

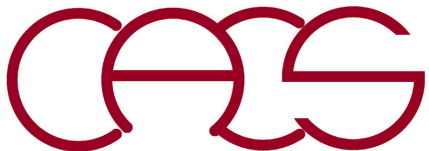
Molecular-Dynamics Machines

Aiichiro Nakano

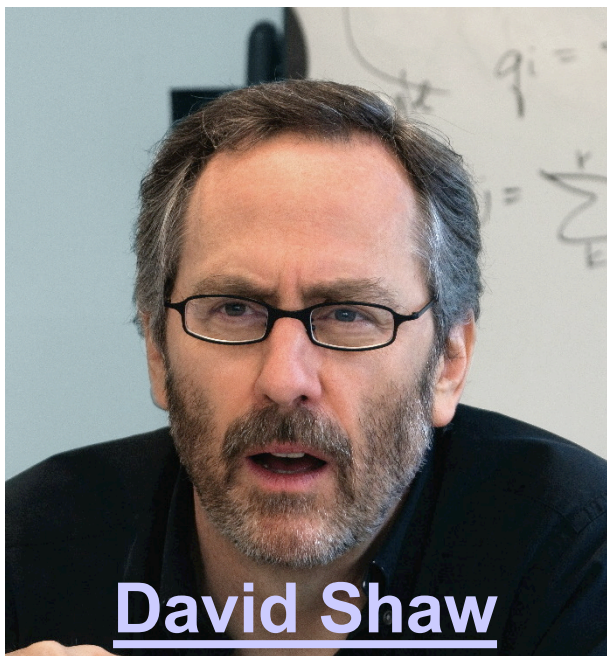
*Collaboratory for Advanced Computing & Simulations
Department of Computer Science
Department of Physics & Astronomy
Department of Quantitative & Computational Biology
University of Southern California*

Email: anakano@usc.edu

Why parallel MD? It's hot in computer science!



Anton: Computational Microscope



16 μ s/day simulation on 512 nodes
(5 μ s/step execution time)



“... make all these discoveries because they were looking at the world in a different way.”

Named after Anton van Leeuwenhoek, who is often referred to as “the father of microscopy”

“... there’s still a lot of juicy, low-hanging fruit in this (molecular simulation) area ...”

D E Shaw Research

A conversation with David E. Shaw, CACM 52(10), 49 ('09)

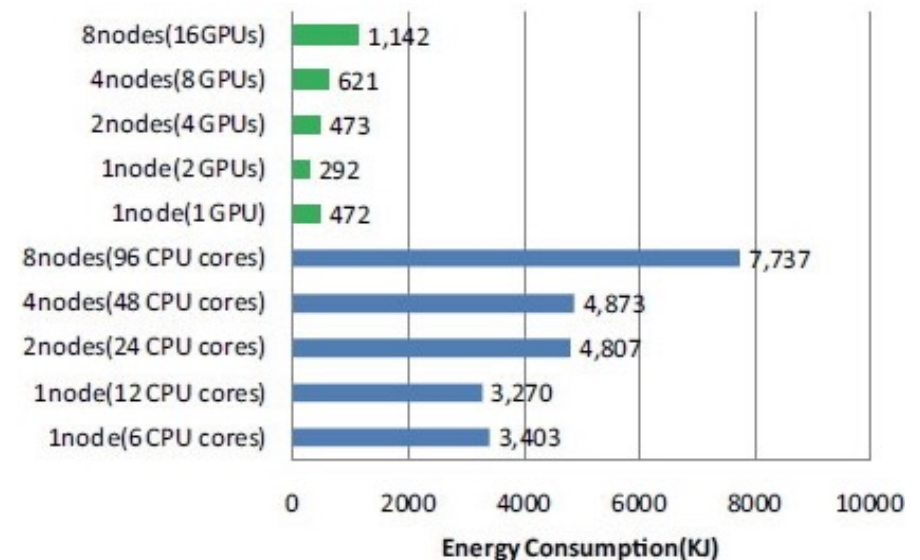
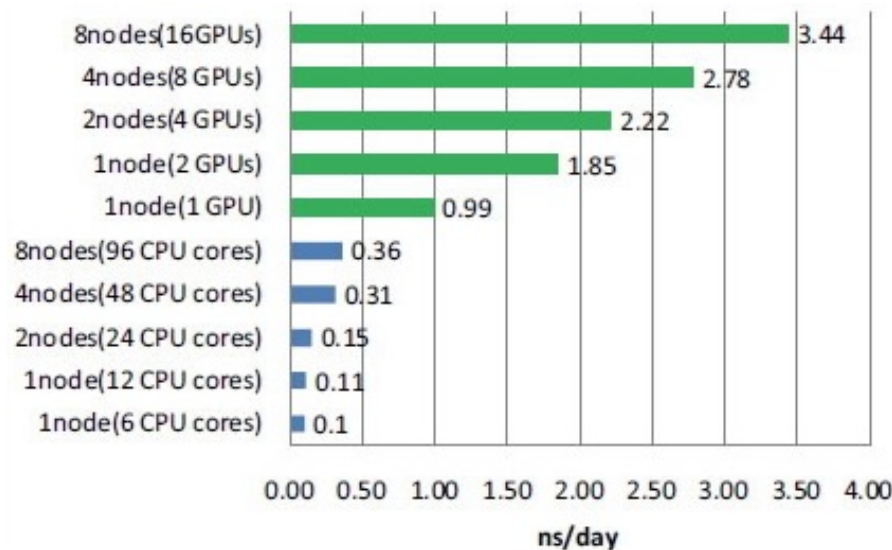
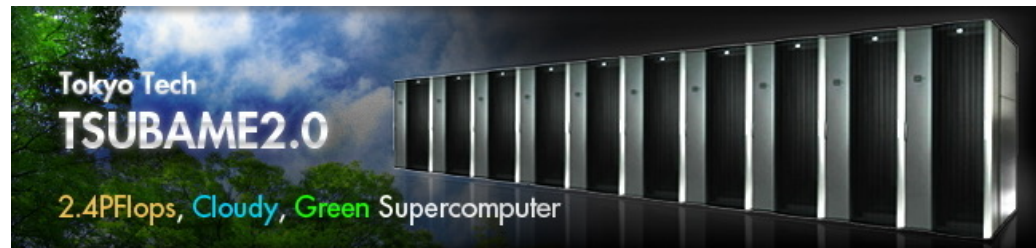
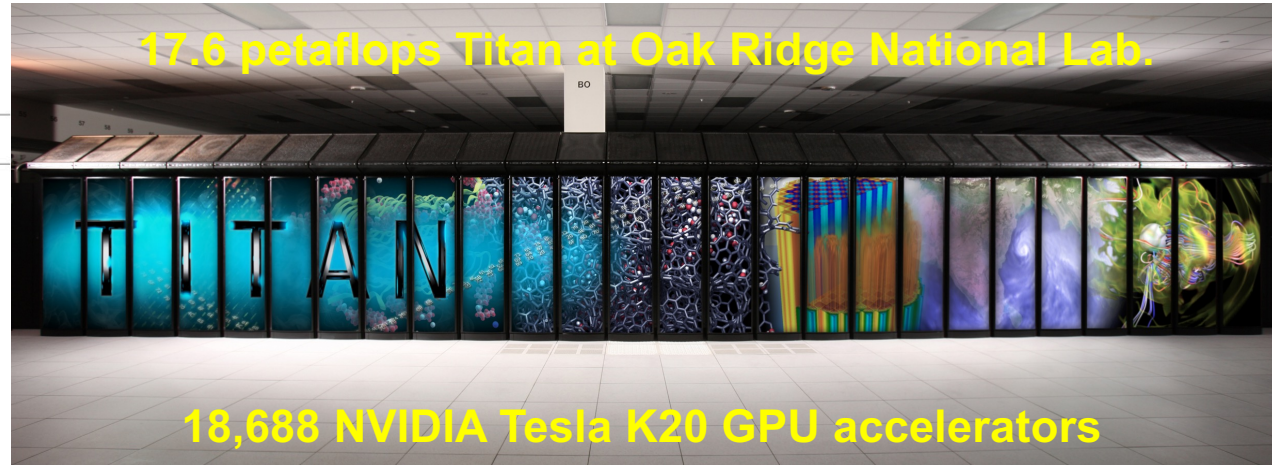
MD on GPU Clusters

