

# High Performance Computing for Biology and Medicine

Srinivas Aluru

School of Computational Science and Engineering  
Georgia Institute of Technology

# Why High Performance Computing

- To solve compute-intensive problems in reasonable time limits
  - e.g. weather prediction, protein folding
- To solve data-intensive problems that don't fit in a single node
  - e.g. genomics, social network analysis
- To simulate what can't be done experimentally
  - e.g. simulating the galaxy

# The Big Data Challenge

Then (2005)



**ABI 3700**

96 ~800 bp reads  
76.8 X 10<sup>3</sup> bases  
~\$1 per kilo base

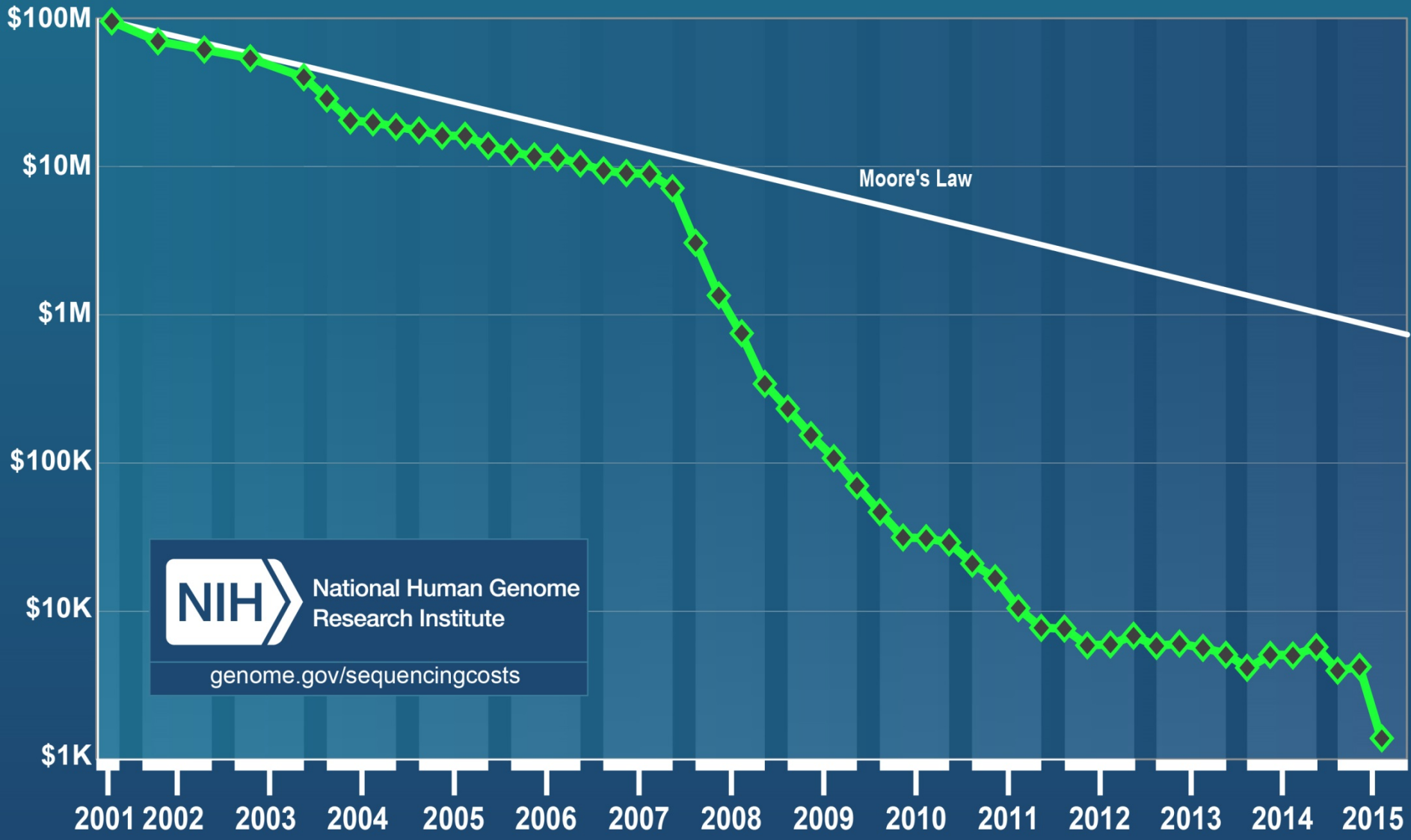
Now



**Illumina HiSeq 2500**

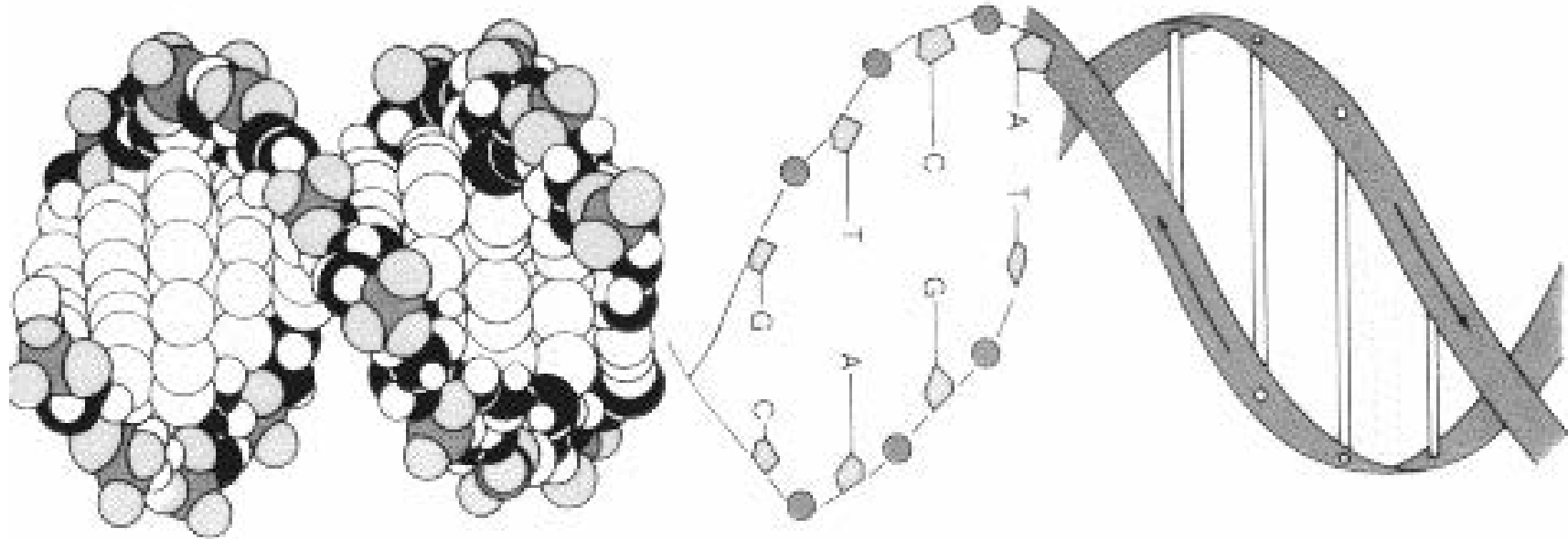
6 billion 100 bp reads  
600 X 10<sup>9</sup> bases  
~\$1 per 200 million  
bases

# Cost per Genome



**NIH** National Human Genome Research Institute  
[genome.gov/sequencingcosts](http://genome.gov/sequencingcosts)

