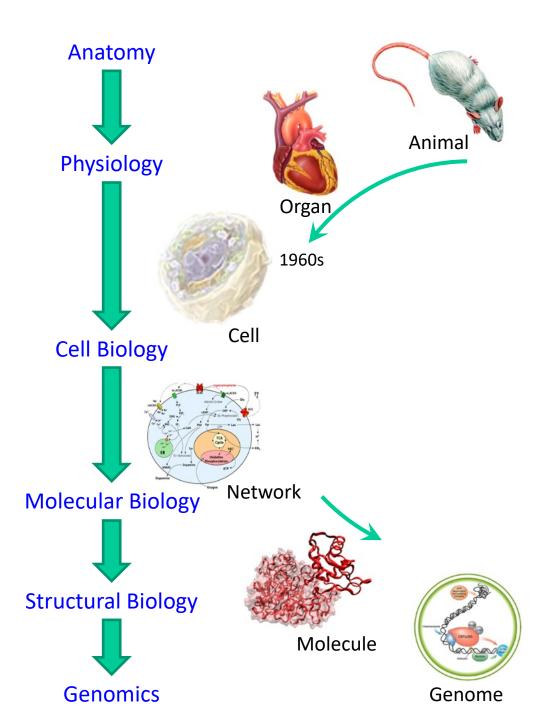
Hermeneutic Circle, noun

[hərmə n(y)oodik sərk(ə)l]

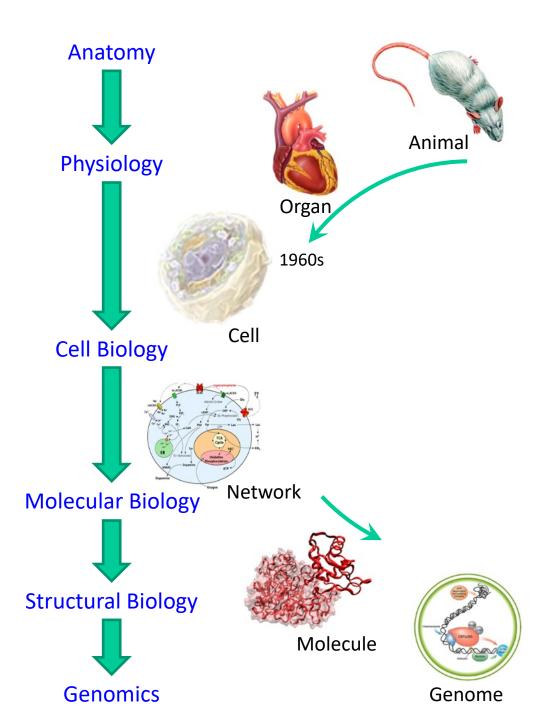
Whole *

One cannot understand the whole until one understands the parts, and one cannot understand the parts until one understands the whole.

