

# How to Teach your Exascale Machine to Do the Data Dance

#### **Kathy Yelick**

Professor of Electrical Engineering and Computer Sciences University of California at Berkeley Associate Laboratory Director for Computing Sciences Lawrence Berkeley National Laboratory

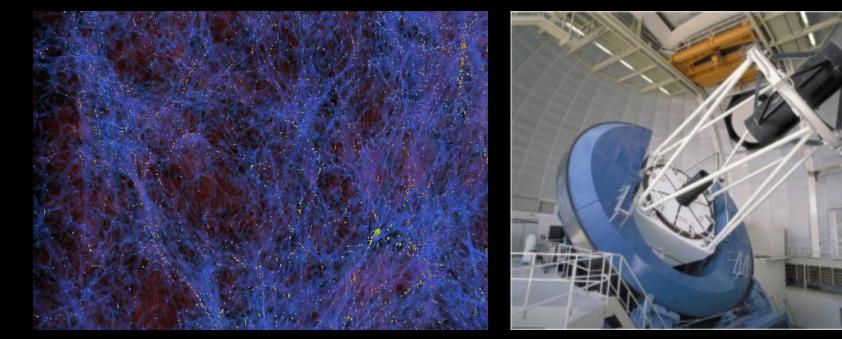
#### "Big Data" Changes Everything...What about Science?







#### **Combine simulation and observation for next Cosmology breakthrough**

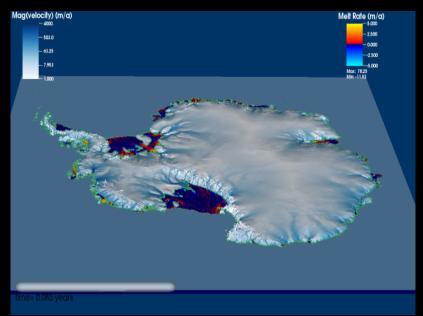


Nyx simulation of Lyman alpha forest using AMR

*Kitt Peak National Observatory's Mayall 4-meter telescope, planned site of the DESI experiment* 

Reduce systematic bias in observation through simulation of ~1 Gigaparsec Baryon Acoustic Oscillations in the Lyman Alpha Forest and ~100 Gigaparsec simulation of galaxy clusters, both requiring adaptive mesh refinement (AMR).

# Climate models and microbial analysis together to predict the future of the environment



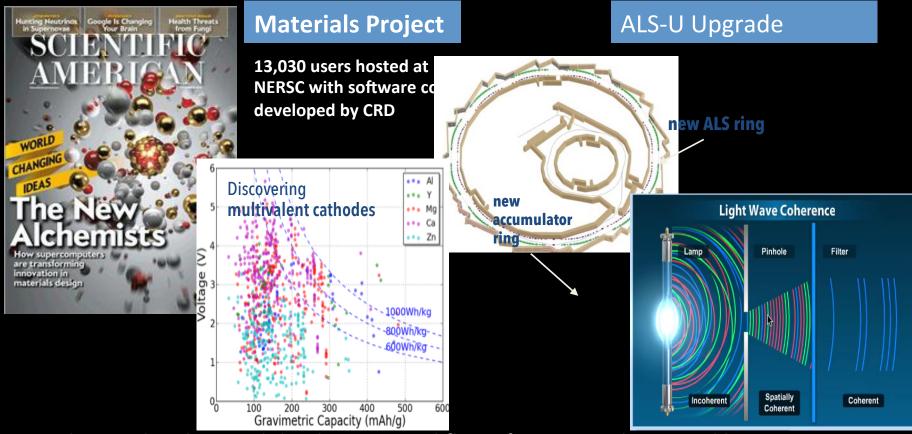
New climate modeling methods, including AMR "Dycore" produce new understanding of ice



Genomes to watersheds Scientific Focus Area

Understand interactions between environmental microbiomes and climate change with *kilometer resolution models* that track dynamic 3D features (with AMR) and *genome-enabled analysis* of environmental sensors.

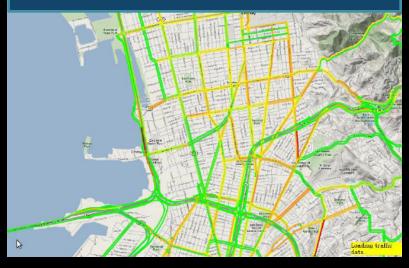
# Understand and control energy with advanced light sources and materials modeling



Understand and control the direction and flow of energy with minimal losses using *advanced instruments, high fidelity models,* and high throughput simulation and analysis for applications in energy, environment and computing,

