

**SEAB TASK FORCE REPORT
ON BIOMEDICAL SCIENCES**

September 22, 2016

**Steven Koonin & Harold Varmus
Co-chairs**

CHARGE TO SEAB FROM SECRETARY MONIZ

(November 21, 2015)

“...form a new Task Force to identify:

- New areas for research by DOE investigators that could... significantly advance the pace of progress in biomedical sciences.
- New mechanisms for conducting research in coordination with scientists from government laboratories (both DOE and the NIH), universities, academic medical centers and industry....”

By September 2016, the Task Force should produce a report... available to the public, Congress, & the current and next Administration.

TASK FORCE MEMBERS

Steven Koonin* (co-chair), NYU
Harold Varmus* (co-chair), Cornell Med
Drew Endy, Stanford
Stuart Feldman, Schmidt Fund
Paula Hammond*, MIT
David Haussler, UCSC
Markus Meister, CalTech
David Piwnica-Worms, MD Anderson
Martha Schlicher*, Malinckrodt

WORKSHOPS

NYU Center for Urban Science, March 10-11, 2016
LBNL, Berkeley CA, July 18-19, 2016

PREMISES

- Biomedical sciences are vital to the nation
- Biomedical research depends on many disciplines
- Scientific responsibilities and agencies are imperfectly aligned
- Relevant technologies could be developed more efficiently
- Despite cultural differences, DOE and NIH are well suited to form partnerships

CULTURES OF THE TWO AGENCIES AND A BRIEF HISTORY OF INTERACTIONS

DOE

- Mission-driven activities
- Technology: primary mission
- Physical sciences
- Large team science
(National Labs, Hubs,
User Facilities, ARPA-E)
- Service to other agencies

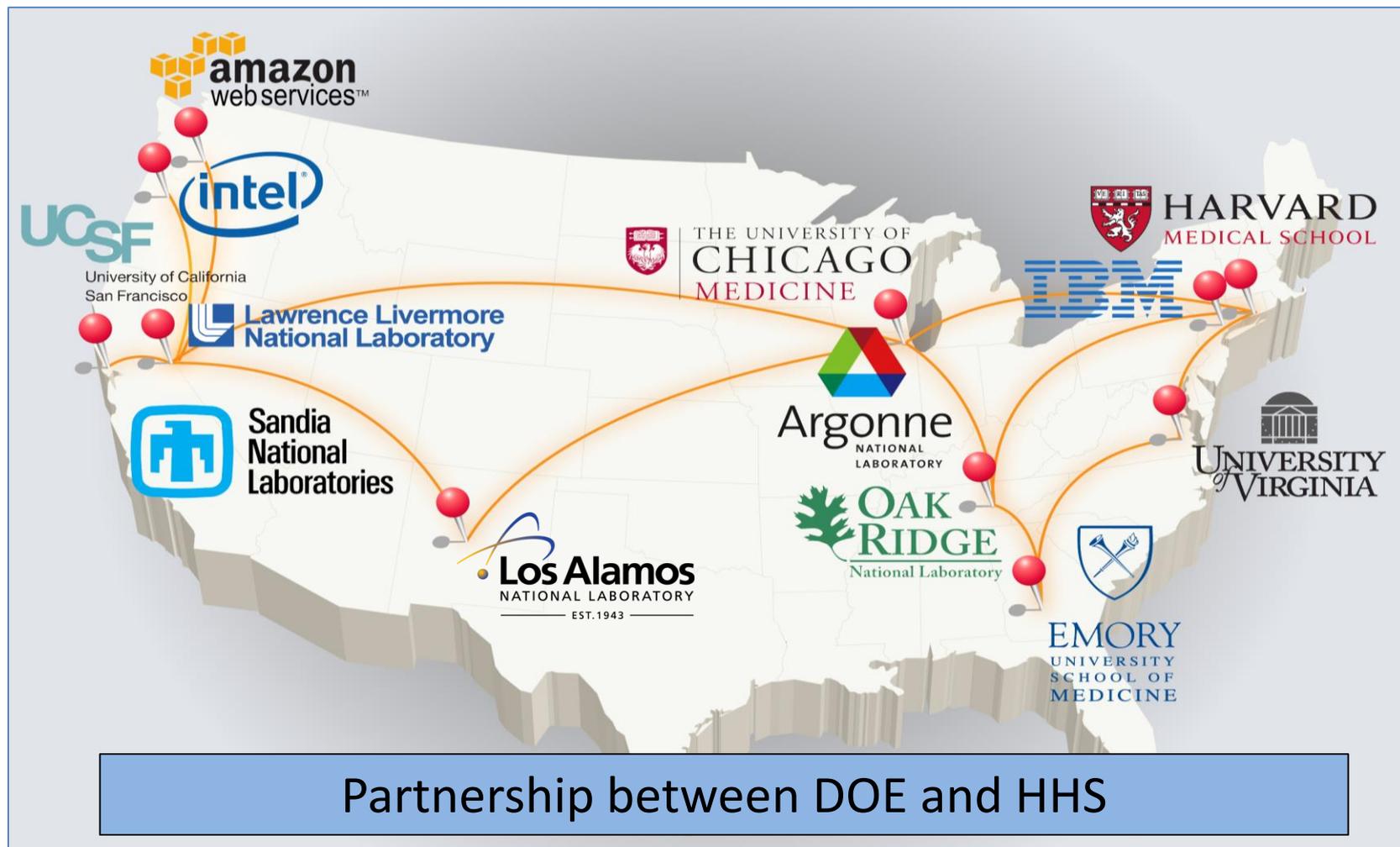
NIH

- Largely investigator-initiated, public health goals
- Technology: secondary
- Biology and medicine,
plus other disciplines
- Individual labs, small teams
- Extramural:Intramural =
10:1
- Large training component

COLLABORATIVE HISTORY:

- Human Genome Project
- DOE beam lines for structural biology
- Contributions to BRAIN, Precision Medicine...
- Other co-funded projects and plans

Initiative to build a national partnership in high performance computing and life sciences



CENTRAL MESSAGE

“With increasing recognition of the utility of DOE-supported technologies in biomedical research and the announcements of national goals for biomedicine, this is an appropriate time to propose the kinds of mechanisms that would make synergistic interactions between the agencies more frequent, less complicated, and more productive.”

CENTRAL GOALS

Expand the utility of DOE capabilities,
enhance the range and speed of NIH discoveries,
broaden perspectives in both agencies,
help to meet nation's scientific goals....

WHERE US GOVERNMENT GOALS FOR MEDICAL SCIENCE CONNECT WITH DOE CAPABILITIES

Administration's Biomedical Initiatives:

BRAIN

Precision Medicine

Cancer Moonshot

Alzheimer's Disease

Microbial drug resistance

Wide Expanse of Modern Research in Biology and Medicine:

Data analysis and management; simulations, etc.

Measurements: instruments, sensors, fabrication...