GPU acceleration and other computer performance increases will offer critical benefits to biomedical science.

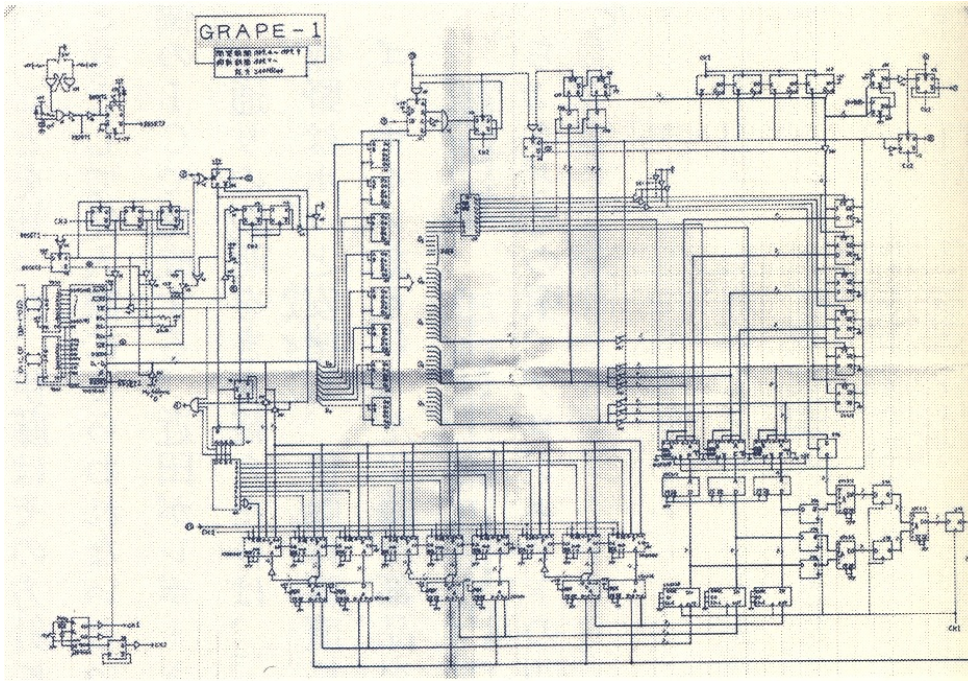
BY JAMES C. PHILLIPS AND JOHN E. STONE

Probing Biomolecular Machines with Graphics Processors

[CACM 52\(10\), 34 \('09\)](#)



GRAPE 1 (\$2K, 1989)



Tomoyoshi Ito & comics he authored

- **GRAPE (GRAVity PipE) = special-purpose computer for the gravitational N -body problem built by astrophysicists at Univ. of Tokyo**
- **GRAPE 1 designed by a 1st-year Ph.D. student (with \$140K/year income)**

Gravitational Pipeline

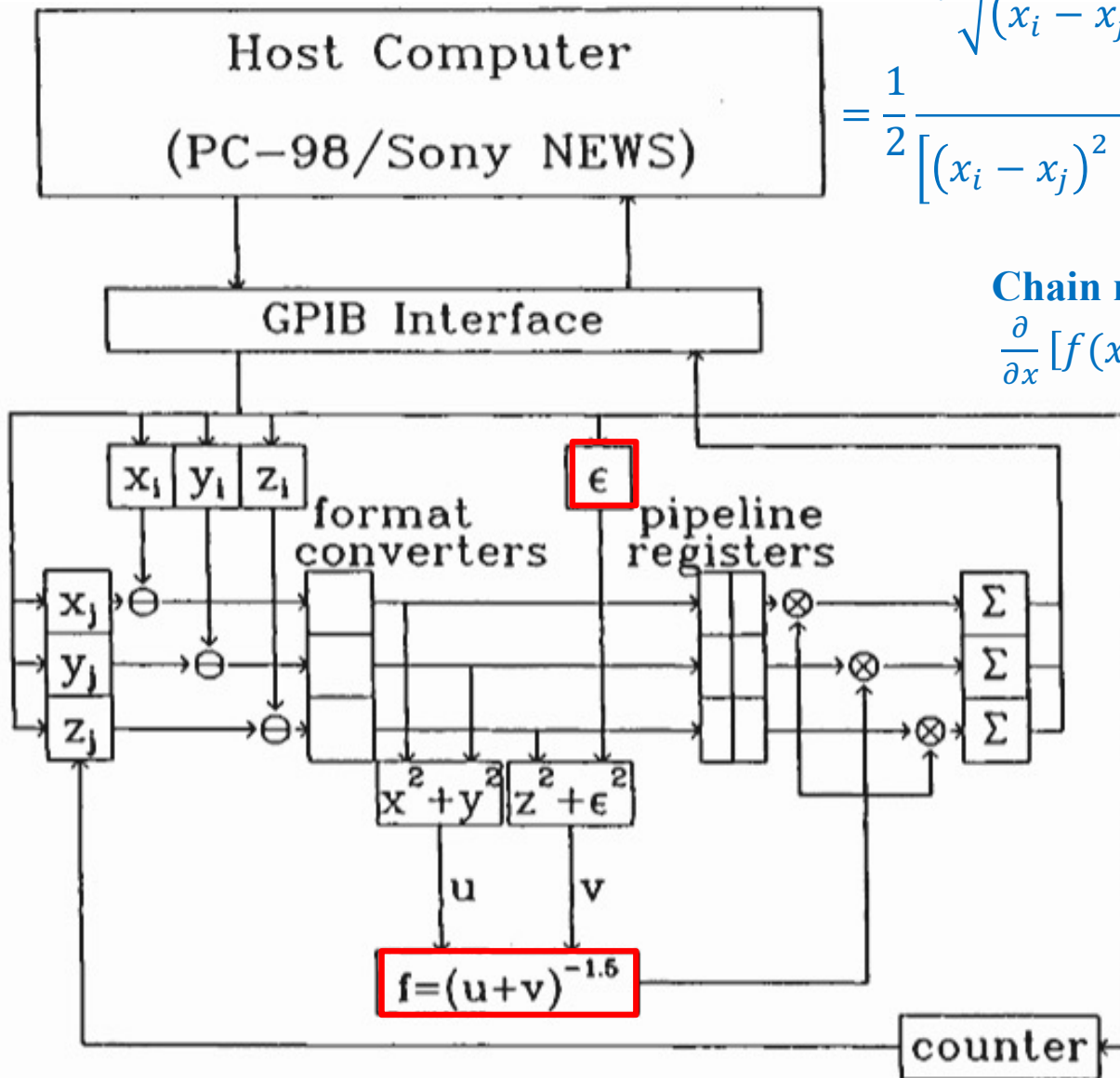
$$\frac{d^2 \mathbf{x}_i}{dt^2} = \mathbf{f}_i = \sum_j \frac{m_j (\mathbf{x}_j - \mathbf{x}_i)}{(r_{ij}^2 + \epsilon^2)^{3/2}}$$

$$-\frac{\partial}{\partial x_i} \frac{1}{\sqrt{(x_i - x_j)^2 + (y_i - y_j)^2 + (z_i - z_j)^2 + \epsilon^2}}$$

$$= \frac{1}{2} \frac{2(x_i - x_j)}{[(x_i - x_j)^2 + (y_i - y_j)^2 + (z_i - z_j)^2 + \epsilon^2]^{3/2}}$$