### Science in embedded sensors: Internet of Things

#### **Transportation Modeling**



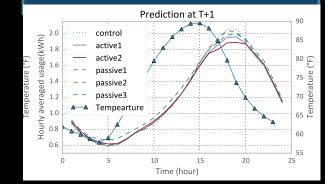
#### Power Grid Modeling



#### Scenario Prediction, Planning



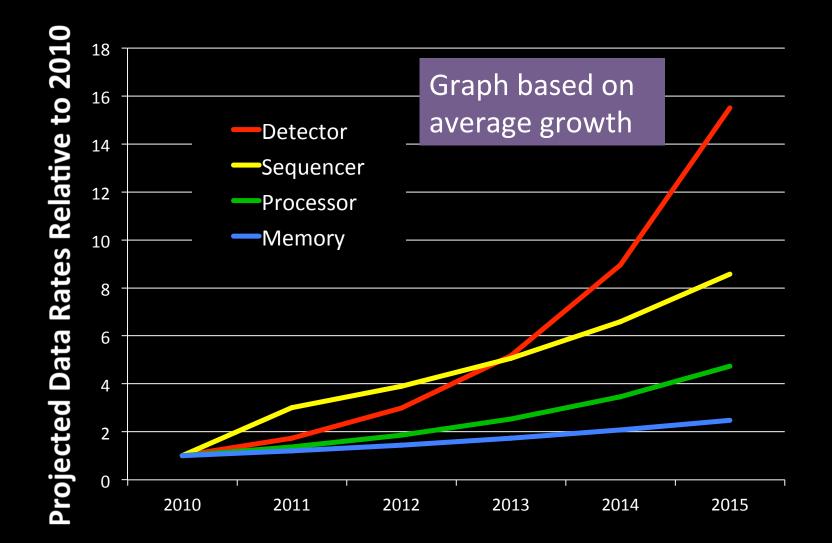
#### **Decision Science**



### Roadmap for this talk

- Science at the boundary of simulation and observation
- Science data challenges
- What do I mean by Exascale?
- Exascale challenges for Data problems
  - Programming models
  - Algorithms
  - Architectures
  - Systems
  - Policies

#### Science Data Growth is Outpacing Computing



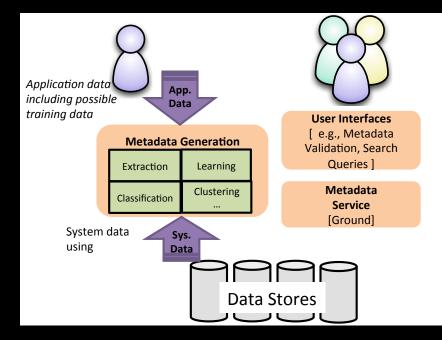
#### **Old School Scientific Data Search**

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### Automated Search, Meta-Data Analysis, and On-Demand Simulation



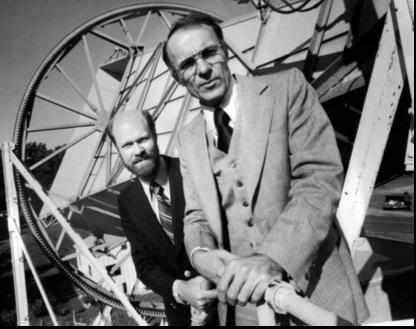
Jobs submitted by "bots" based on queries; algorithms extract informatics for design Automated metadata extraction using machine learning



#### Filtering, De-Noise and Curating Data



AmeriFlux & FLUXNET: 750 users access carbon sensor data from 960 carbon flux data years



Arno Penzias and Robert Wilson discover Cosmic Microwave Background in 1965

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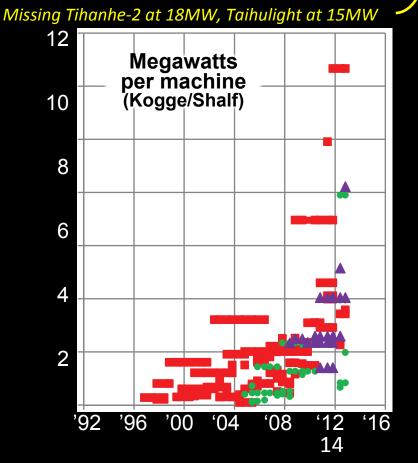
Exascale = Orders of magnitude increase in performance at all scales

Continued growth at constant energy
 Continued growth at constant cost

#### **Computing is energy-constrained**

At ~\$1M per MW, energy costs are substantial

- 1 petaflop in 2008 used 3 MW
- 1 exaflop in 2018 at 200 MW "usual chip scaling"



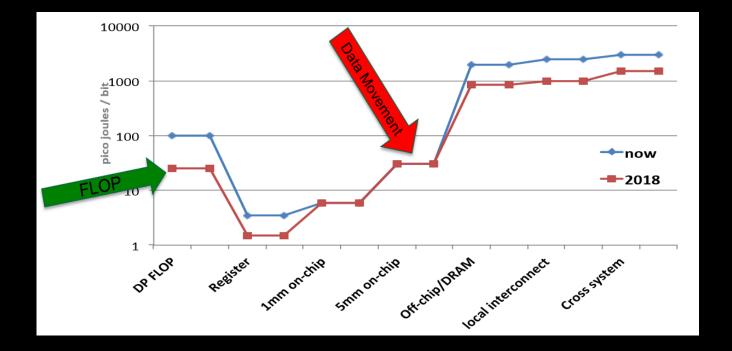
Goal: 1 Exaflop in 20 MW = 20 pJ / operation

#### Note: The 20 pJ / operation is

- Independent of machine size
- Independent of # cores used per application
- But "operations" need to be useful ones

## **Challenge: Communication is expensive**

#### Communication is expensive in time and energy



#### Hard to change: Latency is physics; bandwidth is money!

### Roadmap for this talk

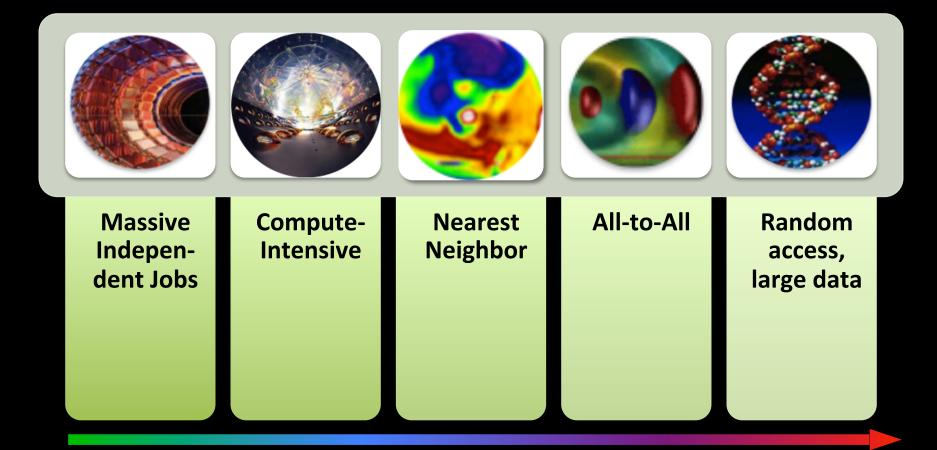
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#### Data vs. Simulation: The Irregularity Spectrum