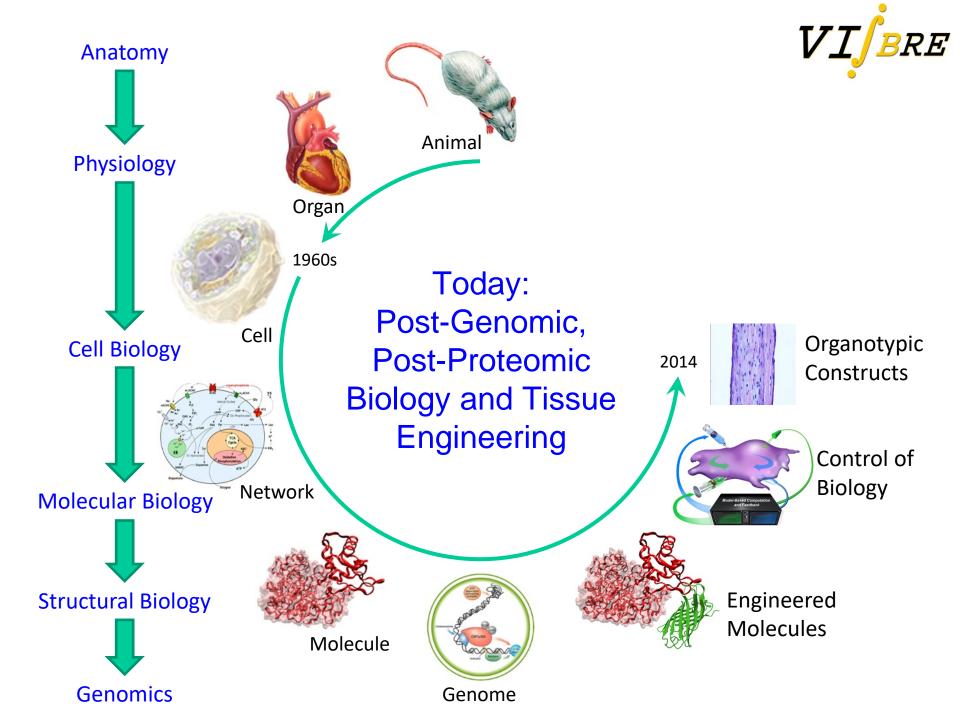
Parts

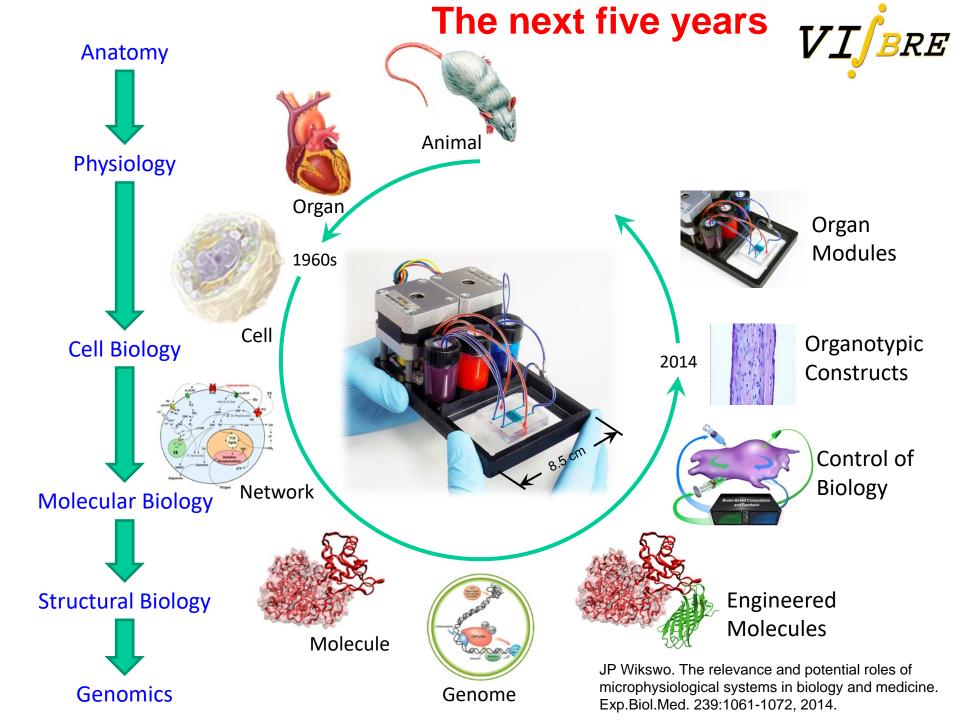






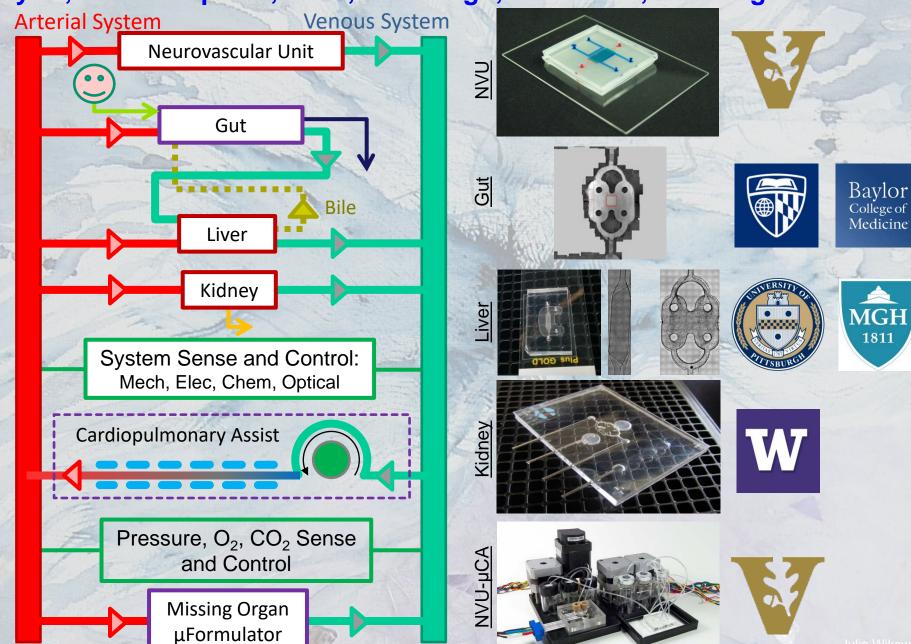






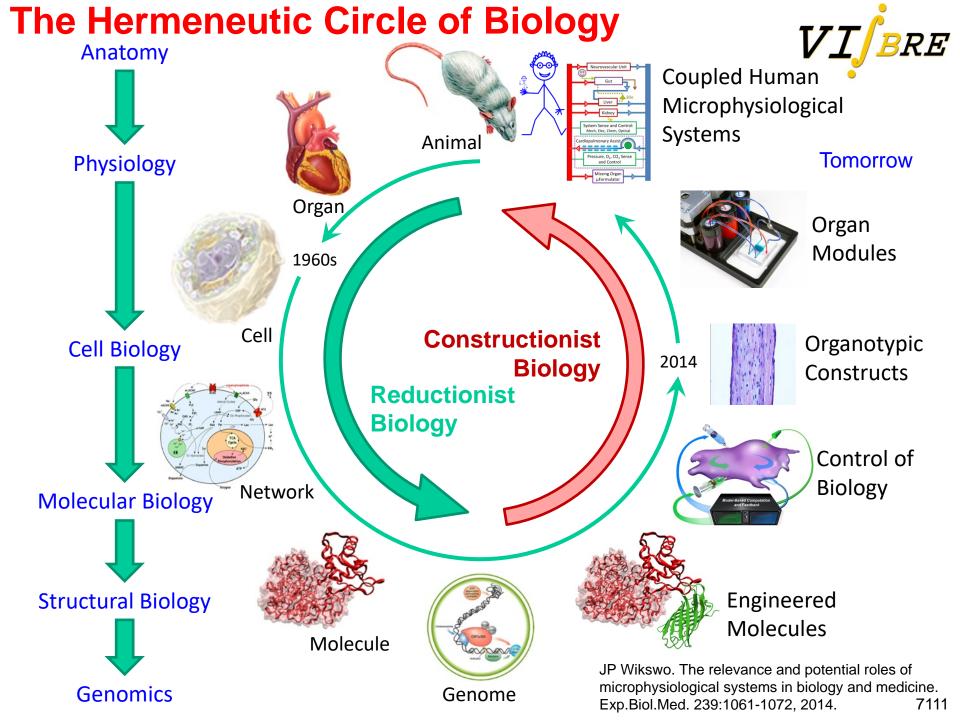
NIH NCATS MPS Integration:

Baylor, Johns Hopkins, MGH, Pittsburgh, Vanderbilt, Washington

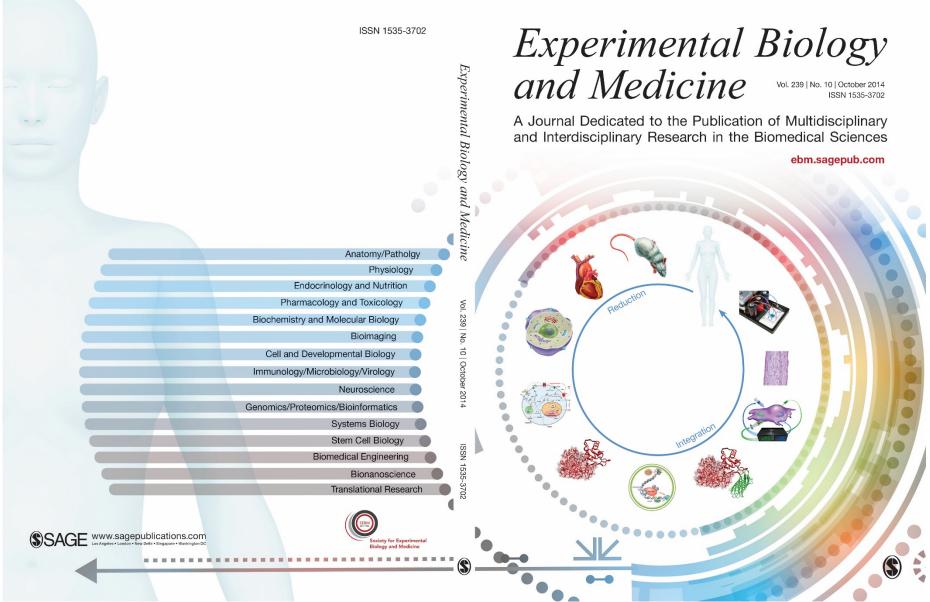


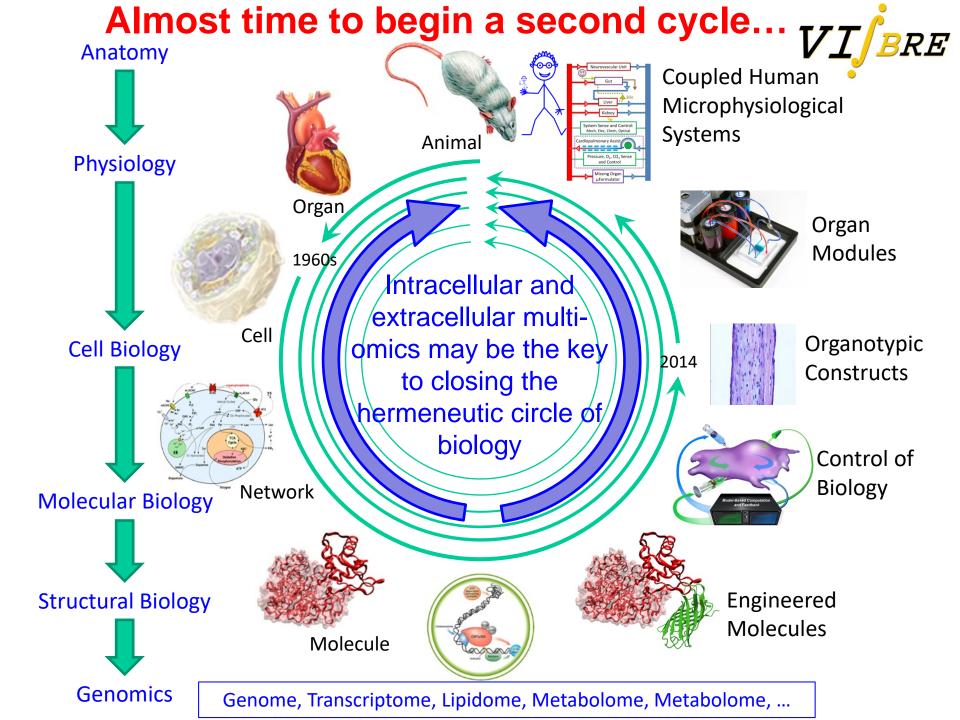
The next five years: Disease Models Anatomy **Coupled Human** Microphysiological Systems **Animal Tomorrow Physiology** Organ Organ Organs on Chips will Modules 1960s contribute to Systems Biology, Cell Organotypic **Cell Biology** Integrative Physiology, 2014 Constructs and **Quantitative Systems** Control of Pharmacology, and **Biology** Network Disease research! **Molecular Biology Structural Biology Engineered** Molecules Molecule JP Wikswo. The relevance and potential roles of microphysiological systems in biology and medicine. **Genomics** Genome Exp.Biol.Med. 239:1061-1072, 2014. 7109

Almost time to begin a second cycle... $VI/_{BRE}$ **Anatomy Coupled Human** Microphysiological Systems **Animal Physiology** Organ Organ Modules 1960s The Hermeneutic Circle of Biology: One cannot understand the Cell Organotypic **Cell Biology** organism until one understands 2014 Constructs the parts, and one cannot understand the parts until one understands the organism. Control of **Biology** Network Molecular Biology **Structural Biology Engineered** Molecules Molecule **Genomics** Genome











What's next for me?

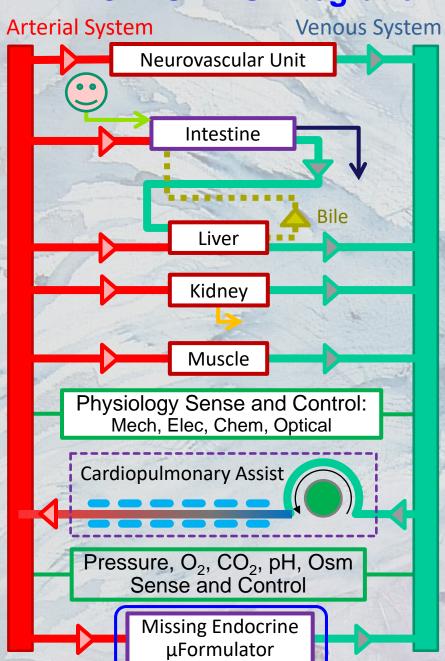
- Missing organs
- Exploring pharmacokinetics (PK) without changing the medicinal chemistry
- Controlling iPSC differentiation

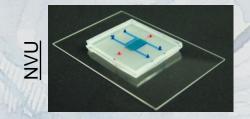
The "Missing Organ" Problem



- The human body has over a hundred organs.
- The Tissue Chips community is building "toy models" of humans, i.e., Homunculi.
 - We cannot include every organ.
 - We should not include every organ.
- For a coupled organ system, there may always be a key organ that has been omitted.
- Missing <u>secretory</u> organs can be replaced with a Missing Organ Microformulator.
 - Hormones, hormones

NIH-NCATS MPS Integration









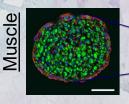
















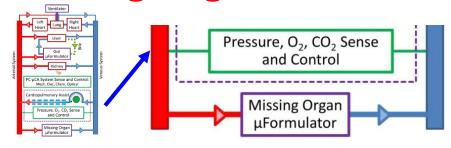




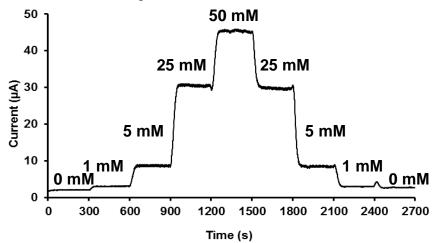


Missing Organ MicroFormulator (µF)

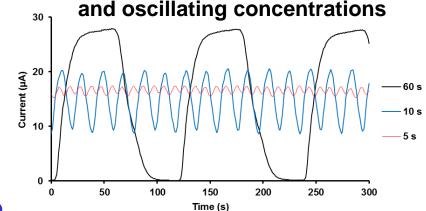




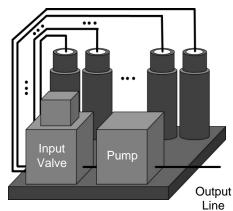
Delivery of desired concentration



Time-division multiplexing and oscillating concentrations



- Cliffel Group:
 - Testing performance with e-chem
 - Reduction of ferricyanide at -0.16V
 vs. Ag quasi-reference.
- Low leakage between ports
- Programming allows rapid switching between ports for dilution, gradients, and calibration of electrochemical sensors





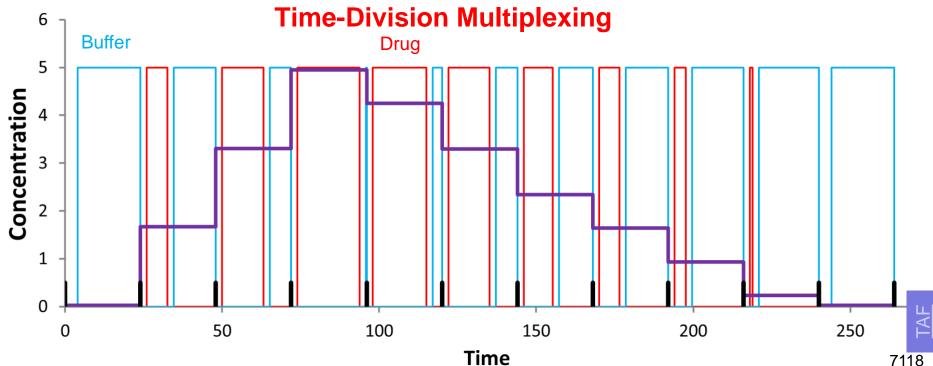
United States Patent, 9,618,129 B2

A normally closed rotary planar valve for microfluidic applications, F. E. Block III, J.R. McKenzie, P. C. Samson, D. A. Markov, and J. P. Wikswo, *In Preparation*.