Images (large and small scale)

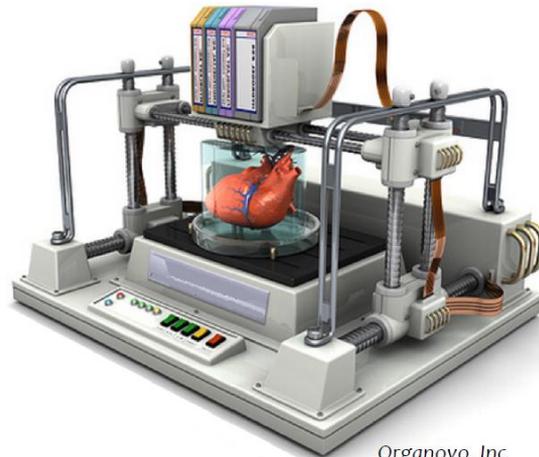
Materials for study, drug delivery, etc

High throughput methods (genomics, etc)

Radiobiology and biothreats

Health care delivery practices

Materials science



Organovo, Inc

Generate realistic tissue models with 3D geometry

Scalable to larger tissue constructs

Reproducible & automated technology

Precision positioning of cells

Nanoarchitecture design principles

- Steric congestion
- Sidechain chirality

Helix

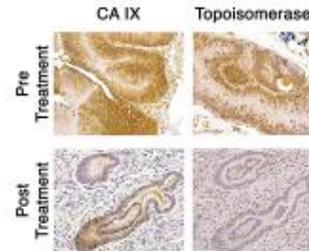
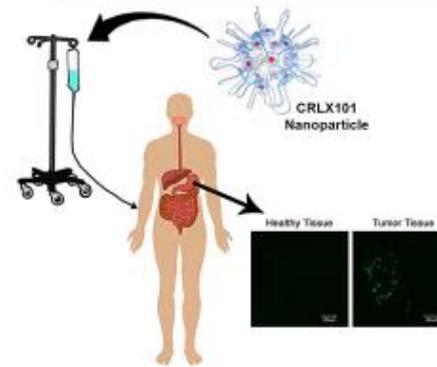
- Electrostatic, pi-pi
- Sequence periodicity
- Sigma strand
- Tiling

Sheet

- Size & shape congruence
- Chemically dissimilar blocks
- Tiling

Tube

Properly Designed Nanoparticles Can Function as Designed in Humans



Therapeutic Inhibits Intended Targets in Tumors of Humans

CRLX101 nanoparticles localize in human tumors and not in adjacent, nonneoplastic tissue after intravenous dosing

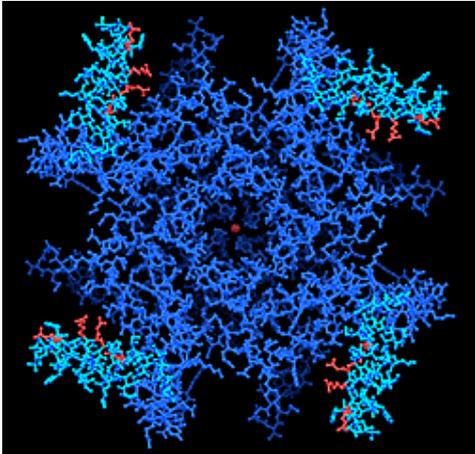
Andrew J. Clark¹, Devin T. Wiley¹, Jonathan E. Zuckerman^{2,3}, Paul Webster⁴, Joseph Chao⁵, James Liu⁶, Yan Yin^{6,7}, and Mark E. Davis^{1*}

¹Division of Chemistry and Chemical Engineering, California Institute of Technology, Pasadena, CA 91125; ²Department of Pathology and Laboratory Medicine, University of California, Los Angeles, CA 90095; ³Oak Crest Institute of Science, Pasadena, CA 91113; ⁴Department of Medical Oncology, City of Hope Comprehensive Cancer Center, Duarte, CA 91010; ⁵Division of Subcutaneous, City of Hope Comprehensive Cancer Center, Duarte, CA 91010; and ⁶Graduate Institute for Cancer Biology and Drug Discovery, Taipei Medical University, Taipei 11051, Taiwan

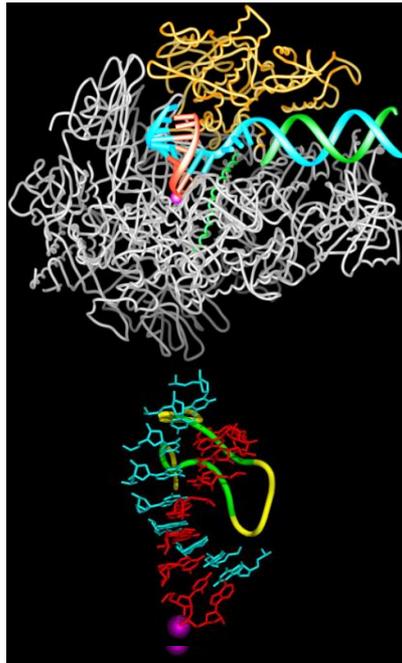
Impact of Synchrotron Structural Biology

- Enabling Nobel Prize Winning Research

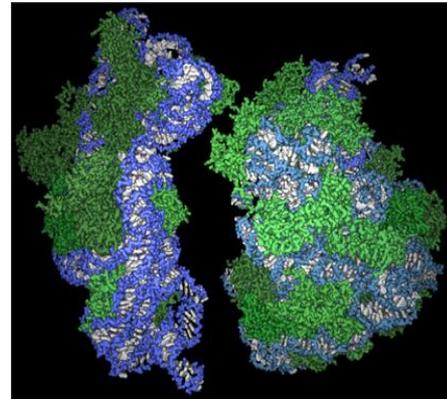
Membrane Channels, Polymerase, Ribosome, GPCRs



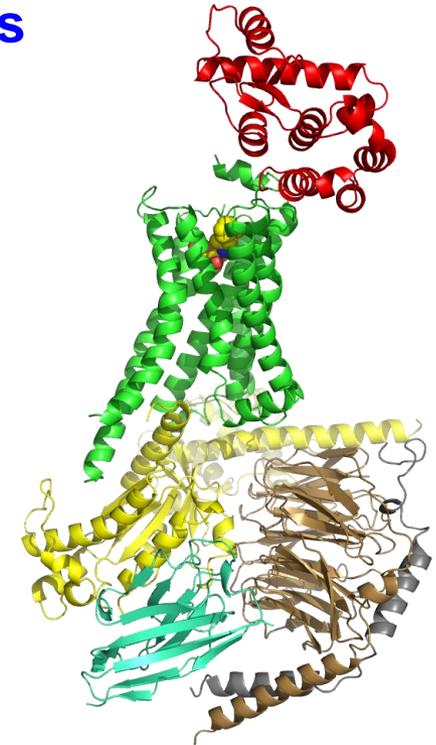
Roderick MacKinnon and Peter Agre – 2003 – K⁺ channel (KcsA)