#### **Different architectures? Programming models?**

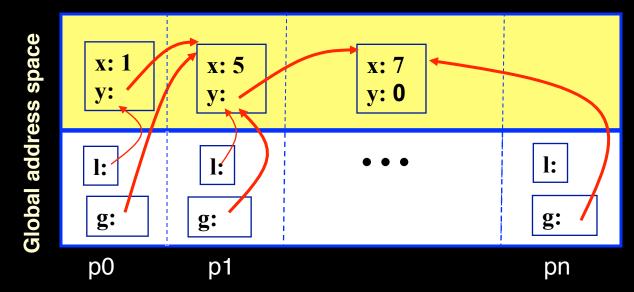
#### **PGAS: A programming model for exascale**

• Global address space: thread may directly read/write remote data using an address (pointers and arrays)

... = \*gp; ga[i] = ...

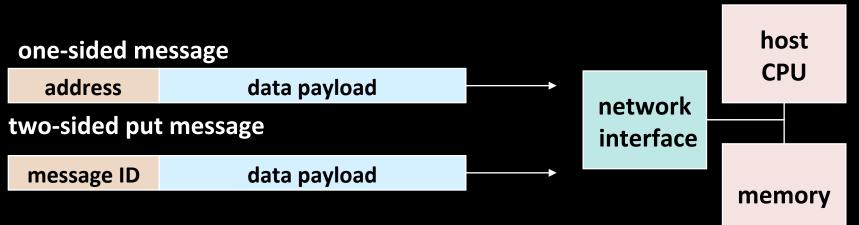
• Partitioned: data is designated as local or global

shared int [ ] ga; and upc\_malloc (...)

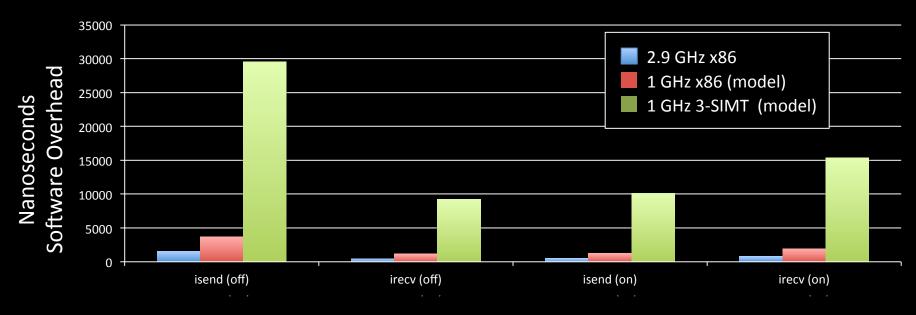


A programming model can influence how programmers think

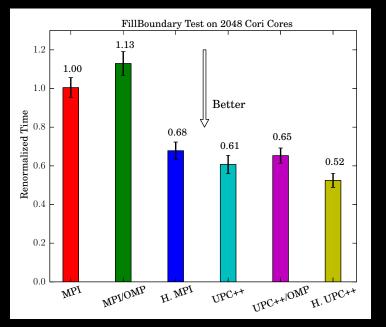
### **One-Sided Communication is Closer to Hardware**

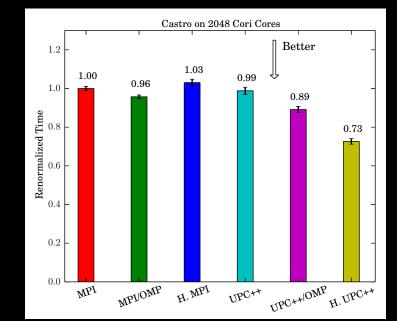


- Hardware does 1-sided communication
- Overhead for send/receive messaging is worse at exascale



### **One-sided PGAS (UPC++) in AMR**



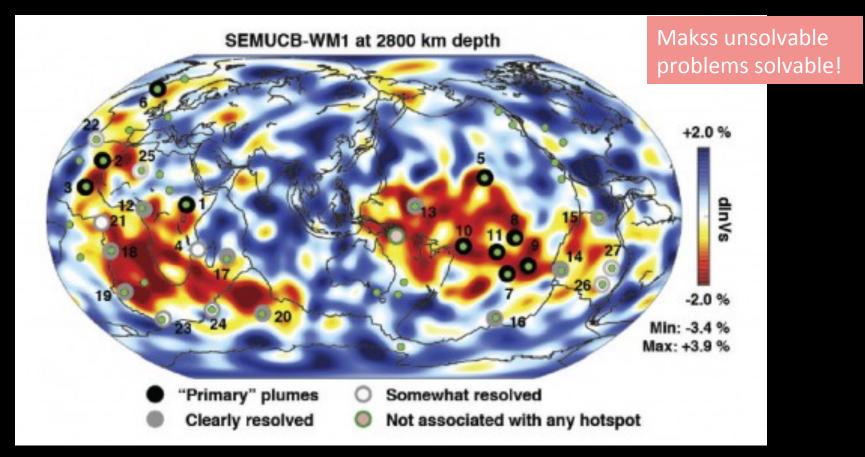


#### Adaptive Mesh Refinement (AMR) using UPC++

- Metadata costs make flat MPI impractical
- Replaced communication (retained most code)
- Hierarchical algorithms (UPC++/UPC++ or MPI/MPI best)