# What is a Genome?



5' A T T C G G G A A T G C A T G C C A 3'  
3' T A A G C C C T T A C G T A C G G T 5'

# How Big are Genomes?

Viruses	50 kb+
Microbes	1 Mb+
Human/mouse/chimp	~3 Gb
Rice	450 Mb
Maize	2.5 Gb
Pine	19 Gb

# DNA to be used for storage?

- *Regenesis: How synthetic biology will reinvent nature and ourselves*
- 53,000 words, 11 pictures, and some javascript
- 70 billion copies in a test tube
- 5.5 Pbits per mm<sup>3</sup>, 10<sup>6</sup> times higher density than disk storage



# Human Genome Progress

- 1,000 genomes project
  - diverse anonymous individuals for study of human genetic variation
- Harvard's Personal Genome Project
  - 100,000 volunteers
  - no anonymity, limited to U.S.
- What are we looking for?
  - SNPs, indels, CNVs, disease associations
  - Race, Ethnic, Gender specific analysis
  - Genome evolution during diseases such as cancer

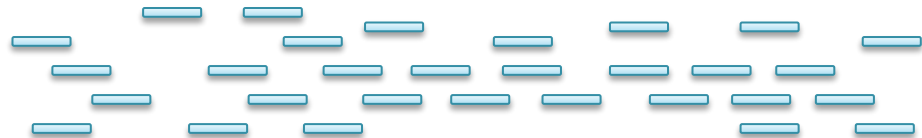
SNPs

... .. ACGTT**A**ACTTA ... .. ACGTT**C**ACGA ... .. GG**C**GAACA ... ..