Chain rule:

$$\frac{\partial}{\partial x} [f(x)]^{-1/2} = -\frac{1}{2} [f(x)]^{-3/2} \frac{\partial f}{\partial x}$$



- Small ϵ to avoid if $(i \neq j)$ & pipeline stall
- $x^{-3/2}$ was implemented as table look-up using read-only memory (ROM)

D. Sugimoto *et al.*,
Nature **345**, 33 ('90)

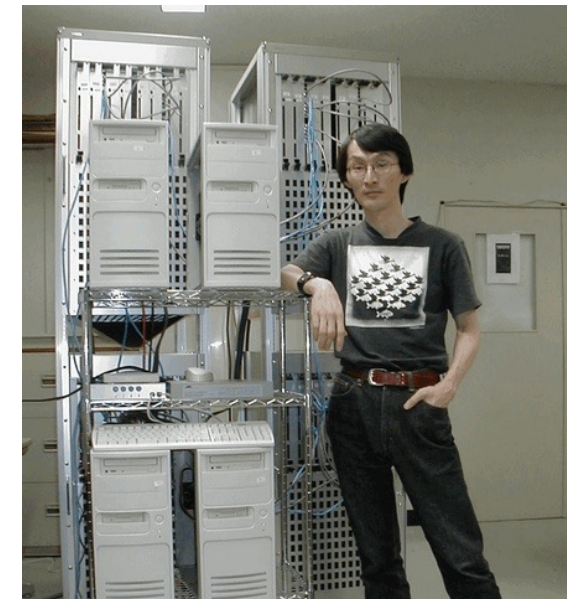
GRAPE & Gordon Bell Prizes

SC2003 Gordon Bell Award
Junichiro Makino
 University of Tokyo
 Performance Evaluation and Tuning of
 GRAPE-6—Towards 40 "Real" Tflops

2003 Gordon Bell Prize, Special Achievement
Performance Evaluation and Tuning of GRAPE-6—Towards 40 'Real' Tflop/s

Junichiro Makino, Hiroshi Daisaka, Eiichiro Kokubo, Toshiyuki Fukushima

*Gordon Bell Prize,
 commonly
 referred to as the
 Nobel Prize of
 Supercomputing
 [Wikipedia].*



**J. Makino
 & Grape 6
 (2001)**

SC2001
 GORDON BELL PRIZE
Junichiro Makino
 Winner, Peak Performance
 A 11.55 Tflops Simulation of Black Holes
 in a Galactic Center on GRAPE-6

2001 Gordon Bell Prize, Winner, Peak Performance
A 11.55 Tflops simulation of black holes in a galactic center on GRAPE-6

Junichiro Makino, Toshiyuki Fukushima

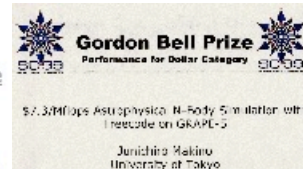
SC2000
 GORDON BELL PRIZE
Junichiro Makino
 Winner, Peak Performance Category
 A 1.34 Tflops Simulation of Black Holes in a Galactic Center on GRAPE-6

2000 Gordon Bell Prize, Winner, Peak Performance Category
A 1.349 Tflops simulation of black holes in a galactic center on GRAPE-6

Junichiro Makino, Toshiyuki Fukushima, Masaki Koga

2000 Gordon Bell Prize, Winner, Peak Performance Category (tie with above)
1.34 Tflops Molecular Dynamic simulation for NaCl with a Special Purpose Computer: MDM
 (MD-GRape system)

Tetsu Narumi, Ryutarō Susukita, Takahiro Koishi, Kenji Yasuoka, Hideaki Furusawa, Atsushi Kawai, Toshikazu Ebisuzaki



1999 Gordon Bell Prize, Price Performance, First Prize
 Astrophysical N-body simulation
 144 Gflops / \$ 1 M on custom-built GRAPE-5 32-processor system

Atsushi Kawai, Toshiyuki Fukushima, and Junichiro Makino



1996 Gordon Bell Prize, Performance, Honorable Mention
 Simulation of the motion of 780,000 stars
 333 Gflops using the Grape-4 machine w/ 1,269 processors

Junichiro Makino, Toshiyuki Fukushima



1995 Gordon Bell Prize, First Place, Special Purpose Machines
 Simulation of the Motion of 10,000 Stars
 112 Gflops using the Grape-4 machine with 288 processors

Astrophysical N-body Simulations on GRAPE-4 Special-Purpose Computer
 Junichiro Makino, Makoto Tajiri

Green500 Rank	MFLOPS/W	Site*	Computer*	Total Power (kW)
1	1684.20	IBM Thomas J. Watson Research Center	NNSA/SC Blue Gene/Q Prototype	38.80
2+	1448.03	National Astronomical Observatory of Japan	GRAPE-DR accelerator Cluster, Infiniband	24.59
2	958.35	GSIC Center, Tokyo Institute of Technology	HP ProLiant SL390s G7 Xeon 6C X5670, Nvidia GPU, Linux/Windows	1243.80
3	933.06	NCSA	Hybrid Cluster Core i3 2.93Ghz Dual Core, NVIDIA C2050, Infiniband	36.00
4	828.67	RIKEN Advanced Institute for Computational Science	K computer, SPARC64 VIIIfx 2.0GHz, Tofu Interconnect	57.96

www.green500.org (Nov. '10)