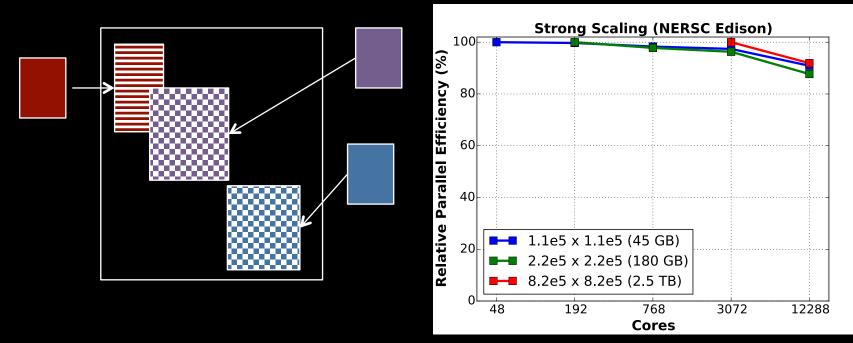
### **Science Impact: Whole-Mantle Seismic Model**

- First-ever whole-mantle seismic model from numerical waveform tomography
- *Finding: Most* volcanic hotspots are linked to two spots on the boundary between the metal core and rocky mantle 1,800 miles below Earth's surface.



Scott French, Barbara Romanowicz, "Broad plumes rooted at the base of the Earth's mantle beneath major hotspots", **Nature, 2015** 

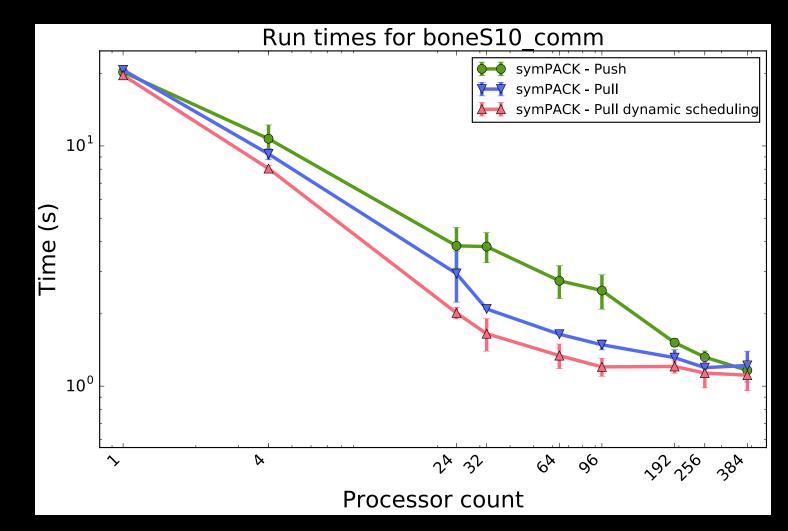
### **Data Fusion for Observation with Simulation**



- Unaligned data from observation
- One-sided strided updates
- Could MPI-3.0 one-sided do this? Yes, but not well so far

Scott French, Y. Zheng, B. Romanowicz, K. Yelick

### **Sparse Cholesky in PGAS (UPC++)**



Fan-both algorithm by Jacquelin & Ng, in UPC++

#### *Unstructured, Graph-based, Data analytics problem: De novo* Genome Assembly

- DNA sequence consists of 4 bases: A/C/G/T
- Read: short fragment of DNA sequence that can be read by a DNA sequencing technology – can't read whole DNA at once.
- De novo genome assembly: Reconstruct an unknown genome from a collection of short reads.
  - Constructing a jigsaw puzzle without having the picture on the box





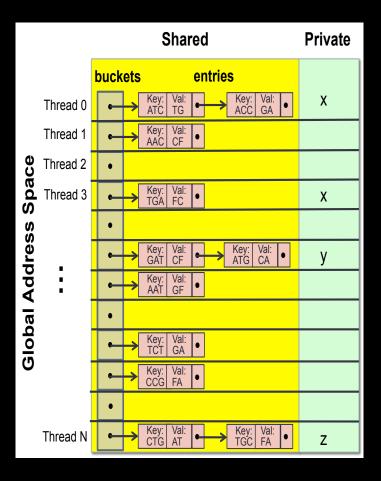
#### **Random Access Graph Analytics**

- Genome assembly "needs shared memory"
  - **Global Address Space** Low overhead communication
- Remote atomics