... .. ACGTT**T**ACTTA ... .. ACGTT**G**ACGA ... .. GG**A**GAACA ... ..



# Genome Resequencing



50X of 3GB human genome  $\approx$  1.5 billion 100bp reads

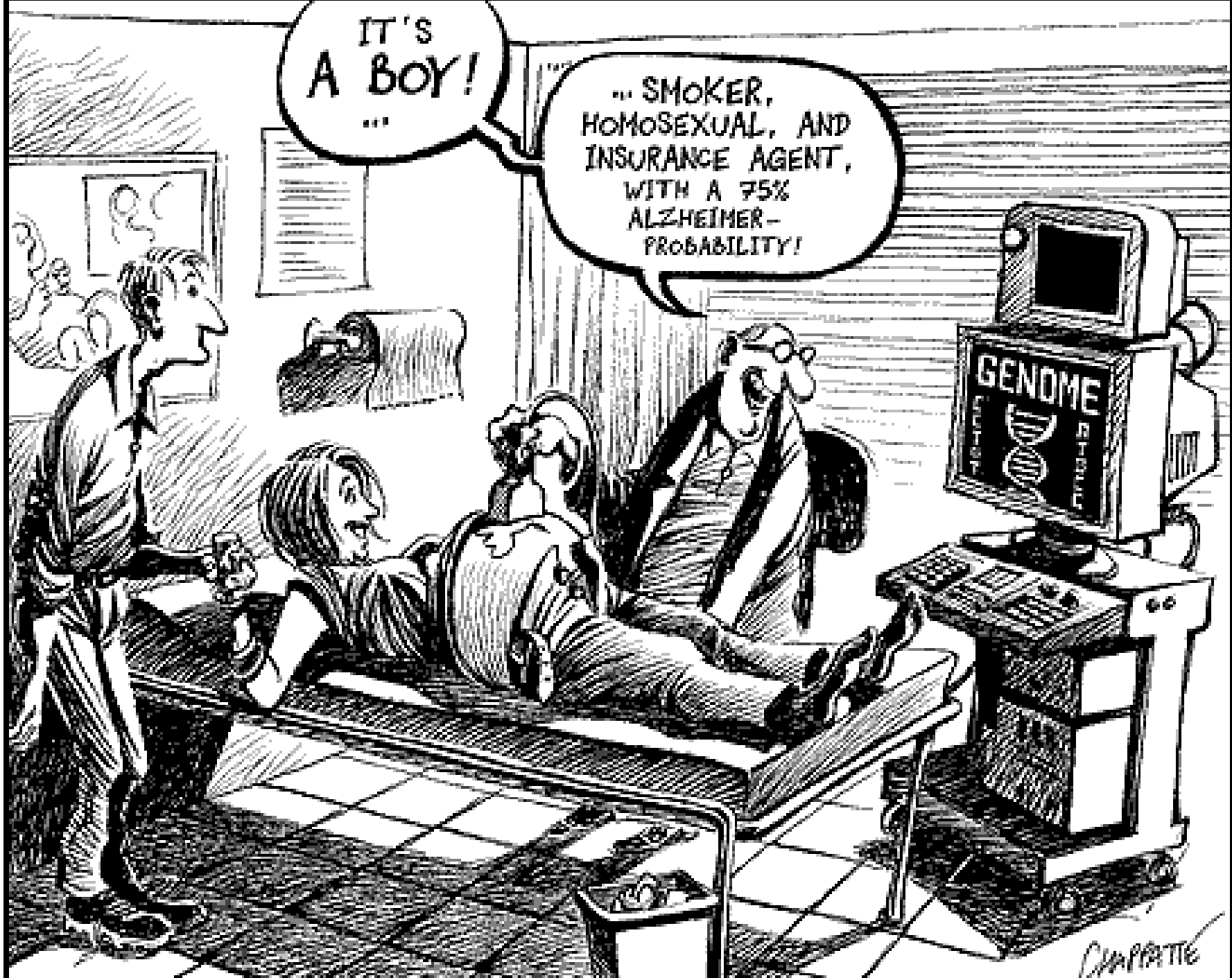
The diagram illustrates the concept of 50X coverage. It shows a horizontal line representing the reference genome. Above this line, numerous short horizontal bars represent sequencing reads. These reads are distributed across the length of the genome, with some overlapping, demonstrating the high density of data generated by 50X coverage.