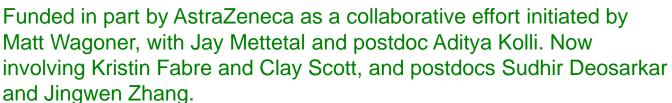
What can you do with a µF-96?



- Matt Wagoner AstraZeneca: "Make me 96 of them! Use time-division multiplexing to create realistic PK drugexposure profiles individualized for each and every well in a 96-well-plate HTS assay!"
 - Conventional cell culture
 - Massively parallel organs on chips
 - Organoid HTS arrays
 - Hanging drop
 - Transwells

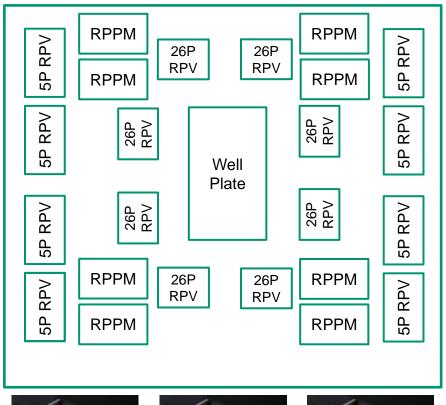


μF-96 v1.0: January 2016









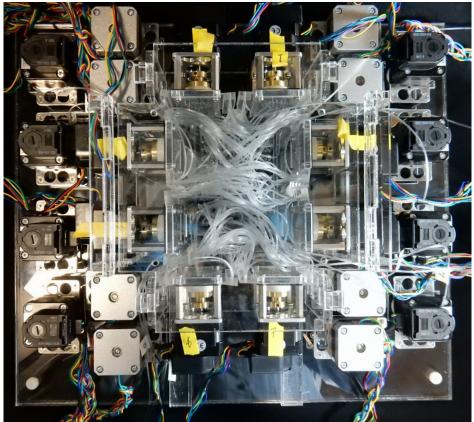












Can individually formulate, deliver, and remove custom media cocktails to each well of a 96-well plate to simulate PK profiles.

96-Channel MicroFormulator (µF-96), v2.0



 For each well, formulate a custom media/drug mixture in real time.

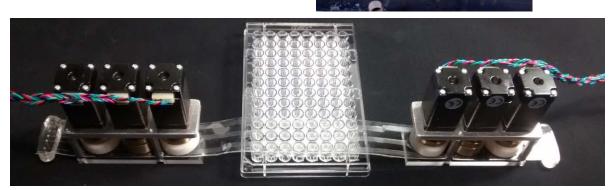
 Change 10% of the fluid in each well 40x/day.

















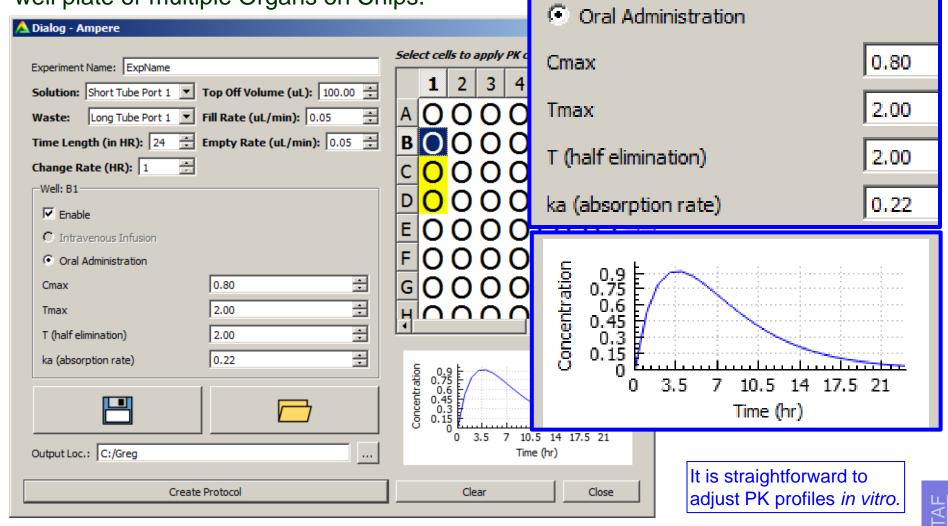


Well Plate Tool

Challenge: Develop a tool for configuring and tracking fluid delivery (including PK exposure profiles) to individual wells in a 96-well plate or multiple Organs on Chips.







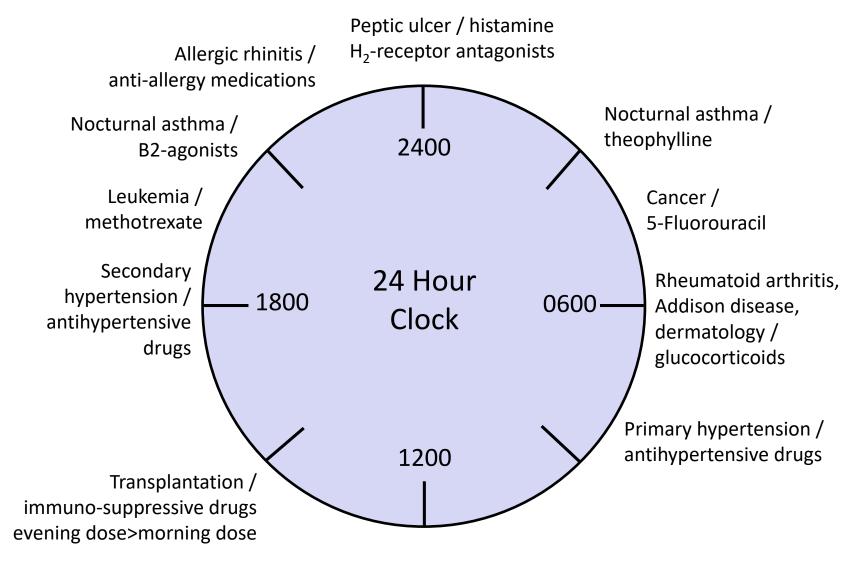
2019: CN Bio Innovations' PharmacoMimix™ VI BRE





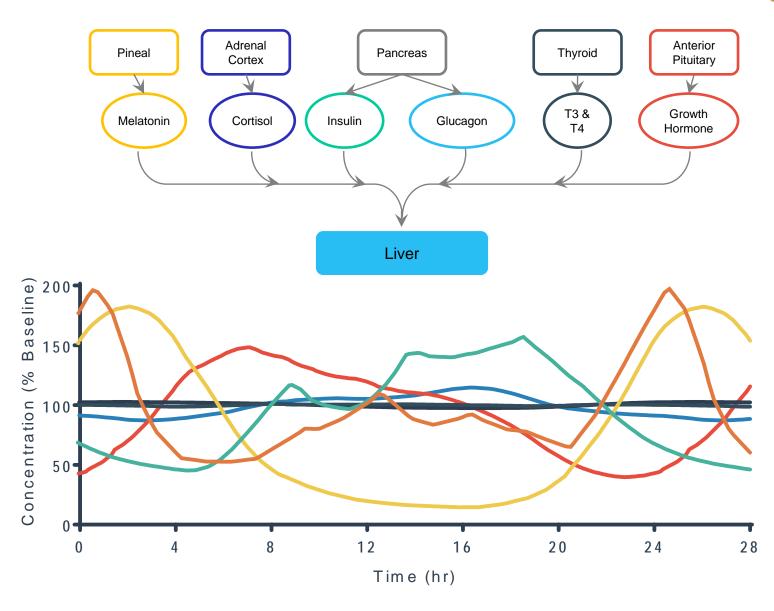
Diseases and Optimal Drug Dosing are Circadian