Enabling Science by Hardware

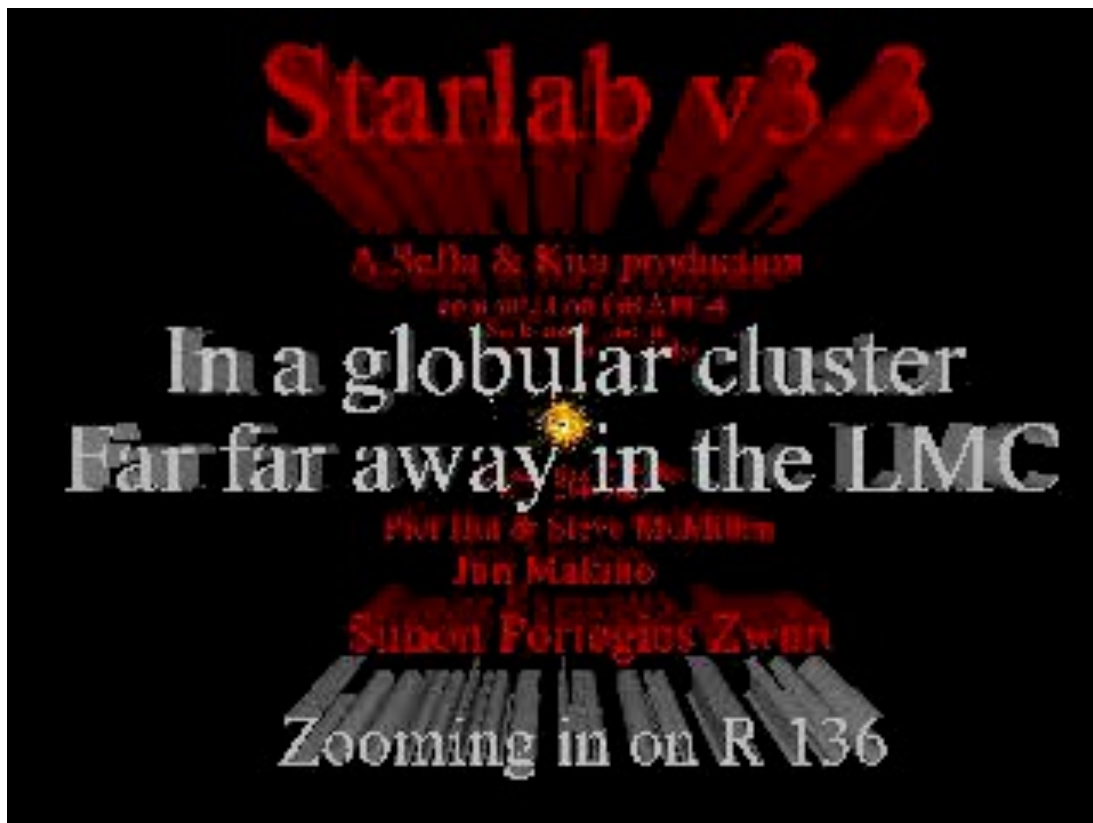
A special-purpose computer for gravitational many-body problems

Daiichiro Sugimoto*, Yoshihiro Chikada†, Junichiro Makino*, Tomoyoshi Ito*,
Toshikazu Ebisuzaki* & Masayuki Umemura†

NATURE · VOL 345 · 3 MAY 1990

33

© 1990 Nature Publishing Group



In Univ. of Tokyo, computer science started as part of physics department (not math or CS)

Computer Physics Communications 60 (1990) 187–194

A special-purpose N -body machine GRAPE-1

Tomoyoshi Ito, Junichiro Makino, Toshikazu Ebisuzaki and Daiichiro Sugimoto
Department of Earth Science and Astronomy, College of Arts and Sciences, University of Tokyo, Tokyo 153, Japan

[CPC homepage](#)

Submit your first paper to *CPC*!

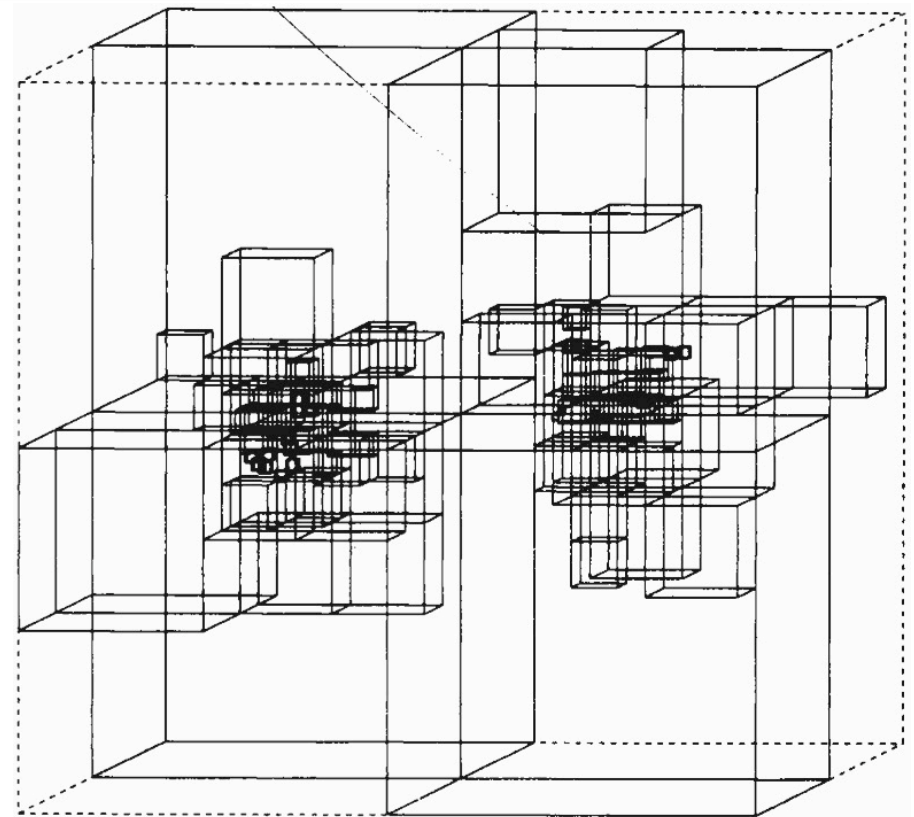
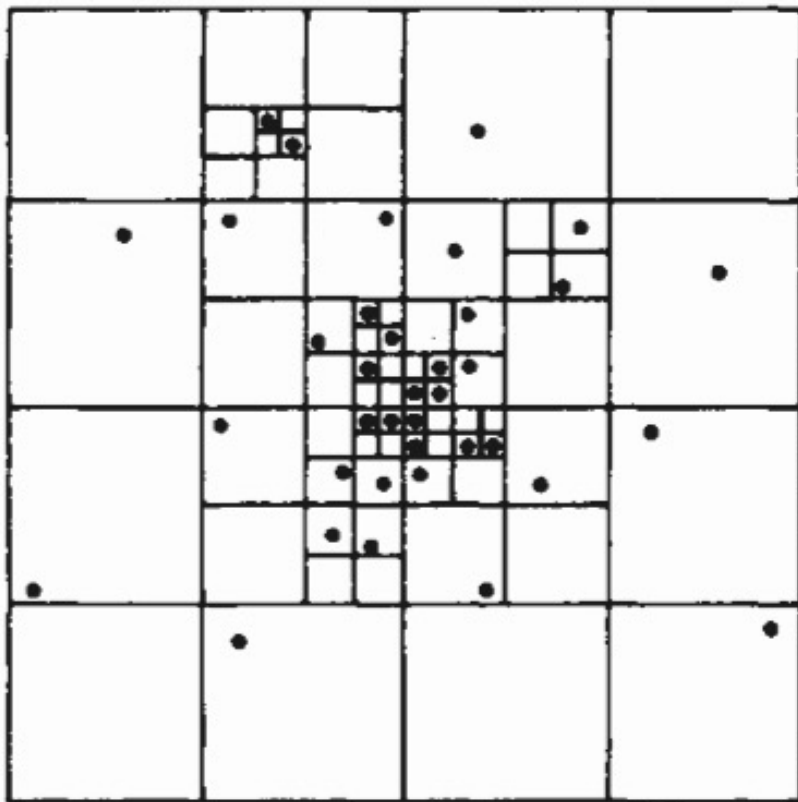
Enabling Science by Algorithm