Reference  
genome



IT'S  
A BOY!

... SMOKER,  
HOMOSEXUAL, AND  
INSURANCE AGENT,  
WITH A 75%  
ALZHEIMER-  
PROBABILITY!



CHAPATTE



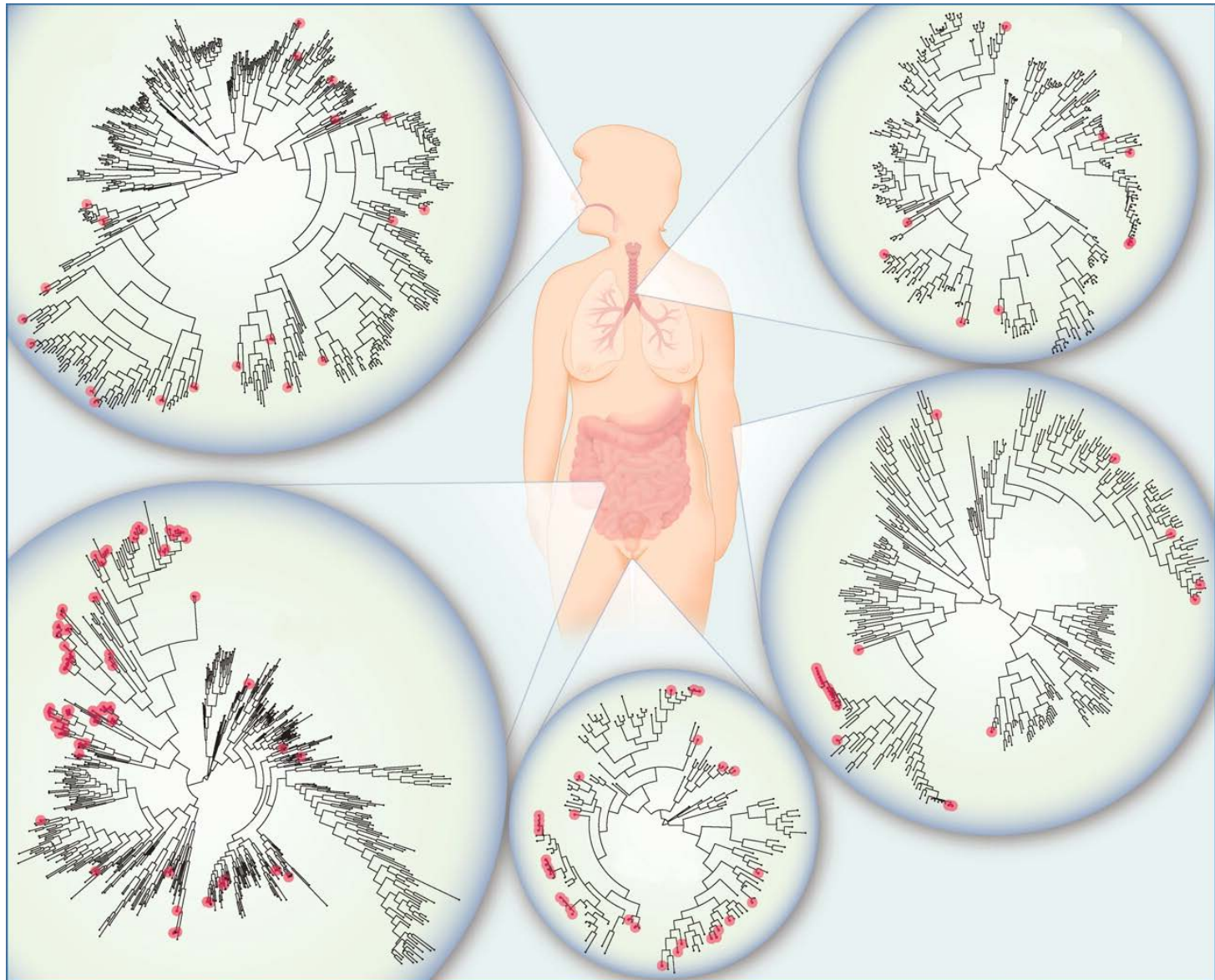