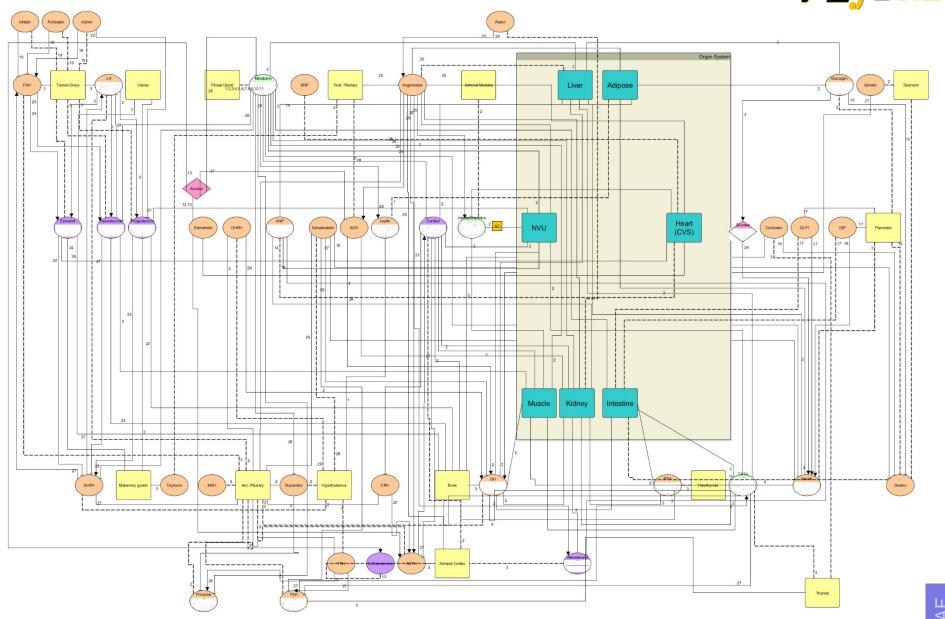
Diurnal Variations of Liver-Regulating Hormones





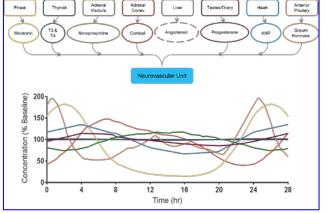
Which endocrine organs / hormones do we need?

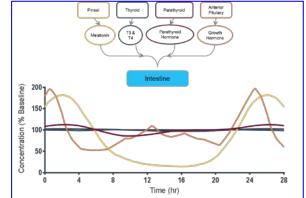




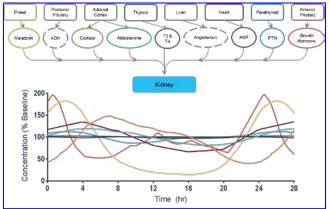


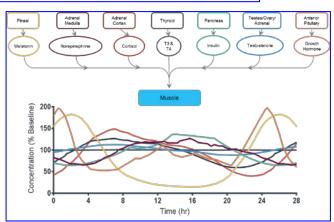


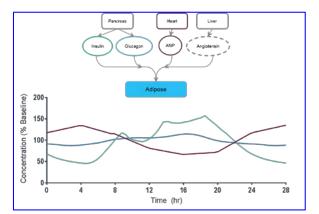


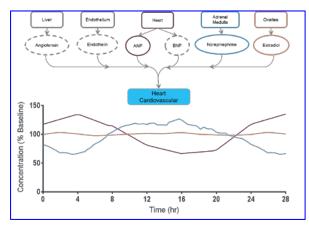


- Neurovascular Unit
- Kidney
- Muscle
- Adipose
- Heart / Cardiovascular



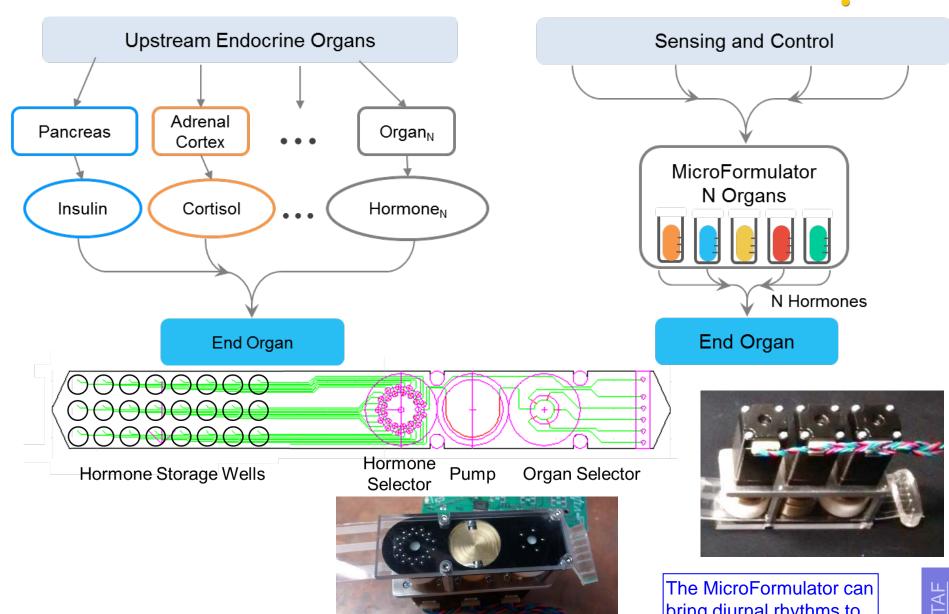






Diurnal Variations of Organ-Regulating Hormones





Cyr, Avaldi, and Wikswo, Experimental Biology and Medicine, 2017

7127

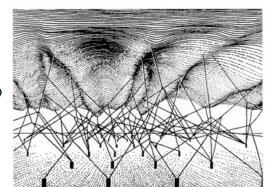
bring diurnal rhythms to

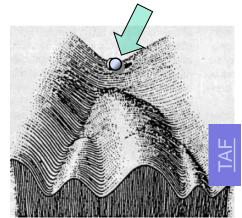
biology on plastic.

What can you do with a µF-96?

- Use time-division multiplexing to create realistic PK
 drug-exposure profiles individualized for each and every
 well in a 96-well-plate HTS assay.
 - Conventional cell culture
 - Massively parallel organs on chips
 - Organoid HTS arrays
- Explore in a massively parallel manner the multitude of combinations of growth factors and other compounds that are needed to guide iPSC differentiation to specific cellular phenotypes.
 - Readily applicable to organoid developmental biology
 - Suitable for machine learning and automated model inference.