Roger Kornberg – 2006 - RNA Polymerase II



Venki Ramakrishnan, Thomas Steitz and Ada Yonath – 2009 - Ribosome



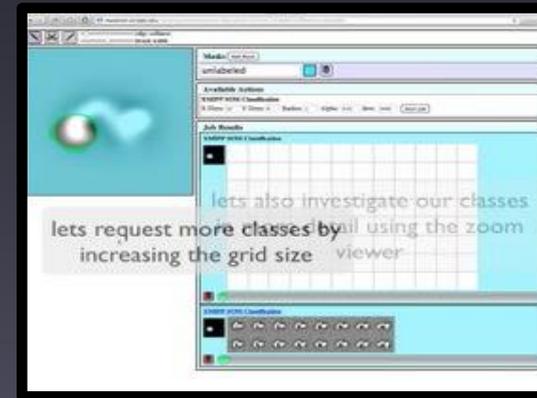
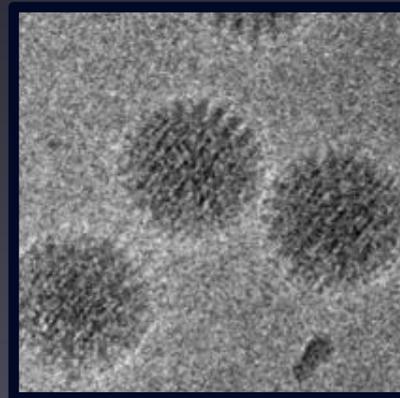
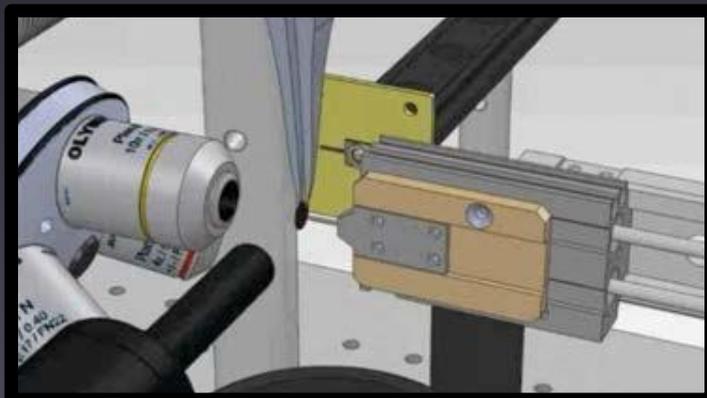
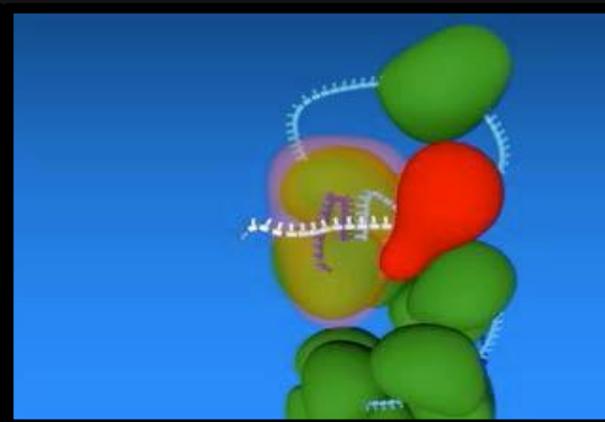
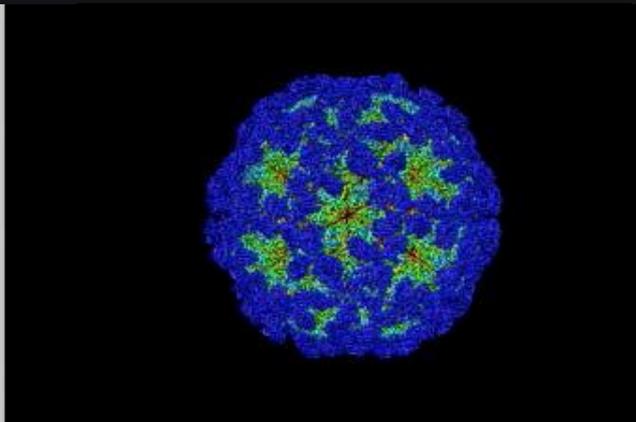
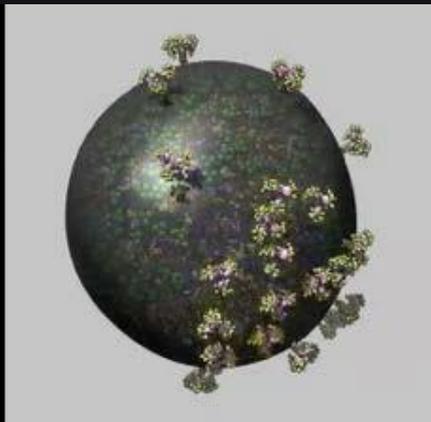
Brian Kobilka and Robert Lefkowitz – 2012 – β_2 AR- beta adrenergic receptor

Most challenging class of problems (often requiring significant effort and time) but yielding remarkable 'atomic level/chemical' insights into some of the most important processes in biology)



Using EM to Understand the Dynamics of Molecular Machines

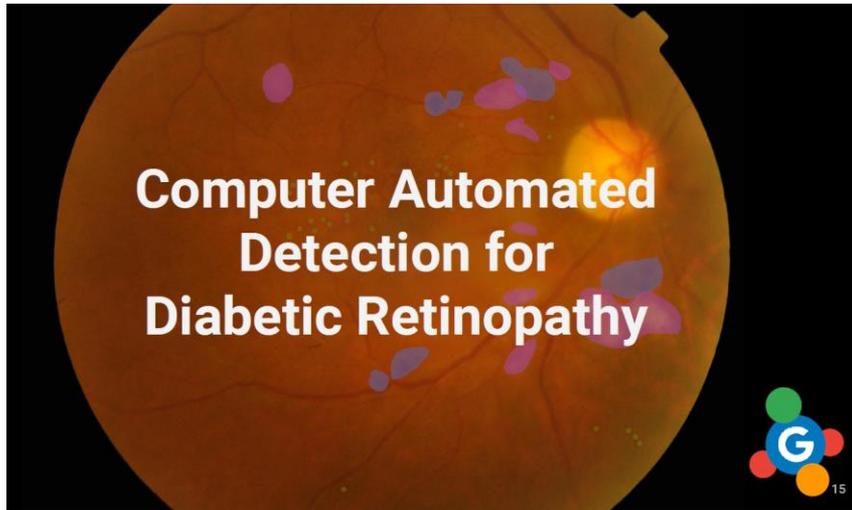
NEW YORK STRUCTURAL BIOLOGY CENTER
Simons Electron Microscopy Center



National Resource for Automated Molecular Microscopy
<http://nramm.nysbc.org>

SEAB Task
Force 10
March 2016

Computational and data sciences



Possible DOE computing role in BRAIN

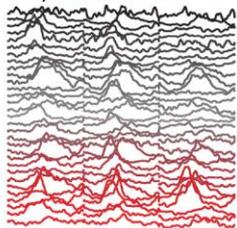


DOE can play a unique role in BRAIN computing through advances in applied mathematics and computer science together with HPC facilities.