NATURE

NATURE VOL. 324 4 DECEMBER 1986

A hierarchical $O(N \log N)$ force-calculation algorithm

Josh Barnes & Piet Hut



ACM Best Theses: Machine vs. Algorithm

DANNY HILLIS

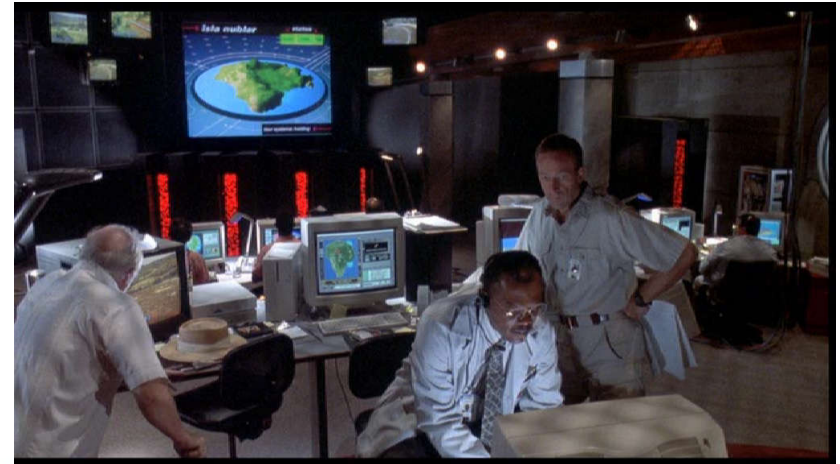
Doctoral Dissertation Award
United States – 1985

CITATION

For his dissertation "*The Connection Machine.*"

Watch: Hillis on Richard Feynman

<http://longnow.org/essays/richard-feynman-connection-machine/>

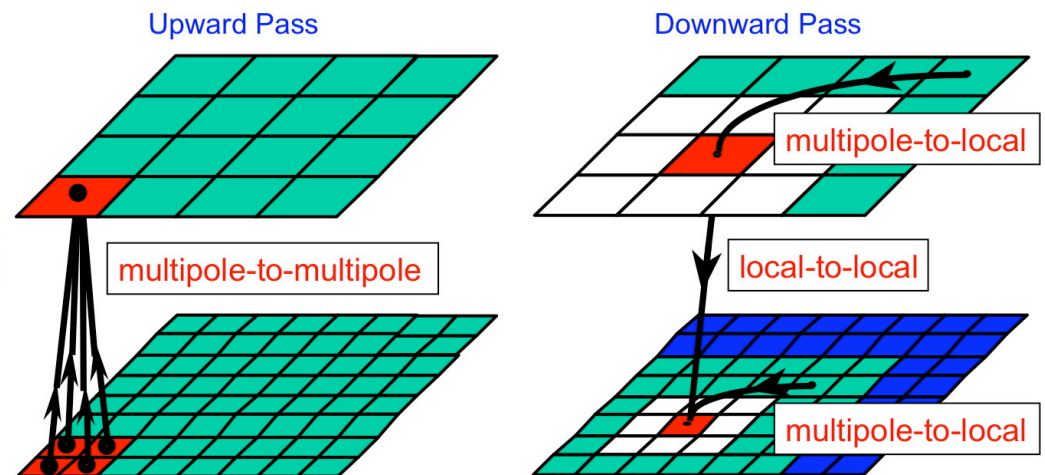


LESLIE GREENGARD

Doctoral Dissertation Award
United States – 1987

CITATION

For his dissertation "*The Rapid Evaluation of Potential Fields in Particle Systems.*"



See lecture notes at <https://aiichironakano.github.io/cs653.html>

A Small Step in Parallel MD

VOLUME 71, NUMBER 1

PHYSICAL REVIEW LETTERS

5 JULY 1993

Structural Correlations in Porous Silica: Molecular Dynamics Simulation on a Parallel Computer

Aiichiro Nakano, Lingsong Bi, Rajiv K. Kalia, and Priya Vashishta

*Concurrent Computing Laboratory for Materials Simulations, Department of Physics and Astronomy,
Department of Computer Science, Louisiana State University, Baton Rouge, Louisiana 70803-4001*

(Received 9 November 1992; revised manuscript received 2 April 1993)

Molecular dynamics simulations of porous silica in the density range $2.2\text{--}0.1\text{ g/cm}^3$ are carried out on a 41 472 particle system using a multiple instruction multiple data computer. The internal surface area, pore surface-to-volume ratio, pore size distribution fractal dimension, correlation length, and mean particle size are determined as a function of the density. Structural transition between a condensed amorphous phase and a low-density porous phase is characterized by these quantities. Various dissimilar porous structures with different fractal dimensions are obtained by controlling the preparation schedule and temperature.

This work was supported by the U.S. Department of Energy, Office of Energy Research, Basic Energy Science, Materials Science Division, Grant No. DE-FG05-92ER45477. The computations were performed using the eight-node iPSC/860 in the Concurrent Computing Laboratory for Materials Simulations (CCLMS) at Louisiana State University. The facilities in the CCLMS were acquired with the Equipment Enhancement Grants awarded by the Louisiana Board of Regents through Louisiana



More N -body Simulations at SC

42 TFlops Hierarchical N -body Simulations on GPUs with Applications in both Astrophysics and Turbulence

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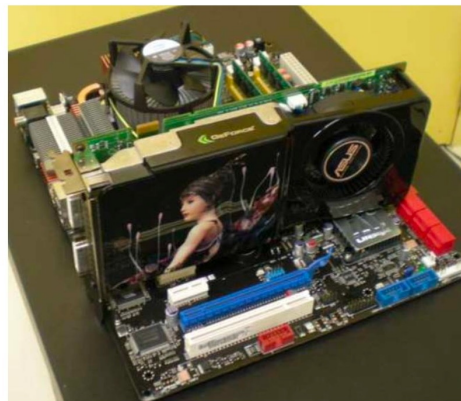
Keigo Nitadori
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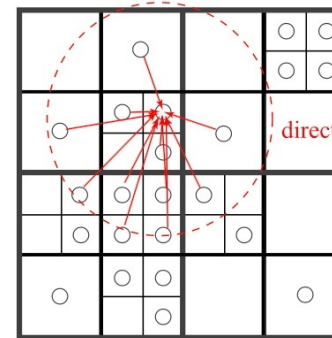
2009 Gordon Bell Prize Price/Performance Category

Table 2: Price of the GPU cluster

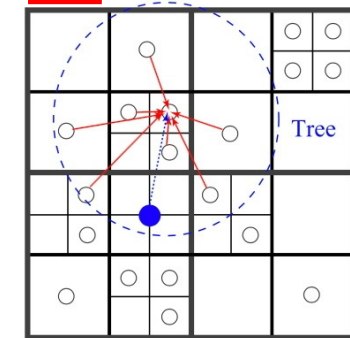
Elements	Quantity	Price (JPY)	Price (\$)
GPUs	256	12,160,000	\$ 118,345
Host PCs	128	10,716,032	\$ 104,292
Network switch	4	644,800	\$ 6,275
Total		23,520,832	\$ 228,912



