

# Avoiding, Hiding and Managing Communication at the Exascale

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U.S. DEPARTMENT OF  
**ENERGY**

Office of  
Science

Computing Sciences Area

# Deploy Exascale Systems at NERSC



**NERSC: the broadest, most open, widely used computing center in DOE**

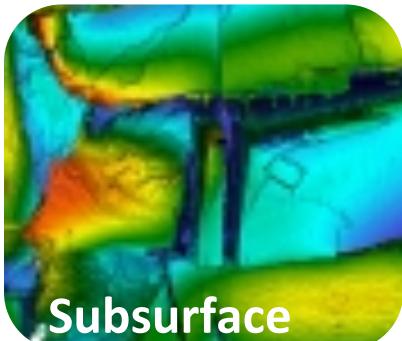
- 7000 users, 2000 publications / year, 700 codes, 5 associated Nobels
- Cori system delivers petascale science for simulation and data

**Exascale goal: broad scientific impact through exascale at NERSC**

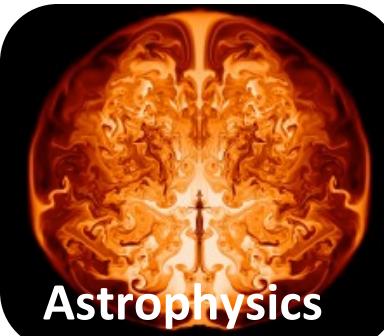
# Berkeley Lab Priorities in Exascale Science