Function

dynamic data



Theory & Models

abstractions

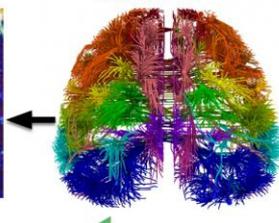
$$T(\xi) \cdot \frac{\partial}{\partial \theta} \ln U(\xi, \theta)$$

$$dx = \int \tau(x) \left(\frac{\partial}{\partial \theta} f(x, \theta) \right) dx$$

$$\theta dx = \int \frac{\partial}{\partial \theta} \tau(x) f(x, \theta) dx$$

Structure

static data



Generation and analysis of raw data

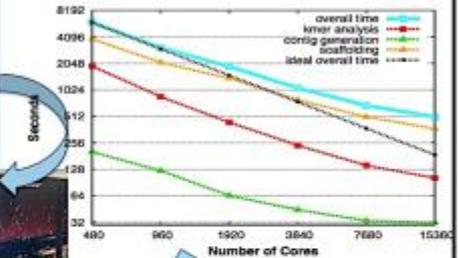
Linking structure to function is a 'grand challenge' in general biology and materials

"HiP-mer" high performance assembler



Computer Science HPC Expertise

- Remote Atomics
- Dynamic Aggregation
- Software Caching (sometimes)
- Clever algorithms and data structures (bloom filters, locality-aware hashing)
- Efficient languages (C vs Perl)
- Fast I/O



HPC systems with high speed interconnect networks

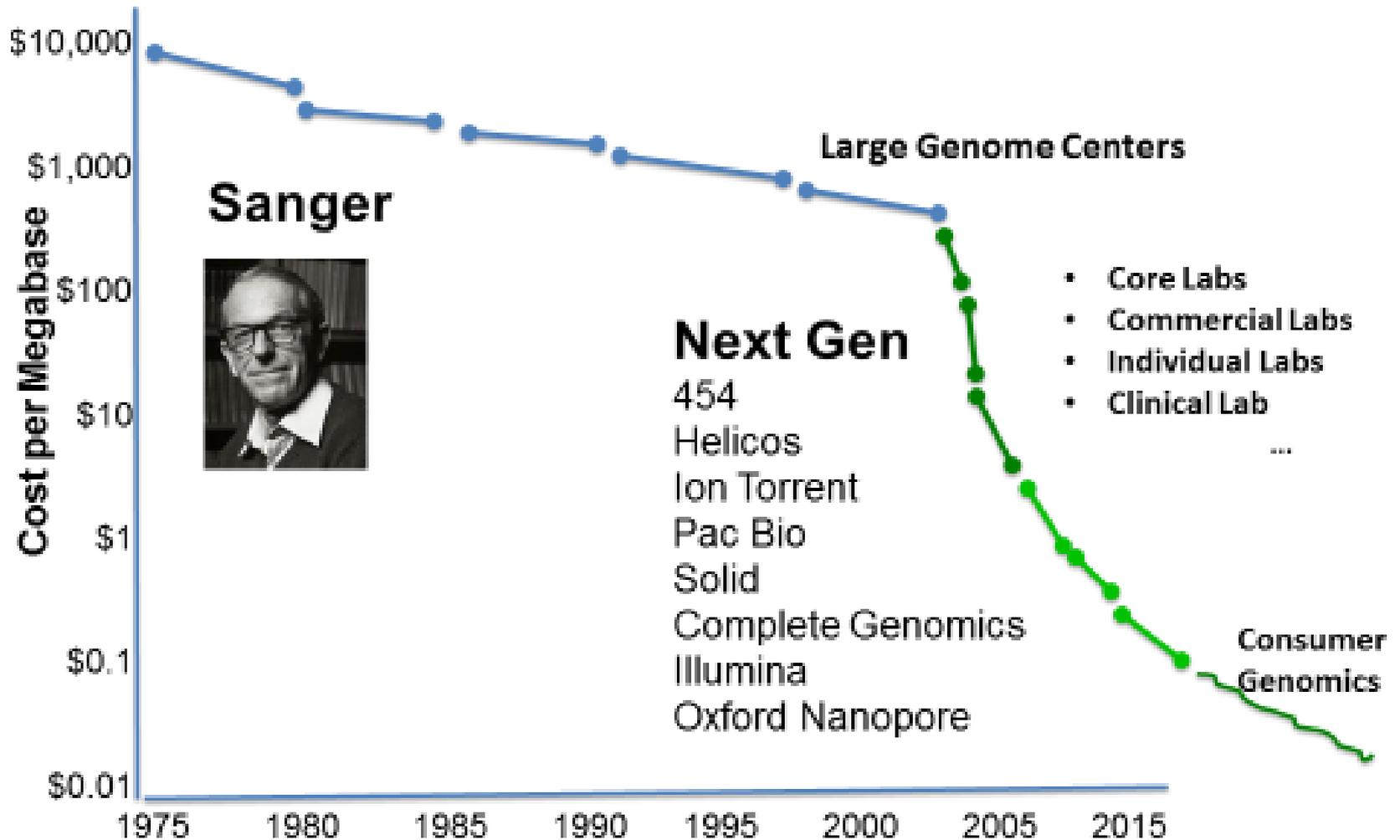
Grad student + software engineers needed to get to production

De novo Human genome assembly in 8 minutes! (15,000 cores)
Enables assembly of wheat (6x size of human!) in 40 minutes