P³M



TreePM



4.45 Pflops Astrophysical N -Body Simulation on K computer - The Gravitational Trillion-Body Problem

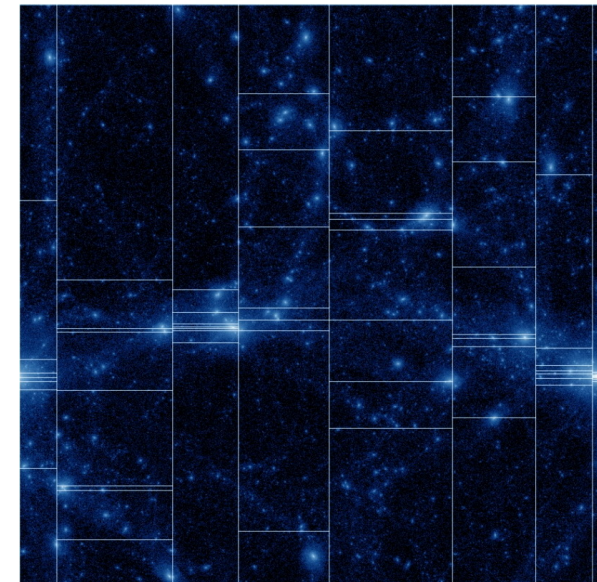
Tomoaki Ishiyama
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Tokyo Institute of Technology
makino@geo.titech.ac.jp

IEEE/ACM supercomputing, SC12

*Machine
&
algorithm!*

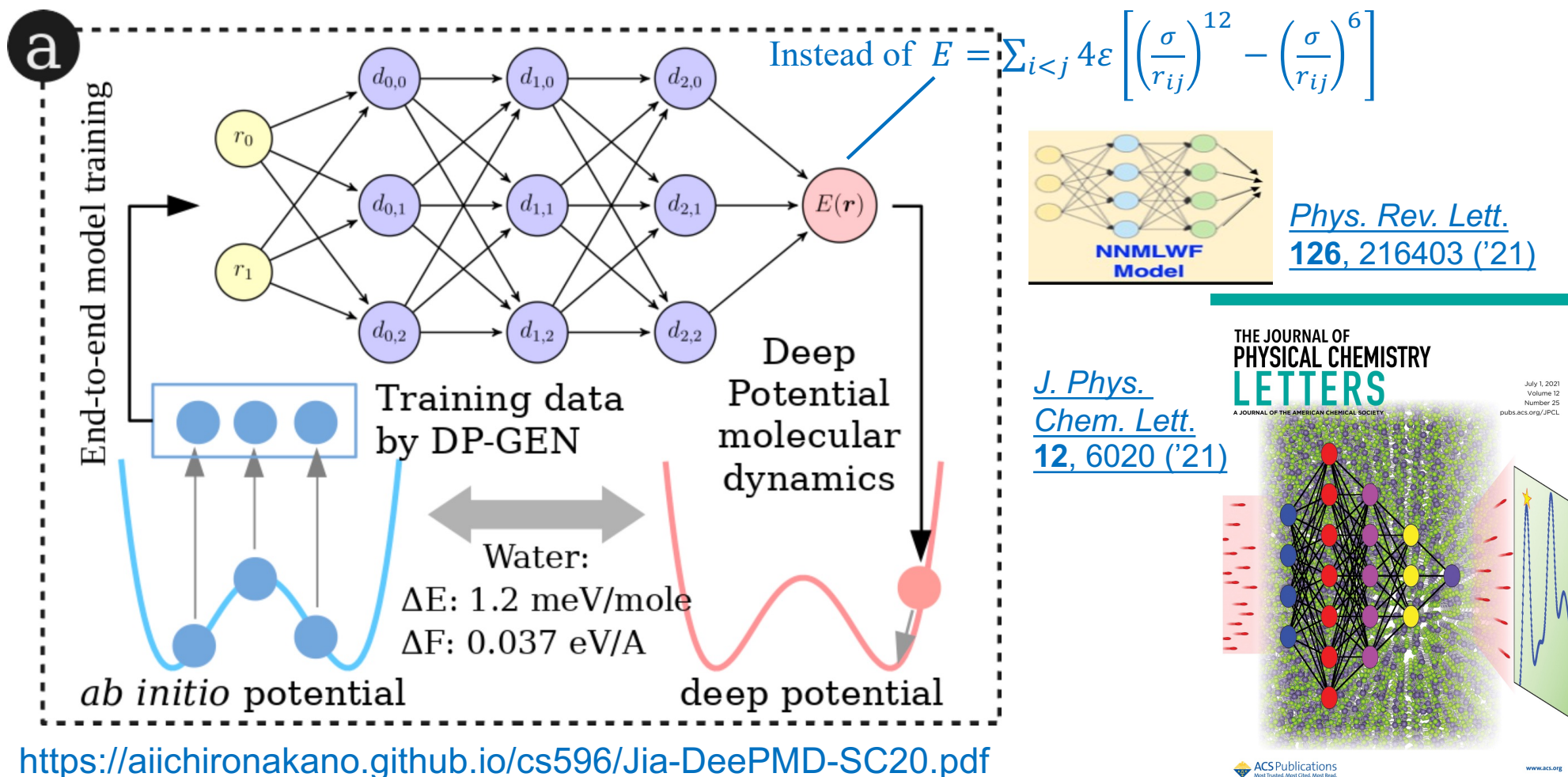


Now It's Deep MD

Pushing the limit of molecular dynamics with ab initio accuracy to 100 million atoms with machine learning

Weile Jia, Han Wang, Mohan Chen, Denghui Lu, Jiduan Liu, Lin Lin, Roberto Car, Weinan E, Linfeng Zhang

Proc. Supercomputing, SC20, Gordon Bell prize



<https://aiichironakano.github.io/cs596/Jia-DeePMD-SC20.pdf>

Enabling Science by Online Game

nature

Vol 466 | 5 August 2010 | doi:10.1038/nature09304

LETTERS

Predicting protein structures with a multiplayer online game

Seth Cooper¹, Firas Khatib², Adrien Treuille^{1,3}, Janos Barbero¹, Jeehyung Lee³, Michael Beenen¹, Andrew Leaver-Fay^{2,†}, David Baker^{2,4}, Zoran Popović¹ & Foldit players

The image shows a screenshot of the Foldit game interface. On the left, there is a green banner with the Foldit logo and the text "Solve Puzzles for Science". Below the banner, a protein structure is shown with a text box that says "Click to learn how you contribute to science by playing Foldit." On the right, the main game area displays a complex protein puzzle with various colored segments and arrows pointing to specific parts. A scoreboard at the top right shows the player's rank (317) and score (2534). A group competition table is also visible, listing group names and scores. At the bottom, there is a toolbar with various actions like "Shake Sidechains", "Wiggle All", "Wiggle Backbone", "Wiggle Sidechains", "Freeze Protein", "Remove Bands", "Disable Bands", "Align Guide", "Reset Structures", "Reset Puzzle", "Help", and "Glossary".

foldit BETA
Solve Puzzles for Science
BLOG PUZZLES FEEDBACK

14:35:22 GMT

Rank: 317 Score: 2534
Soloist Beginner Puzzle 8 (<150): Fruit Fly

Group Competition

#	Group Name	Score
1	Bio Biochemistry	9174
2	Team Commonwealth	9168
3	Ukraine	9088
4	Team Canada	9085
5	Firebird BioChem	9073
6	SETI Germany	9030
7	Bois de	9001

Soloist Competition

#	Player Name	Current	Best
1	Mie Crunching for Physics	9242	9242
2	weitzel	9235	9235
3	ys719	9222	9222
4	markic	9211	9211
5	kaavo_karplus	9186	9186
6	JIMKOR	9183	9183
7	sb.eric	9183	9183

Click to learn how you contribute to science by playing Foldit.