~40 million microbes/gram

~1 million/ml

$5 \times 10^{30}$  estimated  
on the planet



# Human Microbiome

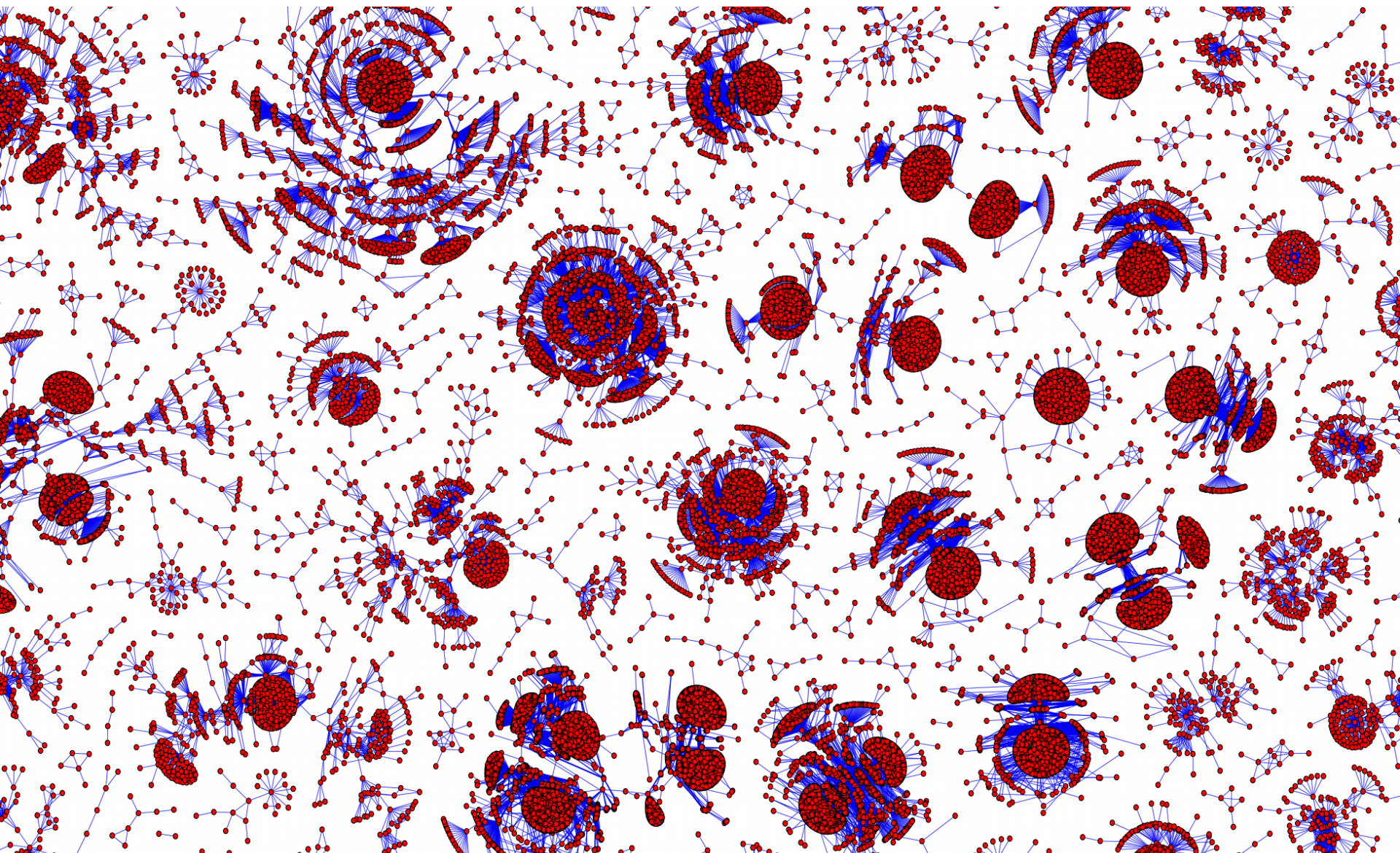