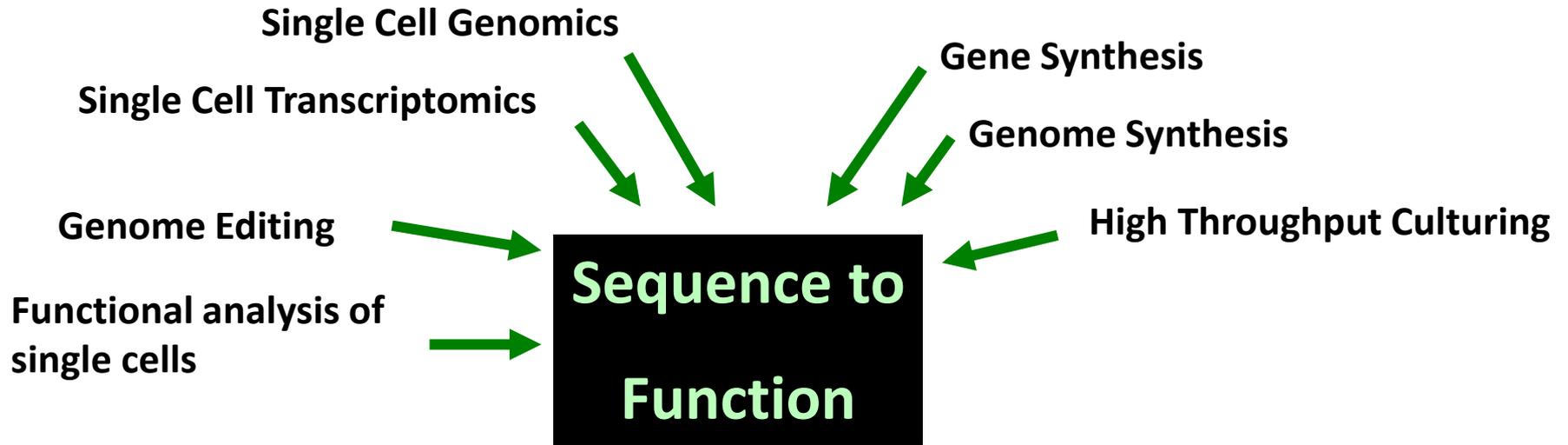


DNA science

Sequencing Technology Eras

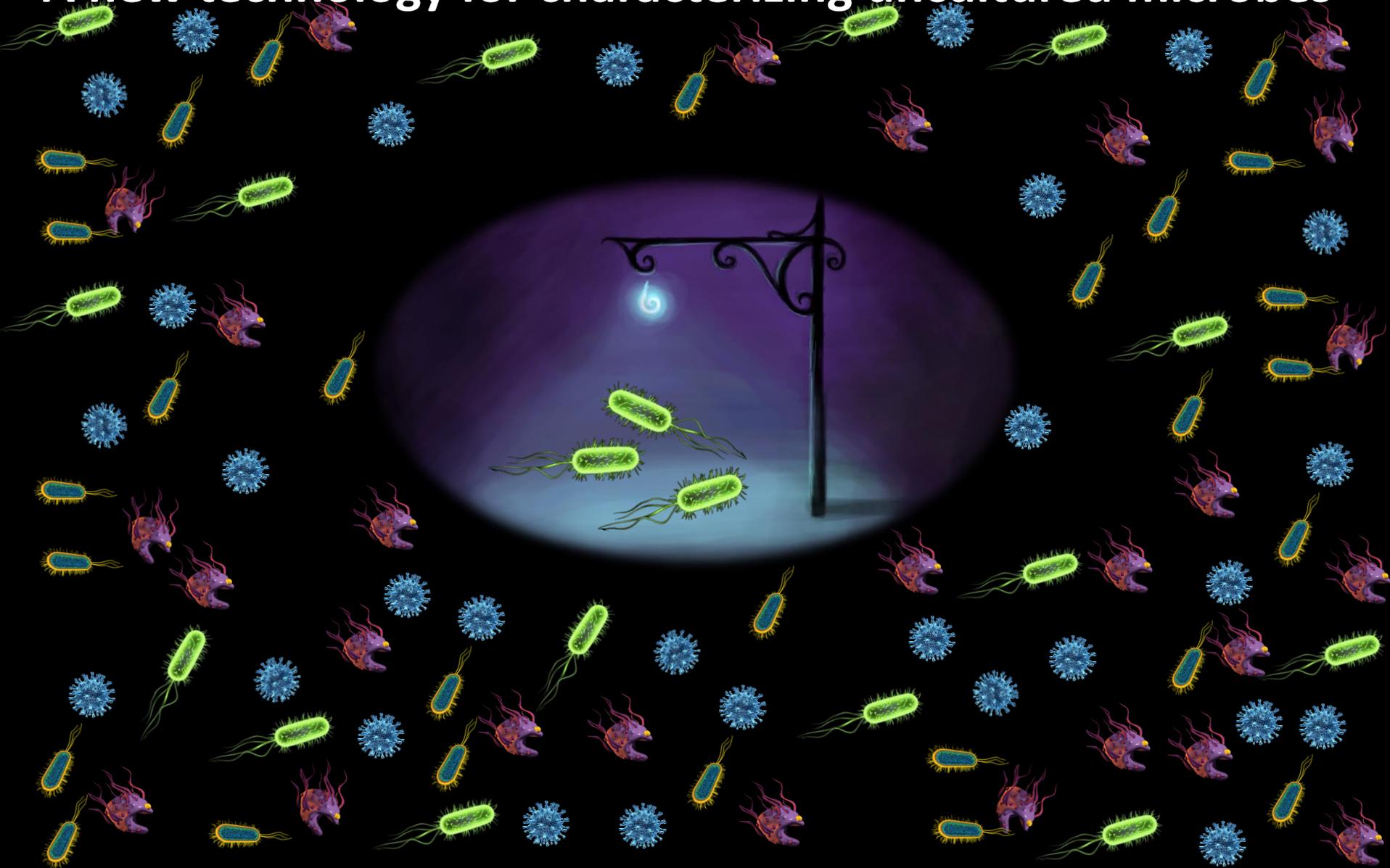


Genomic Sequence to Function: Capabilities

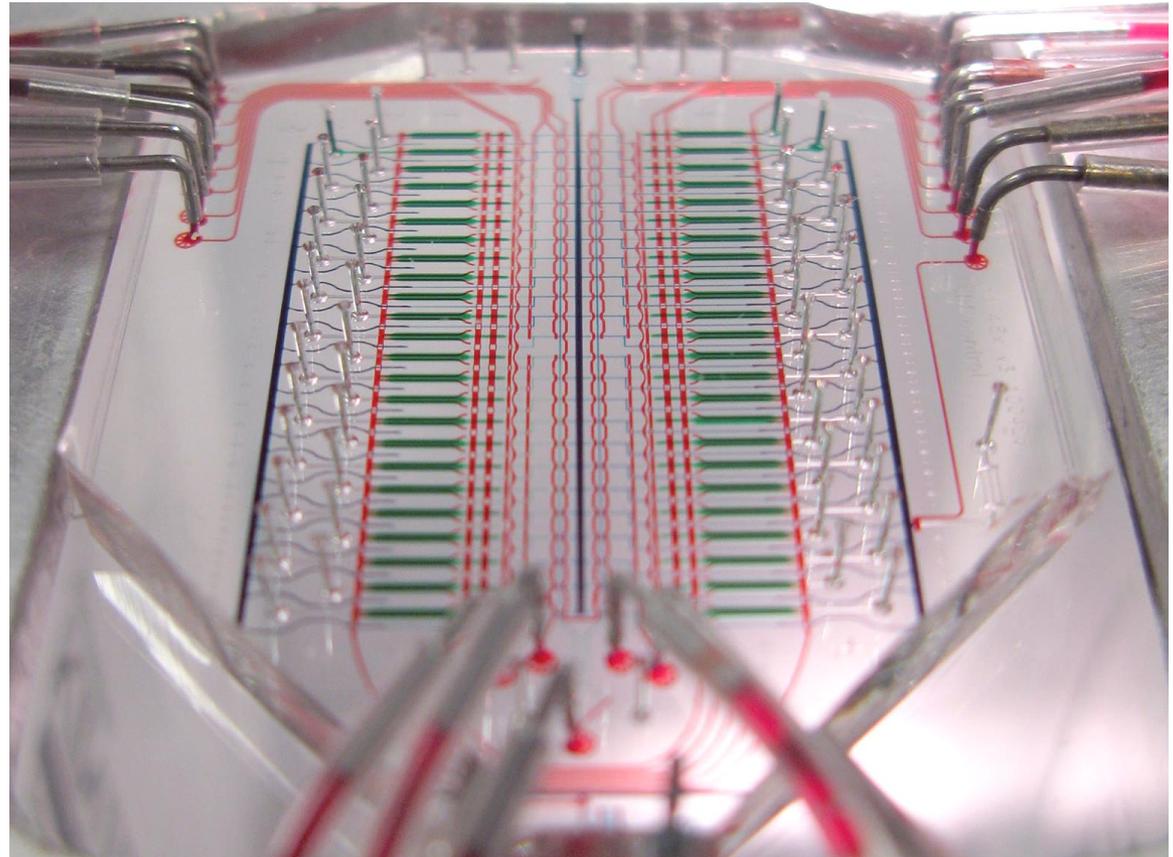
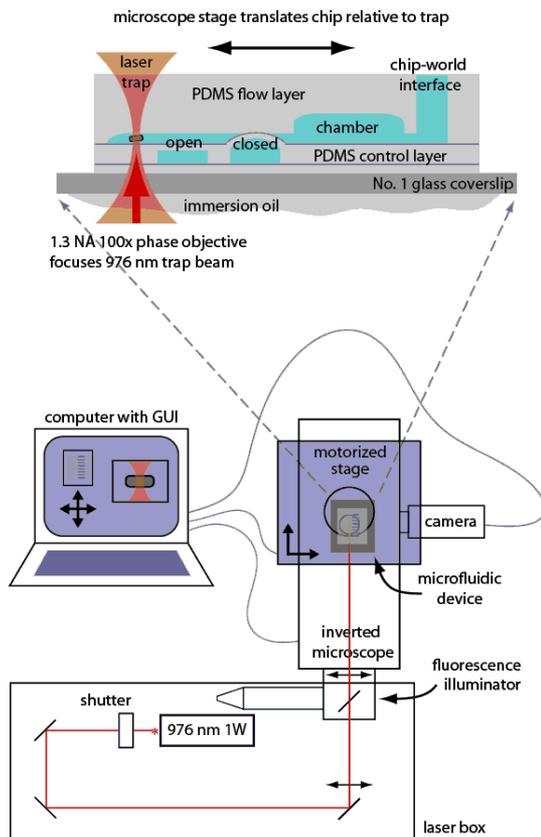