Shake Sidechains Wiggle All Wiggle Backbone Wiggle Sidechains Freeze Protein Remove Bands Disable Bands Align Guide Reset Structures Reset Puzzle Help Glossary

Chat - Group auto show
Chat - Puzzle auto show
Chat - Global auto show
Notifications auto show

Ising Machine



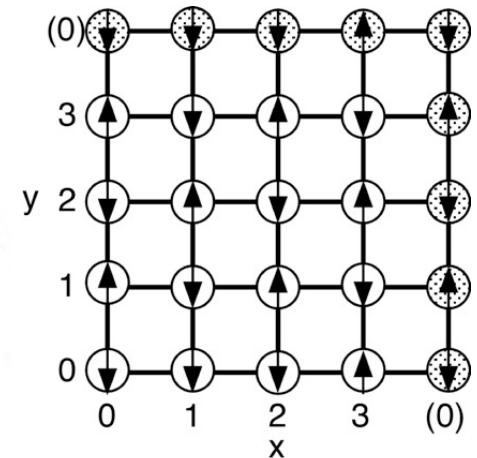
1bit の世界の専用計算機 ——イジング・マシン——

泰地 真弘人
(東京大学教養学部)
(1994年3月2日受理)

Ising Machine:
A Special Purpose Computer for 1-bit Worlds

TAIJI Makoto
(Received 3 March 1994)

$$V(s^N) = -J \sum_{(k,l)} s_k s_l - H \sum_k s_k, \quad s_k = \pm 1$$



<https://aiichironakano.github.io/phys516-lecture.html>

Abstract

This paper describes the development of special-purpose computer systems for Ising models, "Ising Machine" m-TIS 1 and 2. The first two sections explain Ising models and their Monte Carlo simulations. In section 3 and 4, I describe my motivation to build a special-purpose computer and the development of m-TIS 1. In section 5 and 6, the use of field-programmable gate arrays in a special-purpose computer is discussed. In the last two sections I discuss the potential abilities and future prospects of both Ising machine and a special-purpose computer in general. *J. Plasma Fusion Res.* **70**, 332 ('94)

cf. Original GRAPE was a 48-bit machine

USC Quantum Computation Center

- **D-Wave 2X system with 1,098-quantum bits (qubits)**

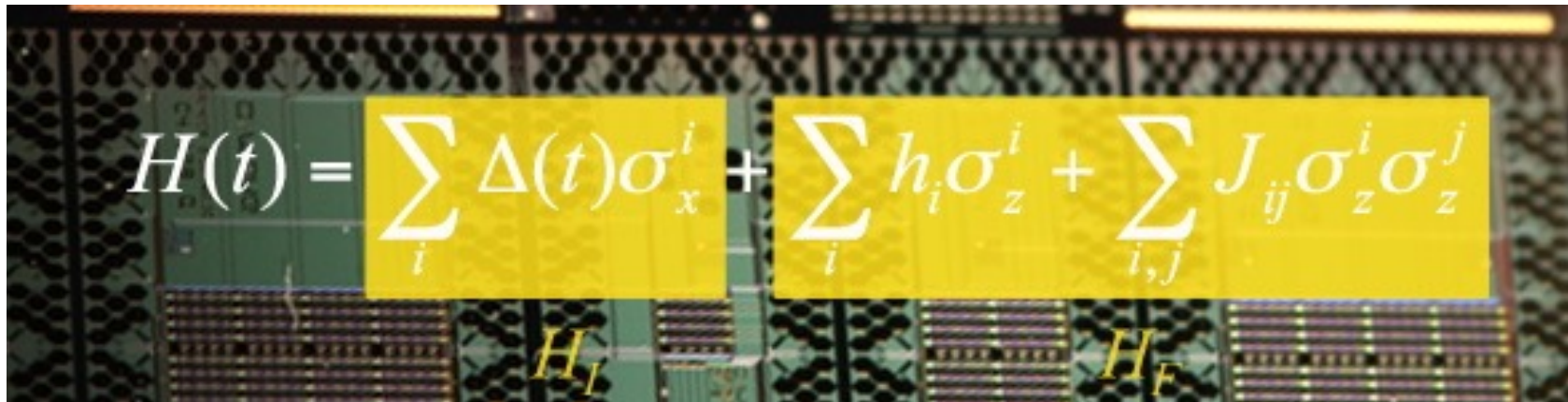
Phase transitions in a programmable quantum spin glass simulator

R. Harris^{1*}, Y. Sato¹, A. J. Berkley¹, M. Reis¹, F. Altomare¹, M. H. Amin^{1,2}, K. Boothby¹, P. Bunyk¹, C. Deng¹, C. Enderud¹, S. Huang¹, E. Hoskinson¹, M. W. Johnson¹, E. Ladizinsky¹, N. Ladizinsky¹, T. Lanting¹, R. Li¹, T. Medina¹, R. Molavi^{1,3}, R. Neufeld¹, T. Oh¹, I. Pavlov¹, I. Perminov¹, G. Poulin-Lamarre¹, C. Rich¹, A. Smirnov¹, L. Swenson¹, N. Tsai¹, M. Volkmann¹, J. Whittaker¹, J. Yao¹

Harris *et al.*, *Science* **361**, 162–165 (2018) 13 July 2018

- **Adiabatic quantum optimization**



A photograph of a quantum circuit board with a complex grid of components. Overlaid on the image is a mathematical equation for a Hamiltonian H(t). The equation is:
$$H(t) = \sum_i \Delta(t) \sigma_x^i + \sum_i h_i \sigma_z^i + \sum_{i,j} J_{ij} \sigma_z^i \sigma_z^j$$
 The terms of the equation are highlighted in yellow boxes. Below the equation, the labels H_I and H_F are visible on the circuit board.