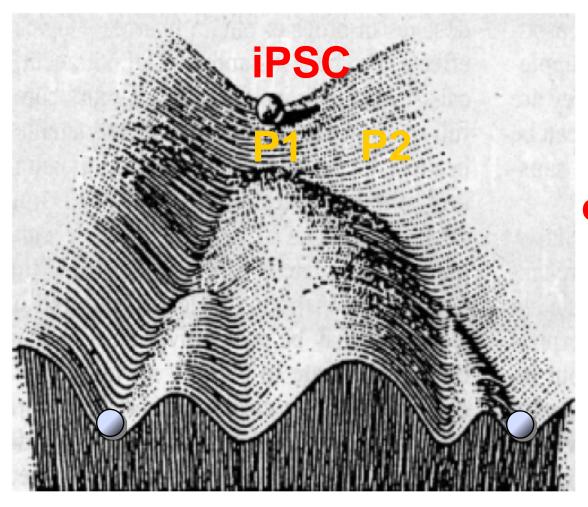
Add and remove growth factors, etc., at will

Stem Cell Differentiation: Cell phenotype rolling down the epigenetic landscape

36



S. Huang and D.E. Ingber / A Non-Genetic Basis for Cancer Progression and Metastasis

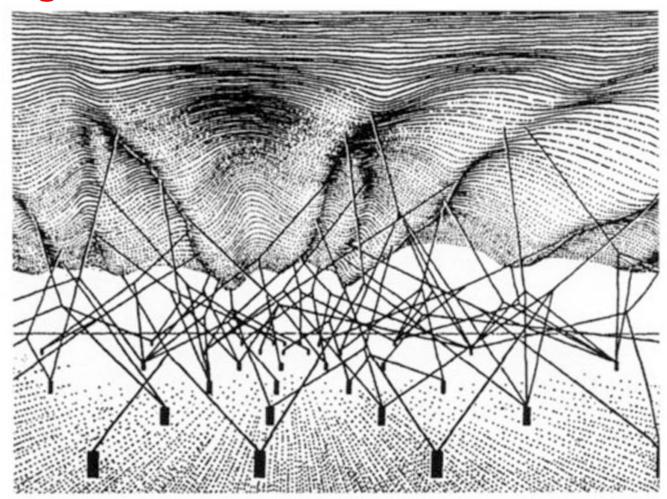


This is a 10² to 10⁵ dimensional surface



Fig. 3. Waddington's epigenetic landscape. Reproduced from C.F. Waddington, 1957 [64]. We postulate here that the metaphoric epigenetic landscape corresponds to the attractor landscape (Fig. 2) that can be reduced to the dynamics of a gene regulatory network.

The epigenetic landscape reflects complex and dynamic genetic control.



- iPSCs
- ESCs
- SCNT-ESCs
- Epigenetic control

Nonequilibrium thermodynamics allows uphill motion.

We need to control the sticks!



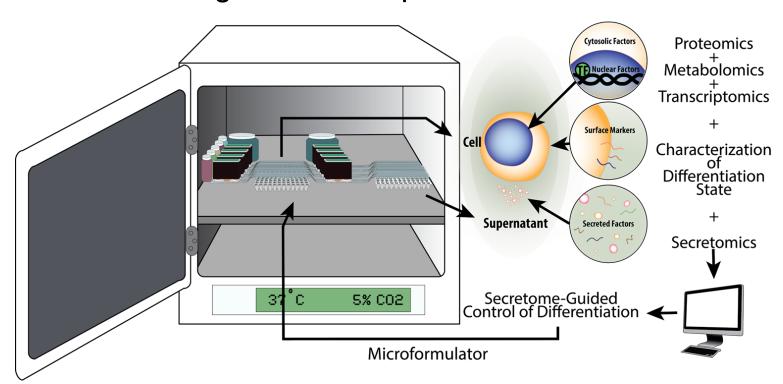


Can the secretome be used to control iPSC differentiation?



ATLAS

- Automated
- Quantitative guidance
- Non-destructive cell monitoring
- Combinatorics through multi-well plates



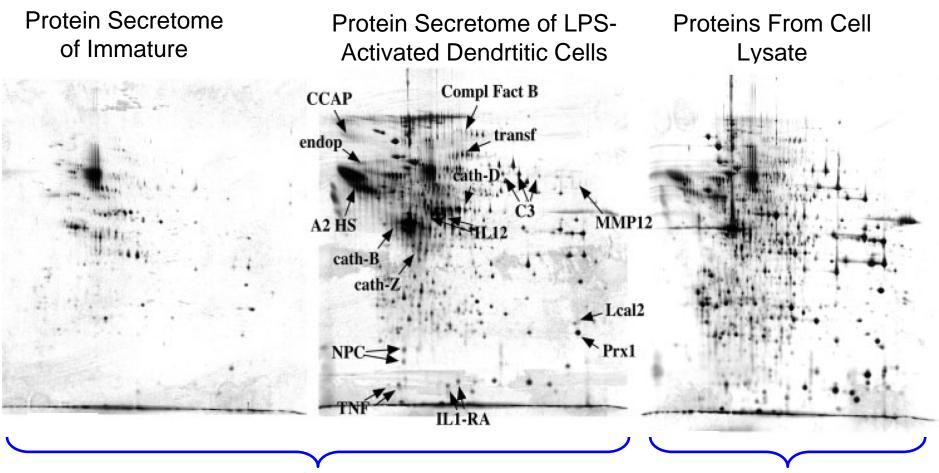




We need a non-destructive signal to control iPSC differentiation!

Proteins in Secretome vs Cytosol





Protein Secretome

Lysate Proteins

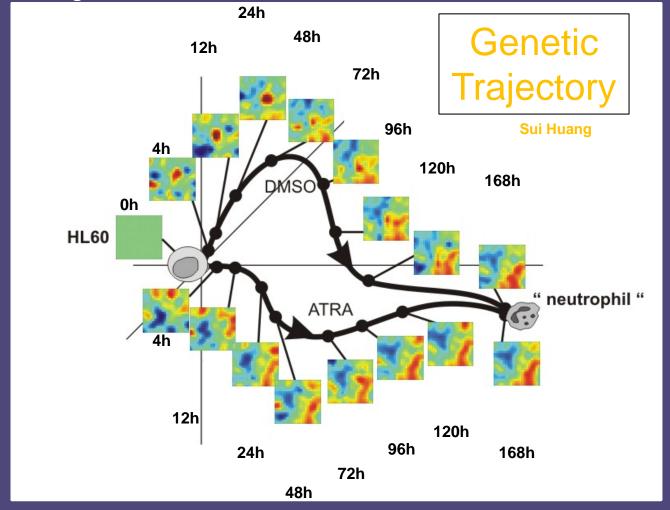
What is the small-molecule, metabolite secretome?

TAF

Chevallet, et al. Proteomics. 2007.