Opportunities of Massive Scale Sequencing

A new technology for characterizing uncultured microbes



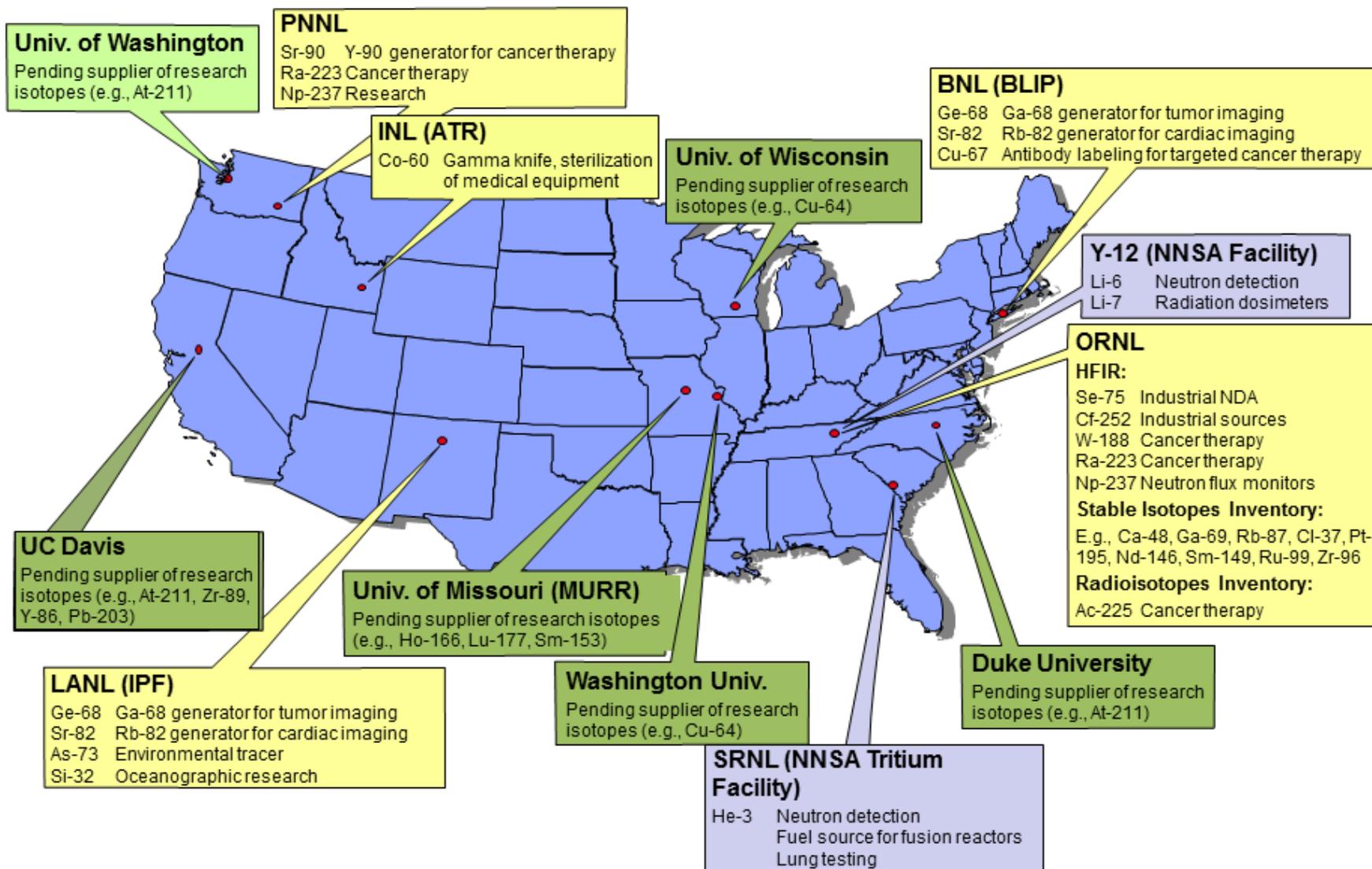
micro-optofluidic cell sorter & MDA amplifier



- reduces contamination (samples, reagents, environment)
- automates reaction set-up with low reagent consumption
- 1 μL sample OK
- aggregated/sticky sample OK