 $\bullet$ 

• Partitions for any structure

### Scales to 15K+ cores Under 10 minutes for human First ever solution

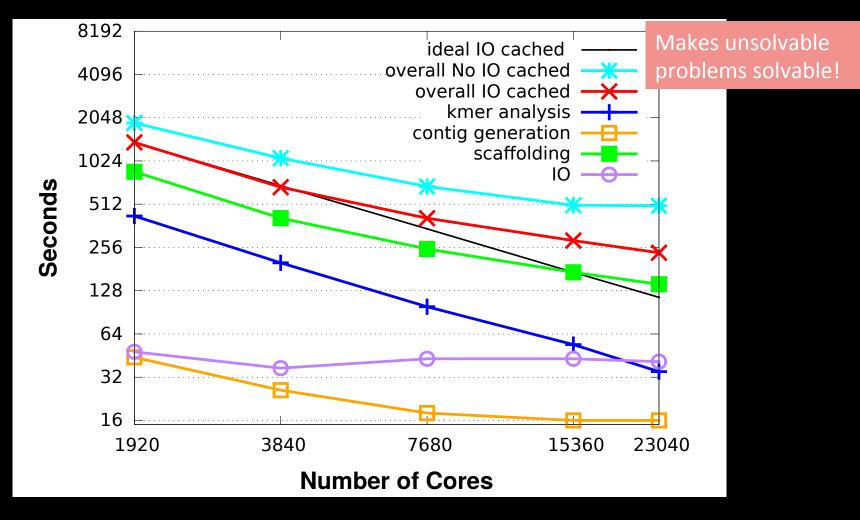


E. Georganas, A. Buluc, J. Chapman, S. Hofmeyr, C. Aluru, R. Egan, L. Oliker, D. Rokhsar, K. Yelick

### Many types of distributed hash tables in HipMer

- Global update only
  - Can aggregate and reorder updates
- Global read only
  - Sometimes with good hash locality so caching helps
- Global read-modify-write of elements in table
  - Remote atomics
- Local read and write
  - Separate all-to-all or reduction phase

# Strong scaling (human genome) on Cray XC30



- Complete assembly of human genome in **4 minutes using 23K cores.**
- **700x speedup over** original Meraculous (took **2,880 minutes** on large shared memory with some Perl code); Some problems (wheat, squid, only run on HipMer version)

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  - Algorithms: Communication avoidance
  - Architectures
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### **Beyond Domain Decomposition**

2.5D Matrix Multiply on BG/P, 16K nodes / 64K cores

#### Surprises:

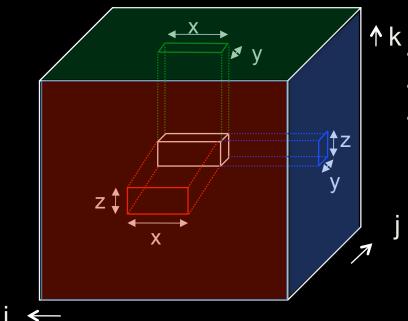
- Even Matrix Multiply had room for improvement
- Idea: make copies of C matrix (as in prior 3D algorithm, but not as many)
- Result is provably optimal in communication

Lesson: Never waste fast memory And don't get hung up on the owner computes rule

Can we generalize for compiler writers?

#### Deconstructing 2.5D Matrix Multiply

Solomonick & Demmel



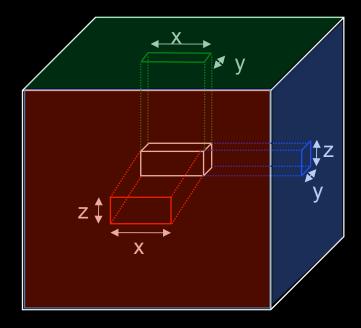
- Tiling the iteration space
- 2D algorithm: never chop k dim
- 2.5 or 3D: Assume + is associative; chop k, which is → replication of C matrix

Matrix Multiplication code has a 3D iteration space Each point in the space is a constant computation (\*/+)

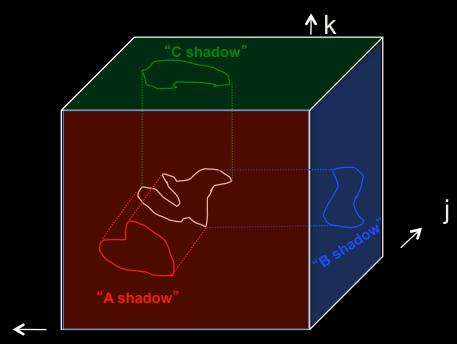
```
for i
for j
for k
C[i,j] ... A[i,k] ... B[k,j] ...
```

#### Lower Bound Idea on C = A\*B

Iromy, Toledo, Tiskin



- Cubes in black box with side lengths x, y and z = Volume of black box
- = x\*y\*z
- = (#A□s \* #B□s \* #C□s )<sup>1/2</sup>
- = ( xz \* zy \* yx)<sup>1/2</sup>



(i,k) is in **"A shadow"** if (i,j,k) in 3D set (j,k) is in **"B shadow"** if (i,j,k) in 3D set (i,j) is in **"C shadow"** if (i,j,k) in 3D set

Thm (Loomis & Whitney, 1949) # cubes in 3D set = Volume of 3D set ≤ (area(A shadow) \* area(B shadow) \* area(C shadow))<sup>1/2</sup>

#### Generalizing Communication Lower Bounds and Optimal Algorithms

- For serial matmul, we know #words\_moved = Ω (n<sup>3</sup>/M<sup>1/2</sup>), attained by tile sizes M<sup>1/2</sup> x M<sup>1/2</sup>
- Thm (Christ, Demmel, Knight, Scanlon, Yelick): For any program that "smells like" nested loops, accessing arrays with subscripts that are linear functions of the loop indices

 $#words\_moved = \Omega$  (#iterations/M<sup>e</sup>)

for some e we can determine

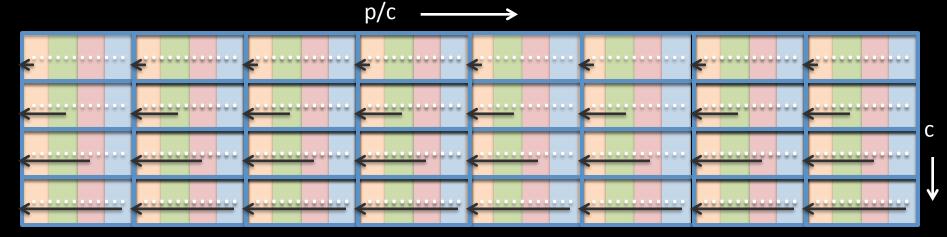
- Thm (C/D/K/S/Y): Under some assumptions, we can determine the optimal tiles sizes
  - E.g., index expressions are just subsets of indices
- Long term goal: All compilers should generate communication optimal code from nested loops