Cell fates as high-dimensional attractor states of complex gene regulatory network

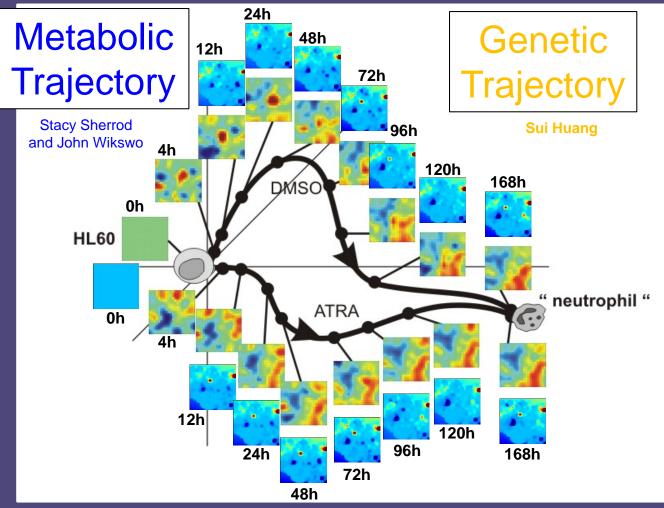
Genome-wide gene regulatory networks govern the behavior of cells (i.e., differentiation, death, etc.). Gene expression profiling can be used to show that two trajectories of neutrophil differentiation converge to a common state from different directions.

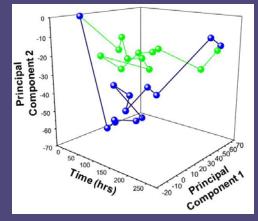




Cell fates as high-dimensional attractor states of complex gene regulatory network

Genome-wide gene regulatory networks govern the behavior of cells (i.e., differentiation, death, etc.). Gene expression profiling can be used to show that two trajectories of neutrophil differentiation converge to a common state from different directions.





Secretome metabolomics can distinguish transitions in intracellular state

Stacy Sherrod and John Wikswo with the support of the Millipore Corporation

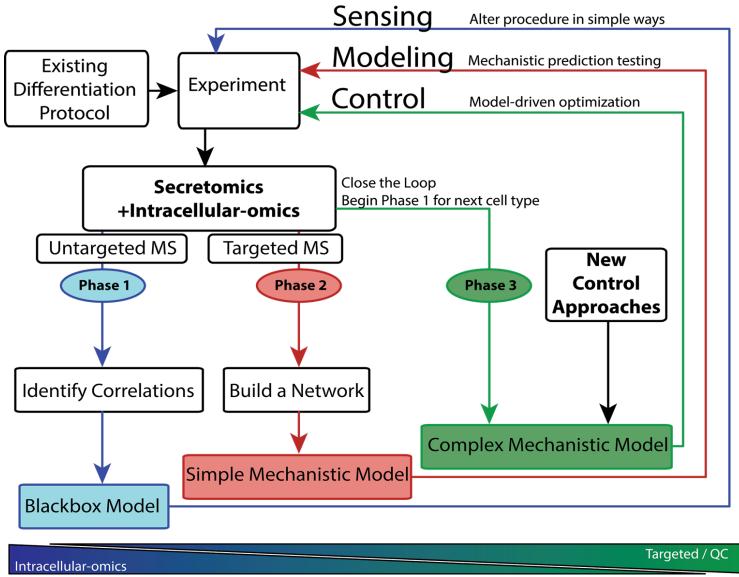






Can the secretome be used to control iPSC differentiation?

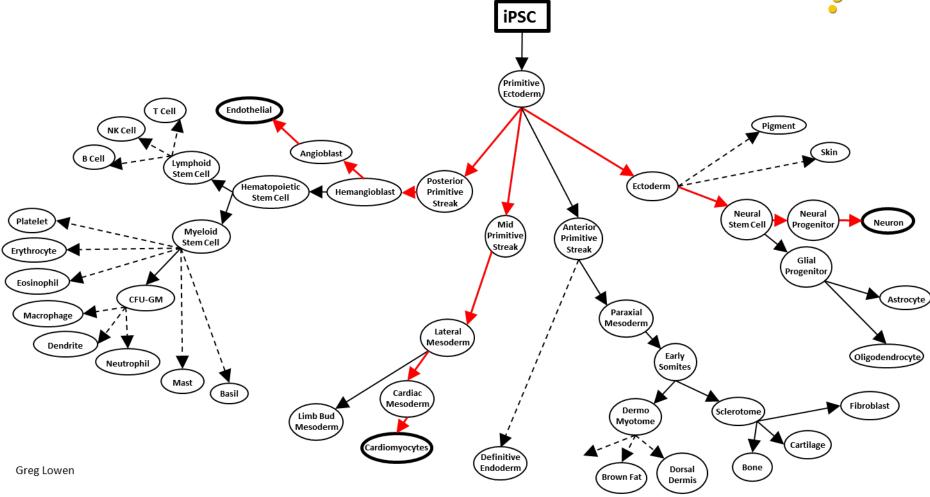






Controlling Cellular Differentiation





Where is Vanderbilt's tactical advantage for MicroFormulator and multi-omic control of cellular differentiation?

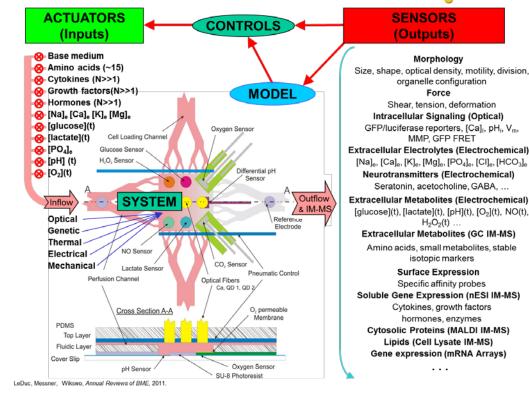
- Neurons?
- GI epithelium?

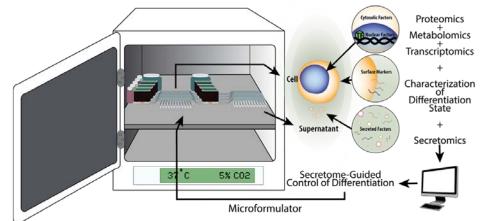


What else might we need?

VIJBRE

- Advances in mathematics
- Genetically coded fluorescent reporters
- Optogenetics
- Addressing cellular heterogeneity
 - Single-cell FISH
 - CRISPR-CAS with Single-cell RNAseq
 - Single-cell mass spectrometry







Almost done!