# You are host to ...

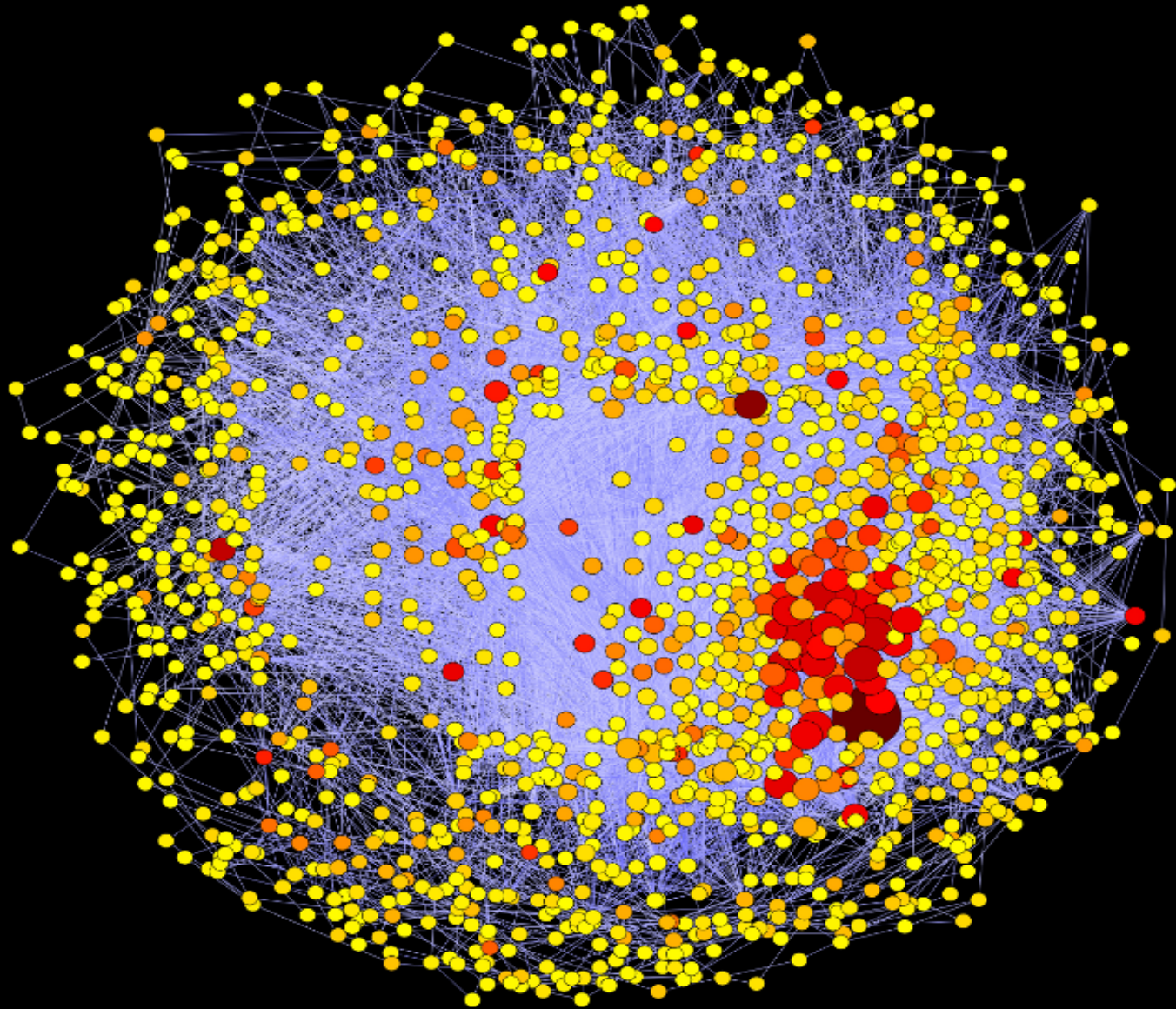
- 10,000 microbial species
- Outnumbering your cells 10 to 1
- About 5 lbs. of your body weight
- Over 8 million genes to your ~30,000