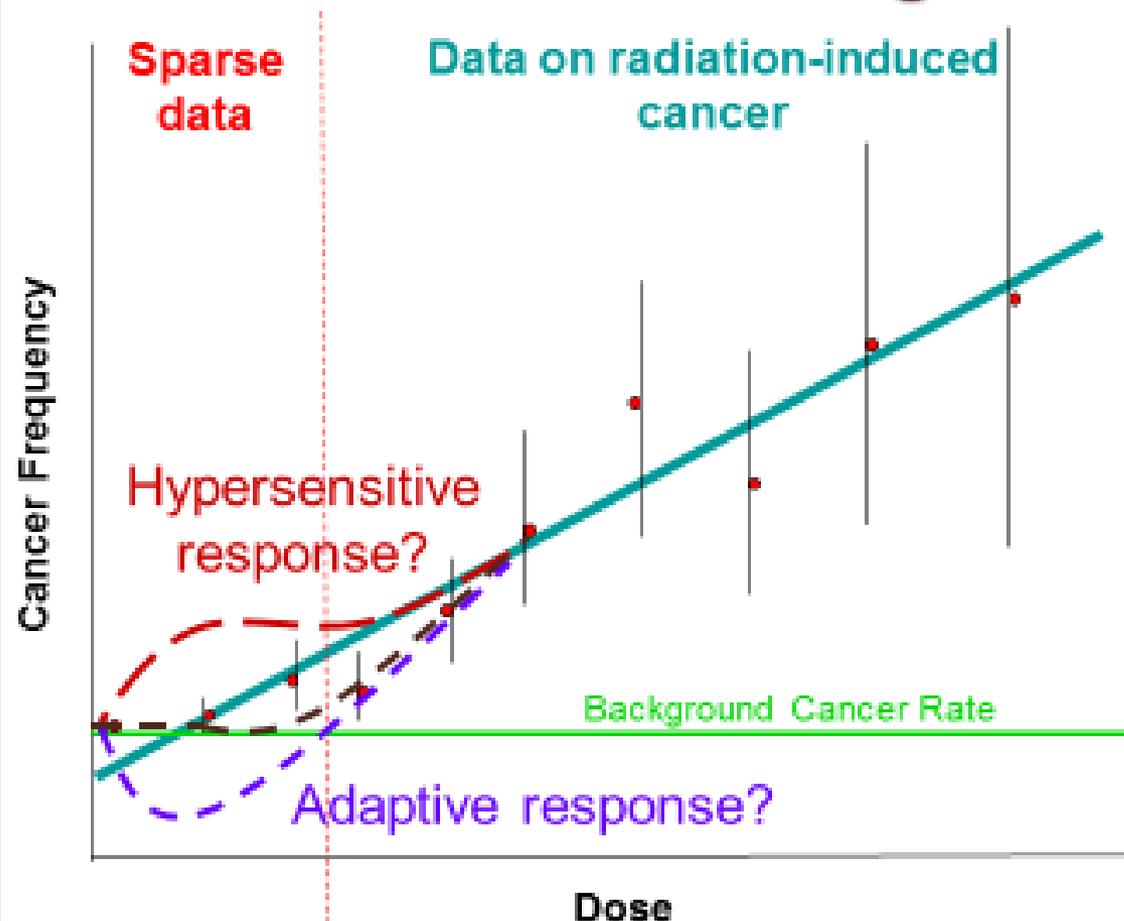
- compatible with optical microscopes
- cell concentrations 10^3 /mL - 10^8 /mL OK
- tiny & large cells OK
- low MDA bias
(Quake & Lasken et al, PLoS Genet 2007)

Isotopes and radio-biology: DOE and university production sites



Radiobiology

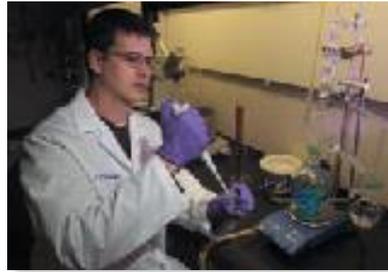
Linear Non-Threshold Radiation Dose-Response Model – Good for High Doses



The NNSA labs have played key roles in all facets of biodefense for decades



Threat Awareness



Prevention and Protection



Surveillance and Detection

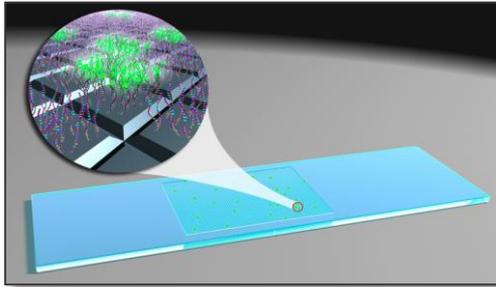


Response and Recovery *

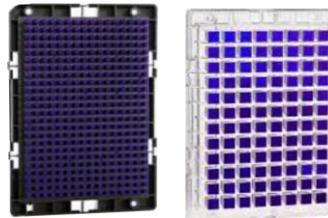


* Four "Pillars" from HSPD 10: Biodefense for the 21st Century

High Density DNA Microarray That Detects Over 10,000 Microbial Species for Wide Range of Applications

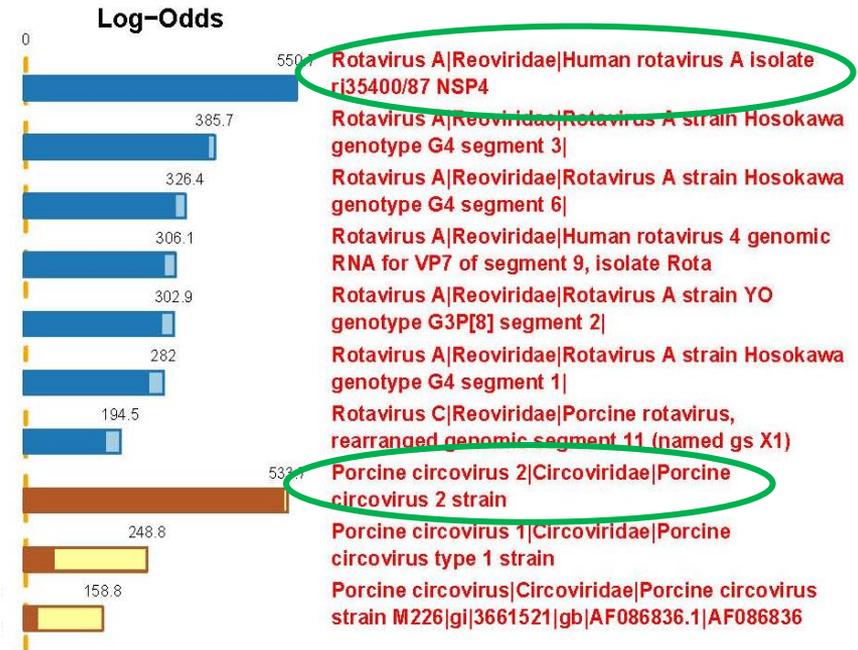


15 years of R&D and applications



384- and 96-sample array formats

Domain	# families	# species	# sequences*
Archaea	29	293	517
Bacteria	265	5,367	24,945
Fungi	101	265	395
Protozoa	32	117	191
Virus	94	4,219	86,931
Total	521	10,261	112,979



Biodefense versions of the arrays include markers for virulence, antibiotic resistance, genetic engineering and other forensic markers.

(D. Lindner, NNSA)

High throughput, high information content and low cost format enables new approaches in surveillance systems

RECOMMENDATIONS AND IMPLEMENTATION PLANS

#1 Identify areas for joint research programs

--DOE and NIH should impanel experts for this purpose on a regular basis

A bottoms-up approach to collaborative work!