#### **Implications for Compilers**

↑	<ul> <li>Much of the work on compilers is based on owner-computes</li> </ul>					
x +=	<ul> <li>For MM: Divide C into chunks, schedule movement of A/B</li> </ul>					
x +=	<ul> <li>Data-driven domain decomposition partitions data; but we can partition work instead</li> </ul>					
	<ul> <li>Ways to compute C "pencil"</li> </ul>					
x +=	1. Serially					
	2. Parallel reduction					
	3. Parallel asynchronous (atomic) updates					
x +=	4. Or any hybrid of these					
	<ul> <li>For what types / operators does this work?</li> </ul>					
	<ul> <li>"+" is associative for 1,2 rest of RHS is "simple"</li> </ul>					
<b>V</b>	<ul> <li>and commutative for 3</li> </ul>					

Using x for C[i,j] here

## Communication Avoiding Version (using a "1.5D" decomposition)



- Divide p into c groups. Replicate particles within group.
  - First row responsible for updating all by orange, second all by green,...
- Algorithm: shift copy of n/(p\*c) particles to the left
  - Combine with previous data before passing further level (log steps)
- Reduce across c to produce final value for each particle
- Total Computation: O(n<sup>2</sup>/p);
- Total Communication: O(log(p/c) + log c) messages,

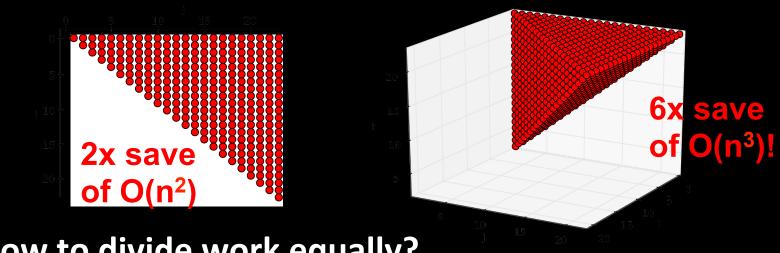
 $O(n^*(c/p+1/c))$  words

Driscoll, Georganas, Koanantakool, Solomonik, Yelick

Limit:  $c \le p^{1/2}$ 

### **Challenge: Symmetry & Load Balance**

- Force symmetry (f<sub>ij</sub> = -f<sub>ji</sub>) saves computation
- 2-body force matrix vs 3-body force cube

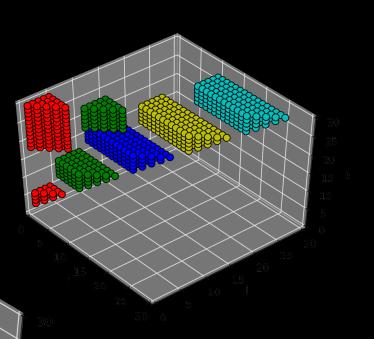


How to divide work equally?

Koanantakool & Yelick

### **3-Way N-Body Animation**

- p=5, n=30
- 6 particles per processor
- 5x5 subcubes



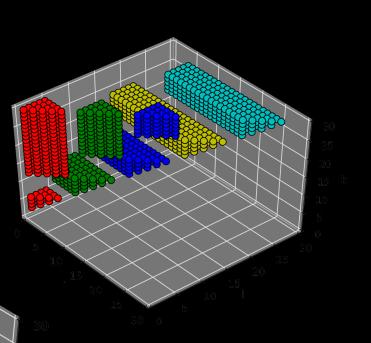
# Equivalent triplets in the big tetrahedron

Actual triplets

Koanantakool & Yelick

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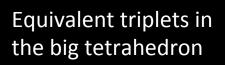


## Equivalent triplets in the big tetrahedron

Actual triplets Koanantakool & Yelick

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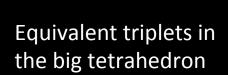
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Actual triplets Koanantakool & Yelick

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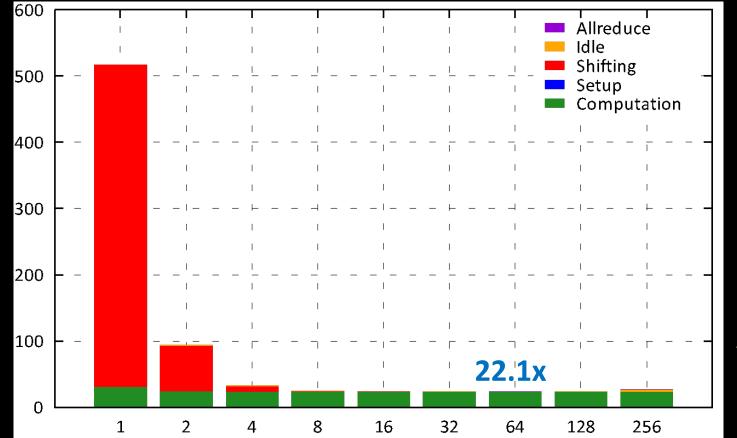
- p=5, n=30
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Actual triplets