# Mouse Gut Microbiome





# Gene Interaction Networks



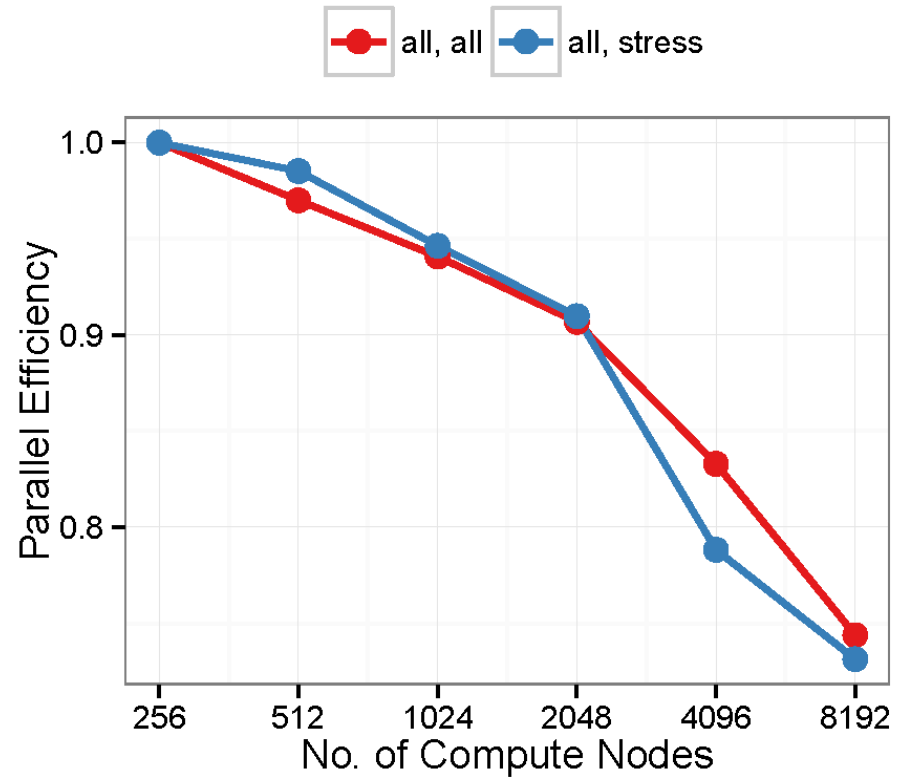
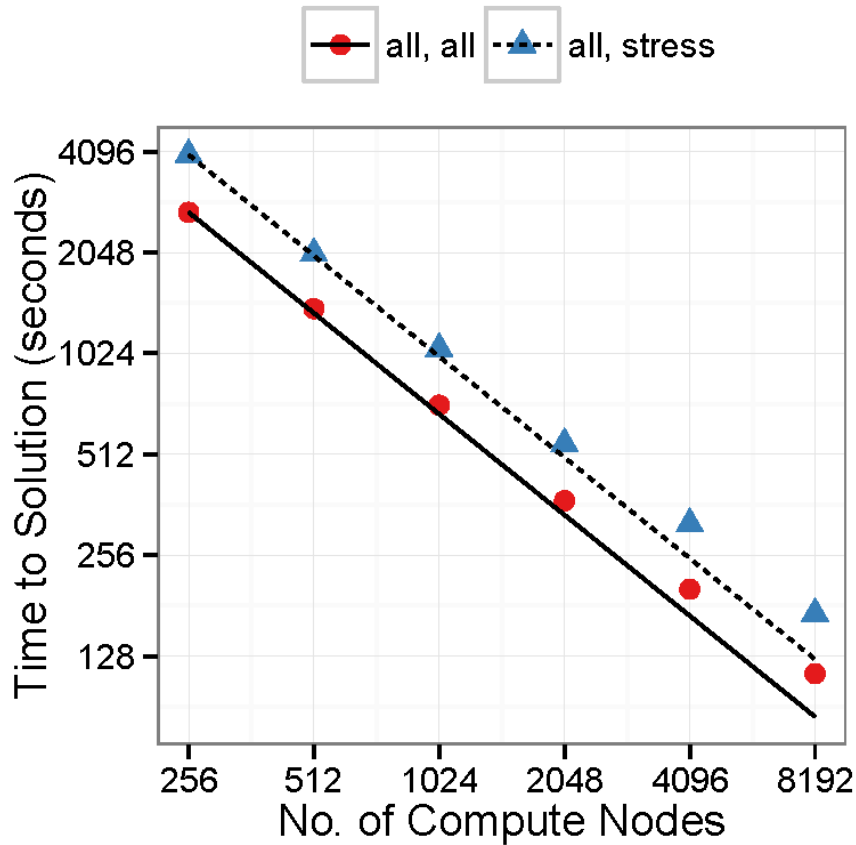
# Target Supercomputers