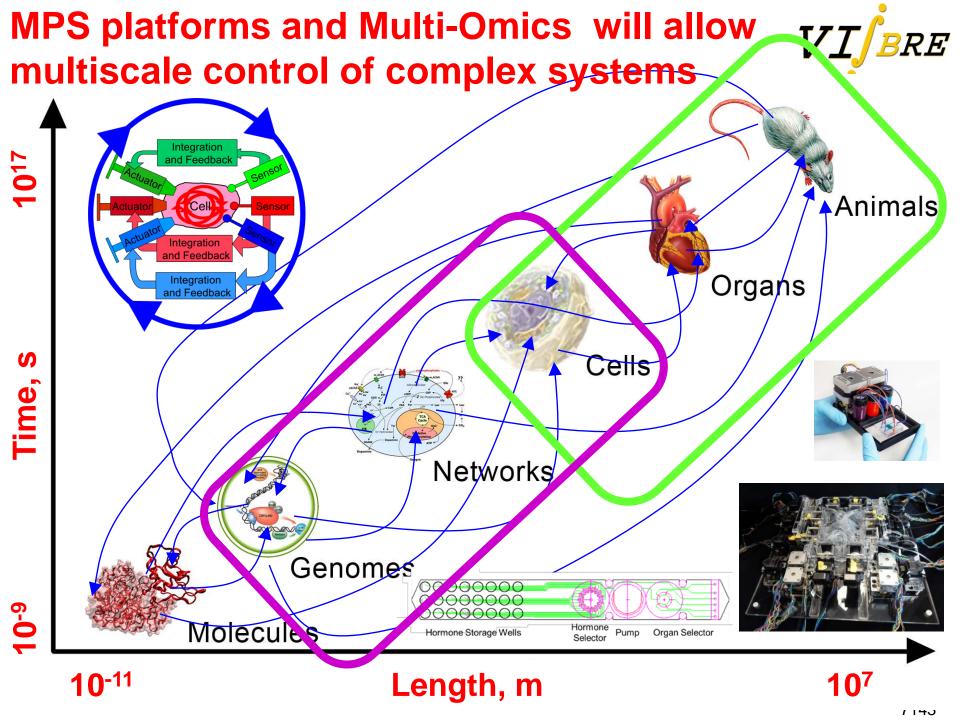
2012: Driving forces for the future of biology VI_{BRE}

- The need for more realistic in vitro experiments
 - Massively parallel, cellular microenvironments for the study of cell-cell, cellcell-drug, and cell-cell-drug-snp interactions
 - Real-time control of biological systems
- The need to control multiple parameters at the same time and measure multiple dynamic variables
 - Cell-scale sensors and actuators
 - Experiments that involve thousands of parameters
- The need to create complex, nonlinear models
 - New mathematics
 - Symbolic regression and exploration-estimation algorithms for machine learning in automated microbioreactors
 - Models to enable control of cellular responses and biomolecule production
- The need to raise research funds from more diverse sources
- The inability of the human mind (or at least those of the reviewers) to understand the complexity of what is being proposed and/or discovered

MicroFormulator

Organs on Chips

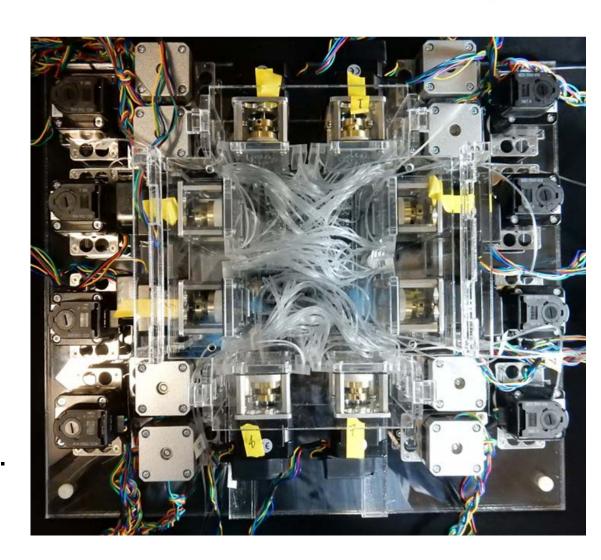
Graph Databases



JPW 2012: The bottom line...



- Ultimately, biology experiments may resemble particle physics experiments.
- Physicists have the requisite training and mindset compatible with large scale, automated biology, but are often bioignorant.
- Computational geometry and topology may be the new mathematics for biology.
- Should we teach
 Physics 101 and
 Topology 101 before
 Biology 101?



JPW January 2016: "Oh, shit. I've done it."

There is yet one more potential problem... VI



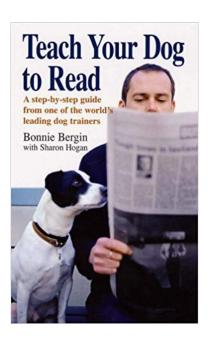
- We may not be able to understand what the computer tells us about biology.
- The next challenge is to create computers that can explain their findings to us....

 It might be as hopeless as explaining Shakespeare to a dog.

Hod Lipson, 2009

JPW 2012: All May Not Be Lost





2006

See Spot Read: Willow the Dog understands written commands

"....the dog can now sit up when a card says 'Sit Up,' plays dead when a card reads 'Bang,' and wave a paw when a sign says 'Wave.'"

http://www.peoplepets.com/news/amazing/see-spot-read-willow-the-dog-understands-written-commands/1



Really Hard Problems



However...

We do not have to <u>fully</u> understand a phenomenon to control or eliminate it. An effective model can accomplish a lot.

John Wikswo



Organs on Chips are highly controlled, interconnected *in vitro* human organ preparations that support intensive data acquisition and control not possible in humans.

Use them accordingly.

VIIBRE Organ-on-a-Chip Collaborators



Vanderbilt

- Vanessa Allwardt
- Frank Block III
- Frank Block Jr.
- Aaron Bowman
- Clayton Britt
- Jacquelyn Brown
- Young Chun
- David Cliffel
- Erica Curtis
- John Scott Daniels
- Jeffrey Davidson
- Anna Davis
- Mona Everheart
- William Fissell
- Greg Gerken
- Lucas Hofmeister
- William Hofmeister
- Orlando Hoilett
- Chaz Hong
- Shane Hutson
- Deyu Li
- Chee Lim
- Ethan Lippmann
- Dmitry Markov

- William Matloff
- Lisa McCawley
- Jennifer McKenzie
- BethAnn McLaughlin
- John McLean
- Dusty Miller
- Karoly Mirnics
- Nicole Muszynski
- Diana Neely
- Kevin Niswender
- Virginia Pensabene
- Ronald Reiserer
- Philip Samson
- David Schaffer
- Kevin Seale
- Stacy Sherrod
- Mingjian Shi
- Matthew Shotwell
- Veniamin Sidorov
- Hak-Joon Sung
- David Tabb
- Adam Travis
- Donna Webb
- Hendrik Weitkamp
- Erik Werner
- John Wikswo

AstraZeneca

 Matthew Wagoner, Jay Mettetal, Kristin Fabre, Aditya Kolli, Sudhir Deosarkar

CFD Research Corporation

- Kapil Pant, Prabhakar Pandian
- Andrzej Przekwas

Charite Hospital, Berlin (2012-2014)

 Katrin Zeilinger, Marc Lubberstadt, Fanny Knöspel

Cleveland Clinic and Flocel Inc.

- Michael Deblock, Kyle Lopin, and Chaitali Ghosh
- Damir Janigro (Flocel)

Harvard/Wyss (2011-2015)

 Don Ingber, Kit Parker, Josh Goss, Geraldine Hamilton, Danny Levner

Johns Hopkins University

Mark Donowitz

Los Alamos National Laboratory (2012-2014)

Rashi Iyer

University of Pittsburgh

• Lans Taylor, Albert Gough, Lawrence Vernetti

University of Texas Medical Branch

Mary Estes

University of Washington

Jonathan Himmelfarb

University of Wisconsin

William Murphy, William Daly