Koanantakool & Yelick

#### Koanantakool & Yelick



## **3-Way N-Body Speedup**

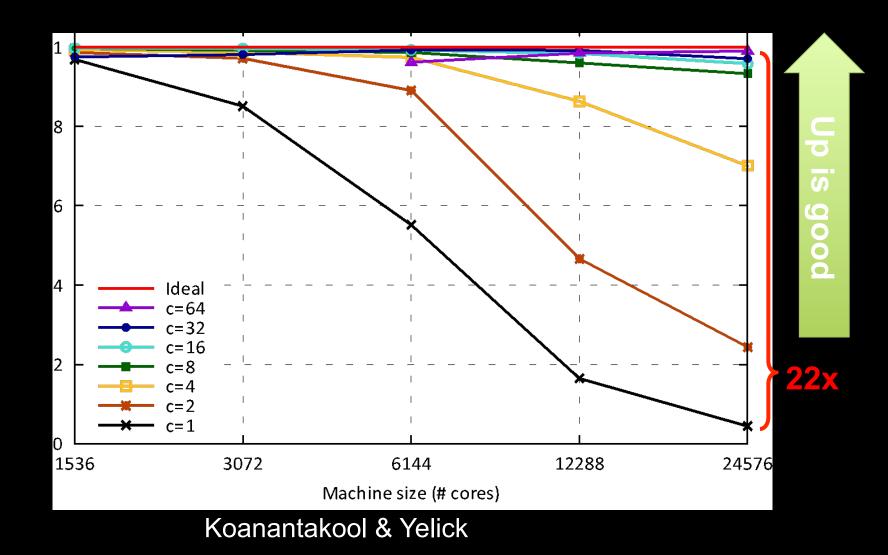
Cray XC30, 24k cores, 24k particles

ood

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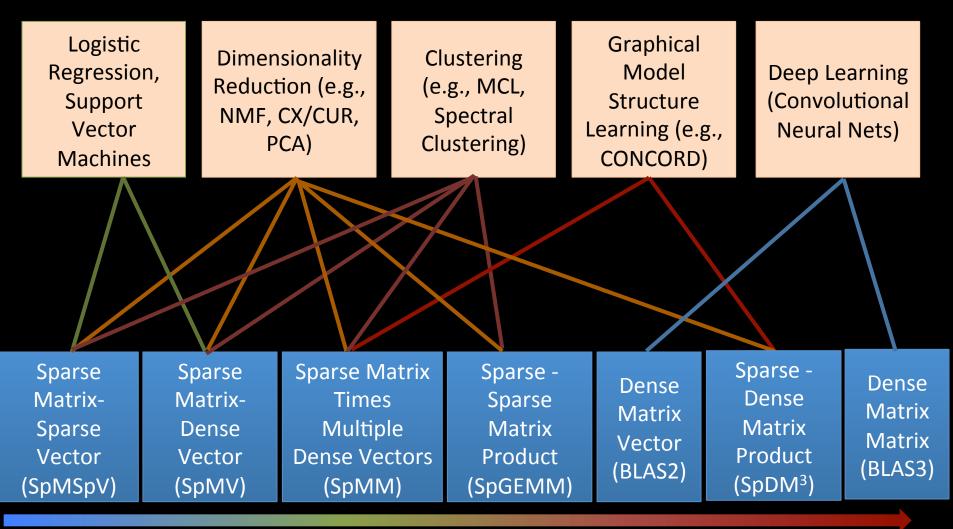
# **Strong Scaling of .5D Algorithns**



## **Analytics vs. Simulation Kernels:**

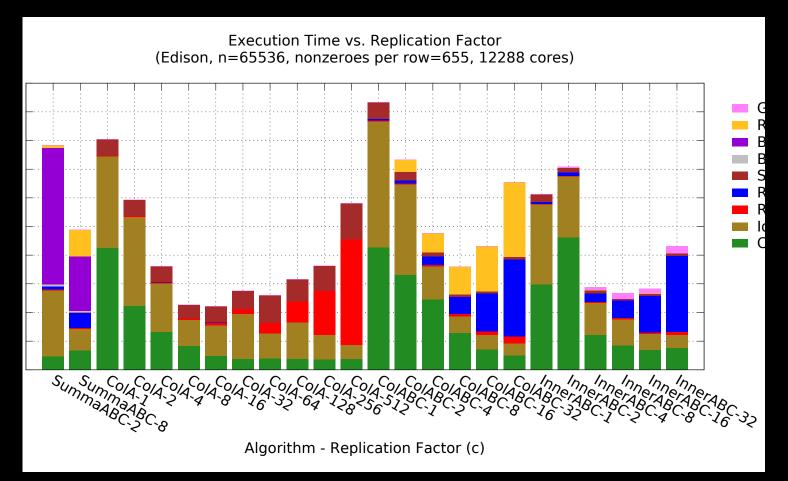
7 Giants of Data	7 Dwarfs of Simulation
Basic statistics	Monte Carlo methods
Generalized N-Body	Particle methods
Graph-theory	Unstructured meshes
Linear algebra 🛛 🗕	Dense Linear Algebra
Optimizations	Sparse Linear Algebra
Integrations	Spectral methods
Alignment	Structured Meshes

## **Machine Learning Mapping to Linear Algebra**



Aydin Buluc

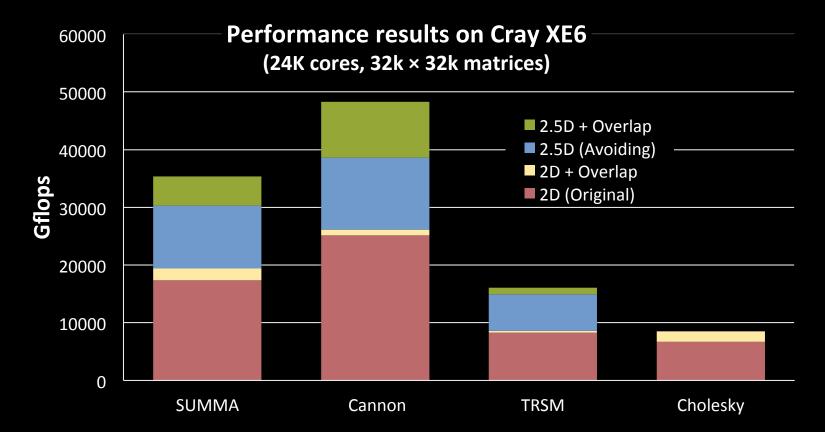
## **Sparse-Dense Matrix Multiply Too!**