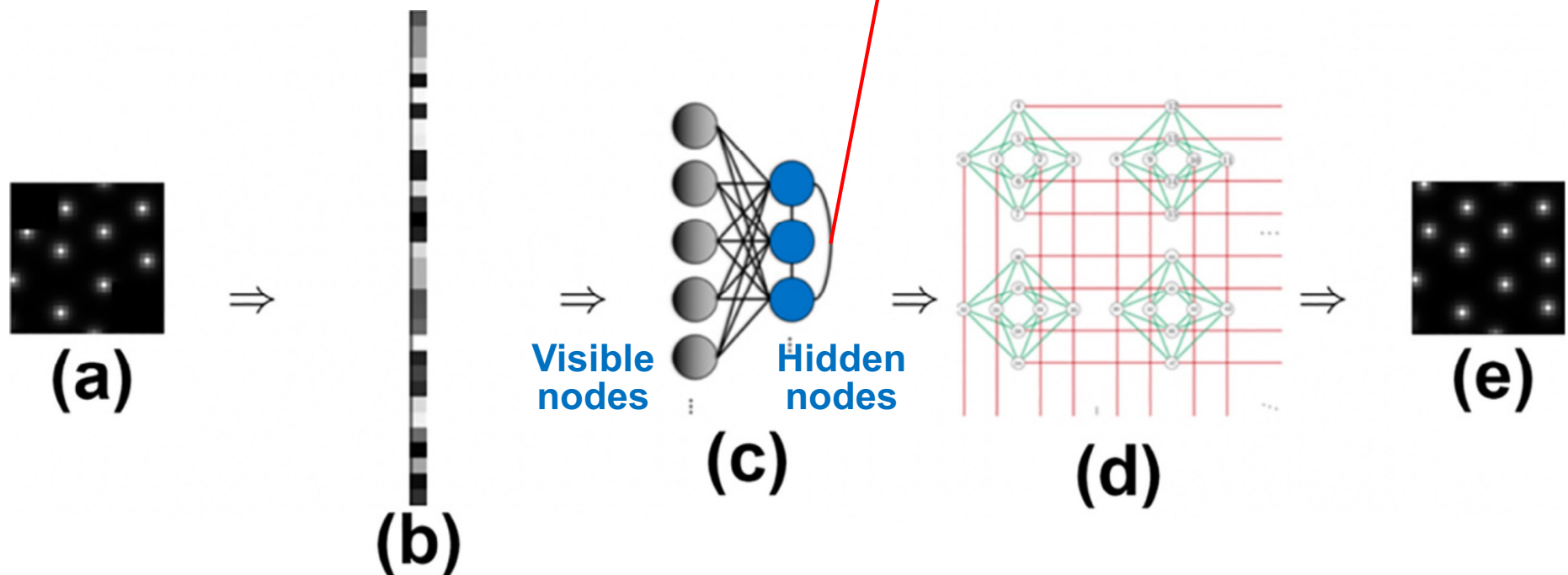
Machine Learning on D-Wave

Boltzmann machine modeling of layered MoS₂ synthesis on a quantum annealer

J. Liu, A. Mohan, R. K. Kalia, A. Nakano, K. Nomura, P. Vashishta, and K.T. Yao

[Comput. Mater. Sci. 173, 109429 \('20\)](#)

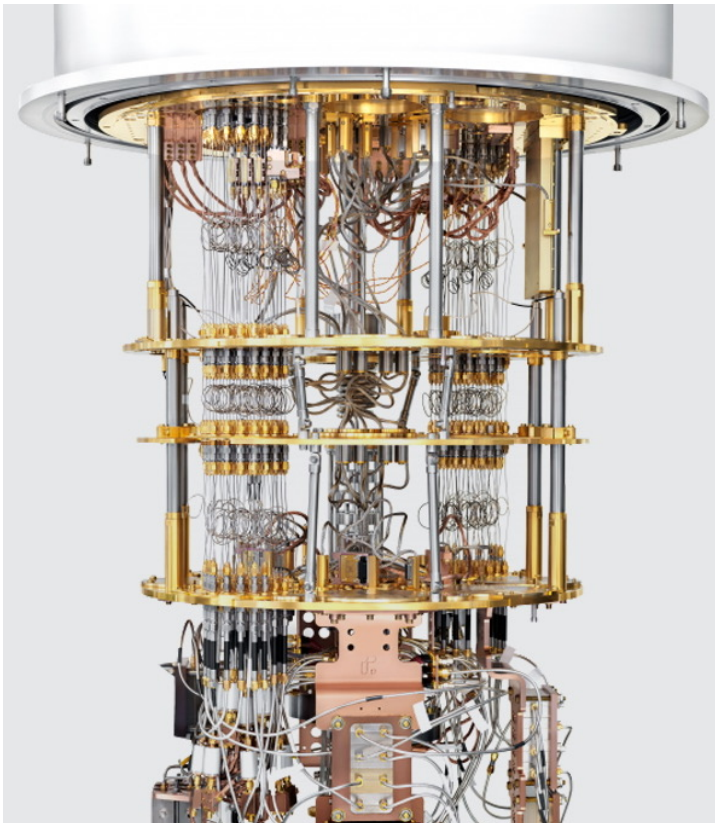
- Computing power of D-Wave allows **unrestricted Boltzmann Machine** to enhance machine learning performance



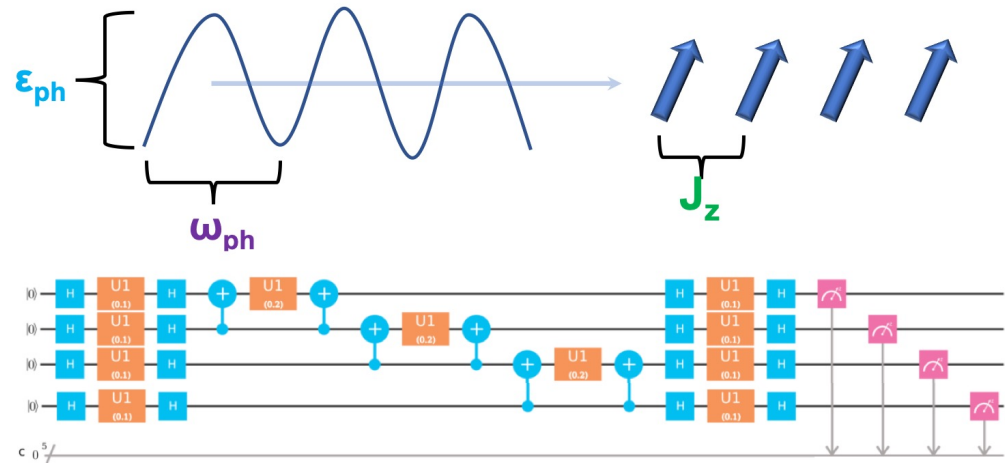
Final project by Ankith Mohan (MSCS) with Jeremy Liu (PhD-CS)

Quantum Computational Science

- **Quantum computing for science: *Universal simulator of quantum many-body systems*** [R. P. Feynman, *Int. J. Theo. Phys.* **21**, 467 ('82); S. Lloyd, *Science* **273**, 1073 ('96)]
- **Successfully simulated quantum many-body dynamics on publicly-available IBM's Q16 Melbourne & Rigetti's Aspen quantum computers** [L. Bassman *et al.*, *Phys. Rev. B* **101**, 184305 ('20)]
- **AI-inspired domain-specific quantum compiler has reduced the circuit size by 30% below that by the vendor's native compiler** [L. Bassman *et al.*, *Quant. Sci. Tech.* **6**, 014007 ('21)]



$$H(t) = -J_z \sum_{i=1}^{N-1} \sigma_z^i \sigma_z^{i+1} - \epsilon_{ph} \sin(\omega_{ph} t) \sum_{i=1}^N \sigma_x^i$$



```

32 | ...#define the two non-commuting terms that comprise the Hamiltonian-
33 | ...Hz := PauliTerm("Z", 0, epsilon_0)-
34 | ...Hy := PauliTerm("Y", 0, epsilon_ph*np.sin(w_ph*t))-
35 | ...#exponentiate the terms of the Hamiltonian for use in Trotter approx-
36 | ...exp_Hz := exponential_map(Hz)(delta_t/(2.0*hbar))-
37 | ...exp_Hy := exponential_map(Hy)(delta_t/hbar)-

```