RECOMMENDATIONS AND IMPLEMENTATION PLANS

#2 Bring diverse researchers together and co-train the young

- Cross-agency assignments
- Summer gatherings
- Grant supplements
- Novel training programs

Overcoming cultural differences!

RECOMMENDATIONS AND IMPLEMENTATION PLANS

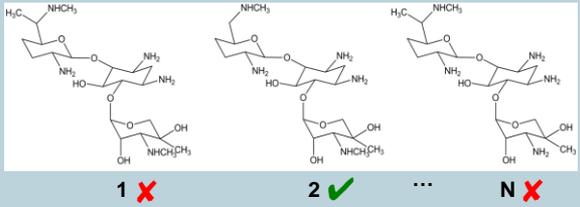
#3 Establish facilities, such as "foundaries," for desirable large scale collaborative projects

- Recognize the virtues of past efforts
- Act on recommendations of informed panels
- Negotiate with relevant parties

Biofoundry: Rapid Production of Antimicrobials



New antibiotic-resistant pathogen



Screen drug variants for efficacy



Stockpiled vials of cells to produce drug variants



Distributed fermentation drug production facilities



Rapid surge production of effective drug variant

Grand Challenge:

- Discover new and improved antimicrobials for human, animal, and plant pathogens
- Rapidly identify an effective antibiotic and surge its production at distributed sites



Office of Science

High Impact Science by Users and Staff...

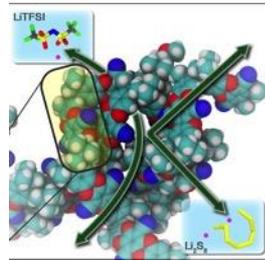
FY 2015

Budget: \$26.4M

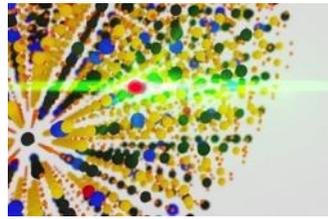
Publications: 326
(41% "high impact")

Proposals: 515

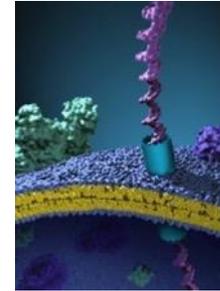
Users: 677



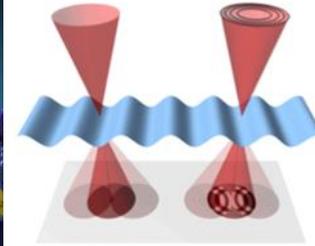
C. Li, *et al.*, *Nano Lett.* (2015)



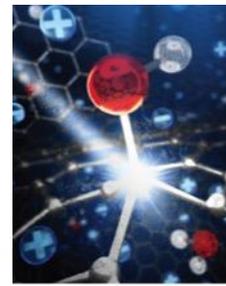
D. J. Gargas, *et al.* *Nature Nano.* (2014)



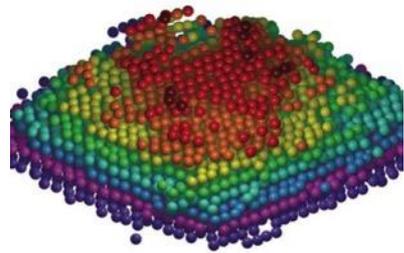
J. Geng *et al.*, *Nature* (2014)



C. Ophus, *et al.* *Nat Commun.* 2016



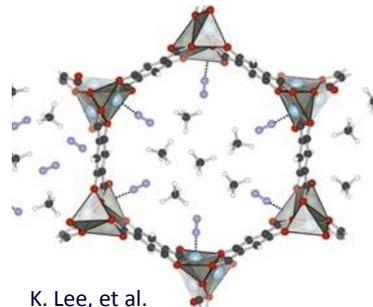
M. Bagge-Hansen, *et al.*, *Adv. Mater.* (2015)



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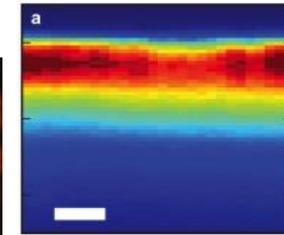
A. Llordés, *et al.* *Nature* (2013)



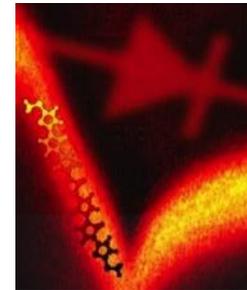
K. Lee, *et al.* *J. Am. Chem. Soc.* (2013)



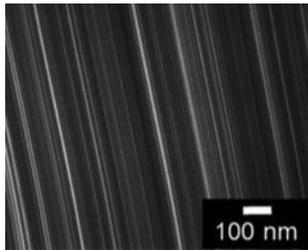
S. Barja, *et al.* *Nature Phys.* (2016)



E. S. Barnard, *et al.* *Scientific Reports* (2013)



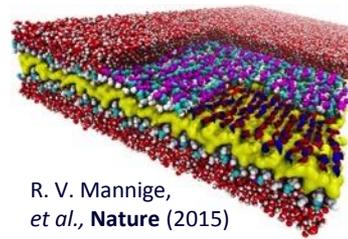
B. Capozzi, *et al.* *Nature Nano.* (2015)



S. Babin, *et al.* *J. Vac. Sci. Technol. B* (2015)
V. V. Yashchuk, *et al.* *Rev. Sci. Instrum.* (2015)



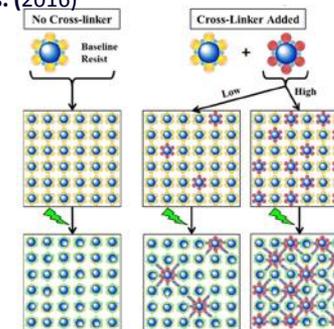
E. S. Cho, *et al.* *ACS Macro Lett.* (2015)



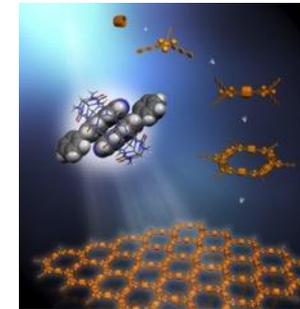
R. V. Mannige, *et al.*, *Nature* (2015)



Mecklenburg *et al.*, *Science* (2015)



P. K. Kulshreshtha, *et al.* *Nanotechnology* (2014)



K. D. Zhang, *et al.* *J. Am. Chem. Soc.* (2013)

...at LBL Molecular Foundry

RECOMMENDATIONS AND IMPLEMENTATION PLANS

#4 Inform OMB, Congress, and the public about strategies, proposed activities, and virtues of enhanced collaboration

ACKNOWLEDGEMENTS

TASK FORCE MEMBERS

WORKSHOP PARTICIPANTS

NIH/DOE STAFF

SEAB (AND ITS CHAIRPERSON)

SECRETARY MONIZ AND DIRECTOR COLLINS