- Tianhe-2, National University of Defense Technology, Changsha.
- Stampede, Texas Advanced Computing Center, Austin

	Tianhe-2(54.9 PF)	Stampede(8.5 PF)
CPU	Intel Xeon E5-2600	Intel Xeon E5-2680
CPU Frequency	2.2 GHz	2.7 GHz
No. of CPUs	2	2
DRAM	64 GB	32 GB
Coprocessors	Intel Xeon Phi 31 S1P	Intel Xeon Phi SE10P
Coprocessors frequency	1.09 GHz	1.09 GHz
No. of Coprocessors	3	1
Coprocessor Memory	8 GB	8 GB
Cores per node	192 (2 * 12 + 3*56)	76 * (2*8 + 60)
Threads per node	696	256



# Parallel Efficiency





# Arabidopsis T-DNA Mutants

**Wild-Type**

**AT1G56500**

**AT5G07020**



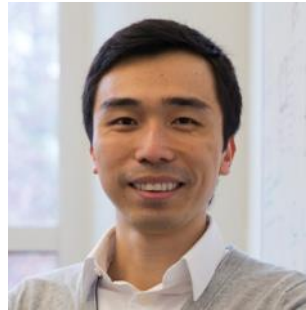
# Faculty: Interdisciplinary Innovators



**Srinivas Aluru**  
*Professor*



**David Bader**  
*Professor and Chair*



**Polo Chau**  
*Assistant Professor*



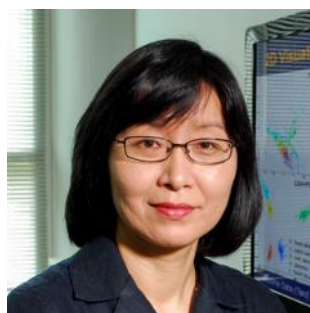
**Edmond Chow**  
*Associate Professor*



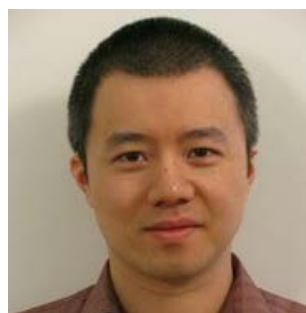
**Bistra Dilkina**  
*Assistant Professor*



**Richard Fujimoto**  
*Regents' Professor*



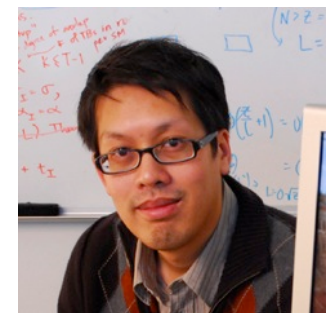
**Haesun Park**  
*Professor*



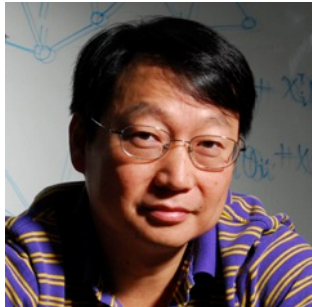
**Le Song**  
*Assistant Professor*



**Jimeng Sun**  
*Associate Professor*



**Richard Vuduc**  
*Associate Professor*



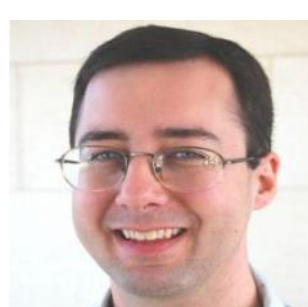
**Hongyuan Zha**  
*Professor*



**Kenneth Brown**  
*Chemistry*



**Mark Borodovsky**  
*BME*



**David Sherrill**  
*Chemistry*



**Surya Kalidindi**  
*Mech. Engr.*