Variety of algorithms that divide in or 2 dimensions

#### Koanantakool & Yelick

## **Communication Overlap Complements Avoidance**



# Even with communication-optimal algorithms (minimized bandwidth) there are still benefits to overlap and other things that speed up networks

SC'12 paper (Georganas, González-Domínguez, Solomonik, Zheng, Touriño, Yelick)

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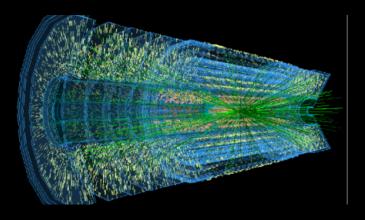
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## Data processing with special purpose hardware

- General trend towards specialization for continued performance growth
- Data processing (on raw data) will be first in science



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Particle Tracking with Neuromorphic chips

Computing in Detectors

Deep learning processors for image analysis

FPGAS for genome analysis

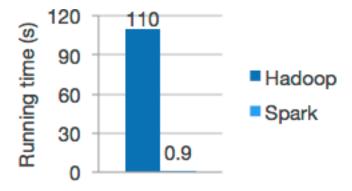
And can we also use these for simulation?

## **Productive Programming**



#### Speed

Run programs up to 100x faster than Hadoop MapReduce in memory, or 10x faster on disk.



- High failure rate
- Slow network
- Fast (local) disk

And Spark is still 10x+ slower than MPI

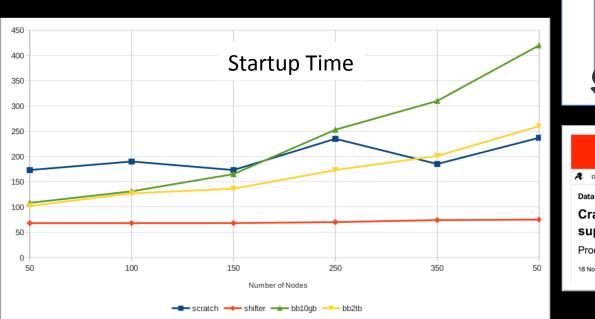
## Systems configured for data-intensive science



NERSC Cori has data partition (Phase 1, Haswell) pre-exascale (Phase 2, KNL preproduction) WAN-to-Cori optimized for streaming data: 100x faster from LCLS to Cori and Globus to CERN

## **Containers for HPC Systems**

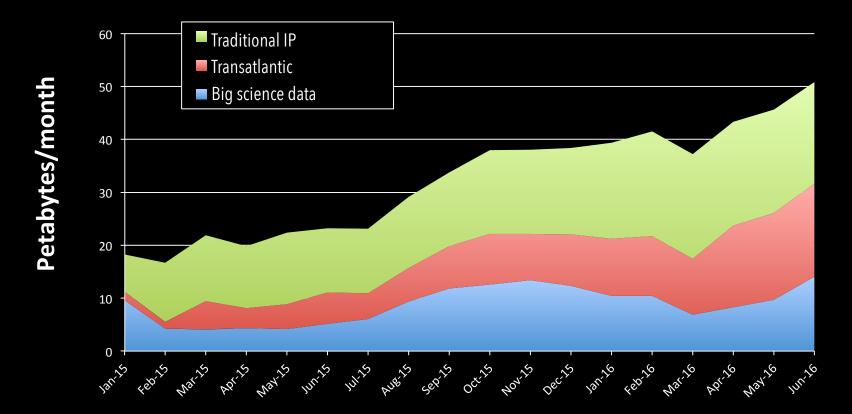
- Data analysis pipelines are often large, complex software stacks
- NERSC Shifter (with Cray), supports containers for HPC systems
- Used in HEP and NP projects STAR, LSST, DESI)





(ATLAS, ALICE,

## ESnet: Exponential data growth drives capacity





Science DMZ to deliver bandwidth to the end users OSCARS for bandwidth reservation



100 Exabytes/year by 2024!

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## **HPC Computing Policies**





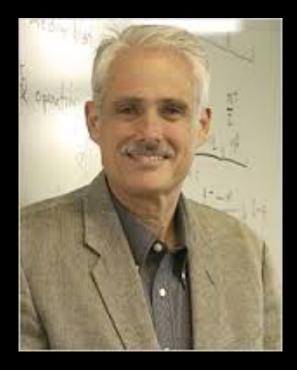
## **Cloud Computing Policies**



## Thanks to many collaborators

Eun-Jin Im Sam Williams Christian Bell Arvind Krishnamurthy Bert de Jong Jason Duell Carelton Miyamoto Nick Knight Horst Simon Jimmy Su Greg Balls Ed Givelberg Dan Rokhsar Chih-Po Wen Khaled Ibrahim Wei Chen Soumen Chakrabarti Lauren Smith Kaushik Datta Leonid Oliker Costin Iancu Ngeci Bowman Hongzhang Shan Edgar Solomonik Jeff Jones Jim Demmel Yili Zheng Siu Man Yau Shoaib Kamil Paul Hargrove Bob Lucas Etienne Deprit Amir Kamil Bill Carlson Eric Roman Noah Treuhaft Dan Bonachea C.J. Lin Tarek El-Ghazawi Rajesh Nishtala Erich Strohmaier Sabrina Merchant Mani Narayanan Randi Thomas Rich Vuduc Harsha Simhadri Deborah Weisser Penporn Kaonantakool Jarrod Chapman Steve Steinberg Evangelos Georganas Brian Kazian Michael Driscoll David Ozog

## And to Ken Kennedy



"Take care of your students and the rest will take care of itself."