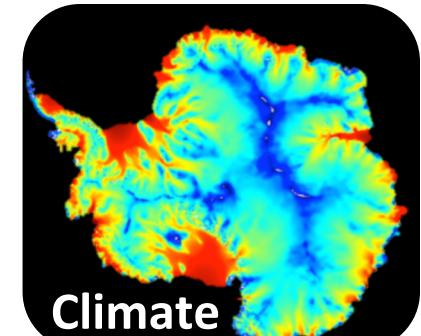
Accelerators



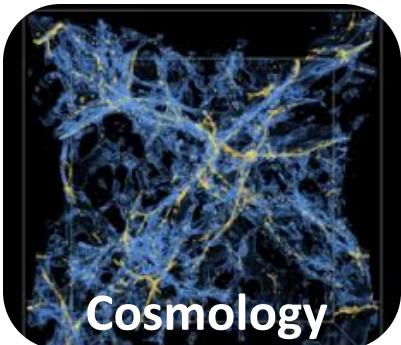
Subsurface



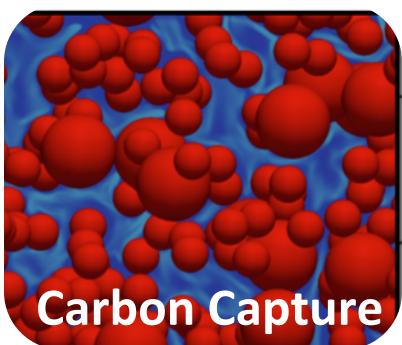
Astrophysics



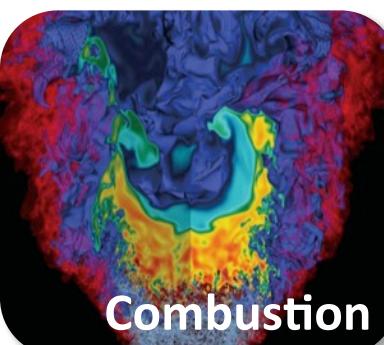
Climate



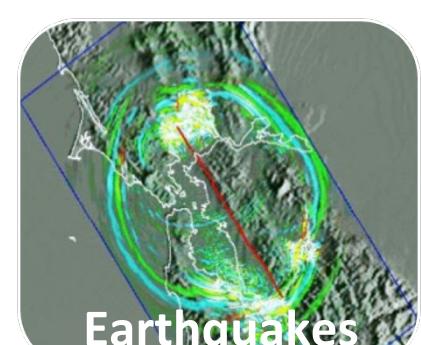
Cosmology



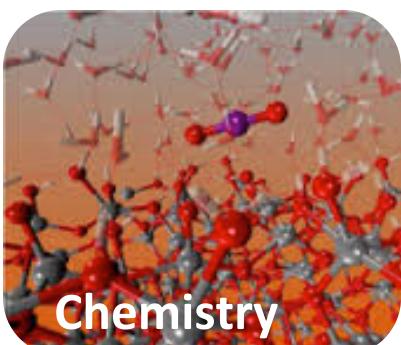
Carbon Capture



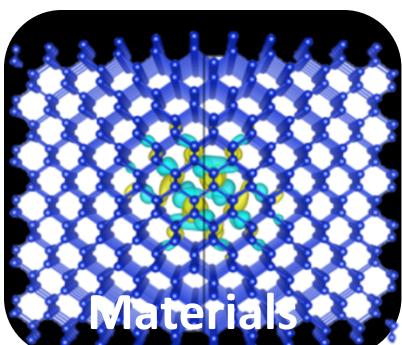
Combustion



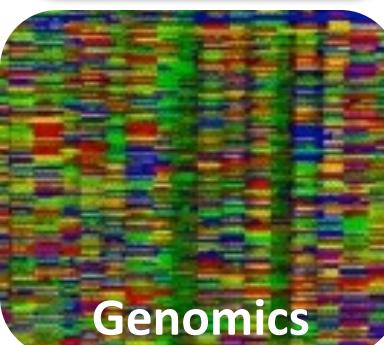
Earthquakes



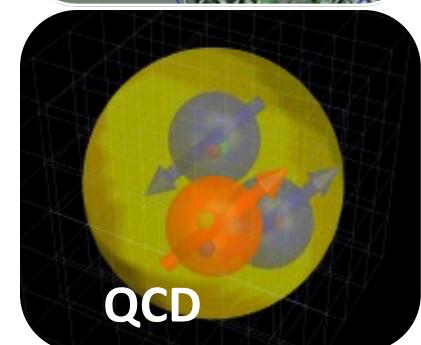
Chemistry



Materials



Genomics

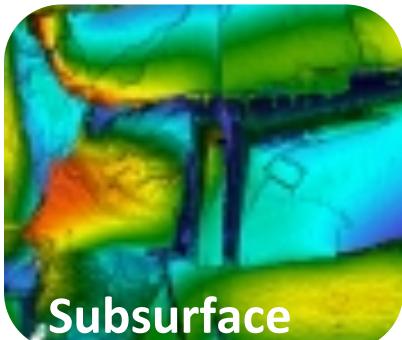


QCD

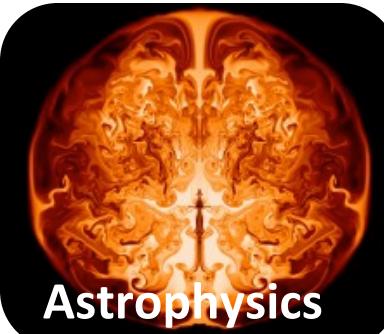
# Berkeley Lab Priorities in Exascale Science



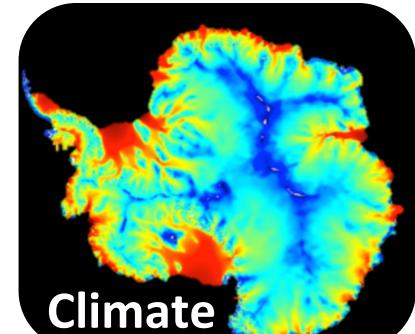
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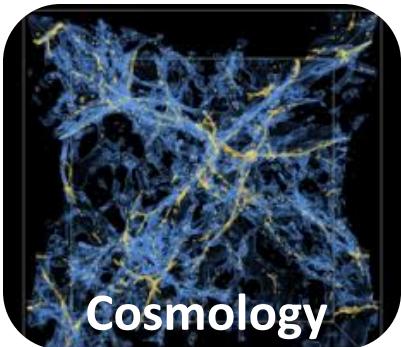
Subsurface



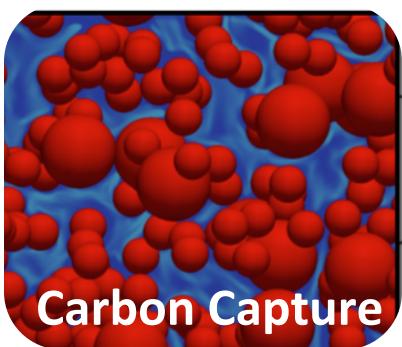
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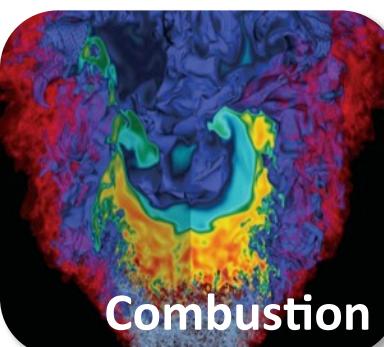
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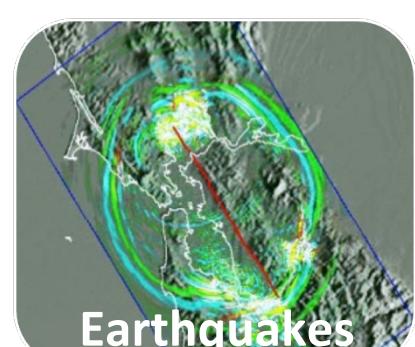
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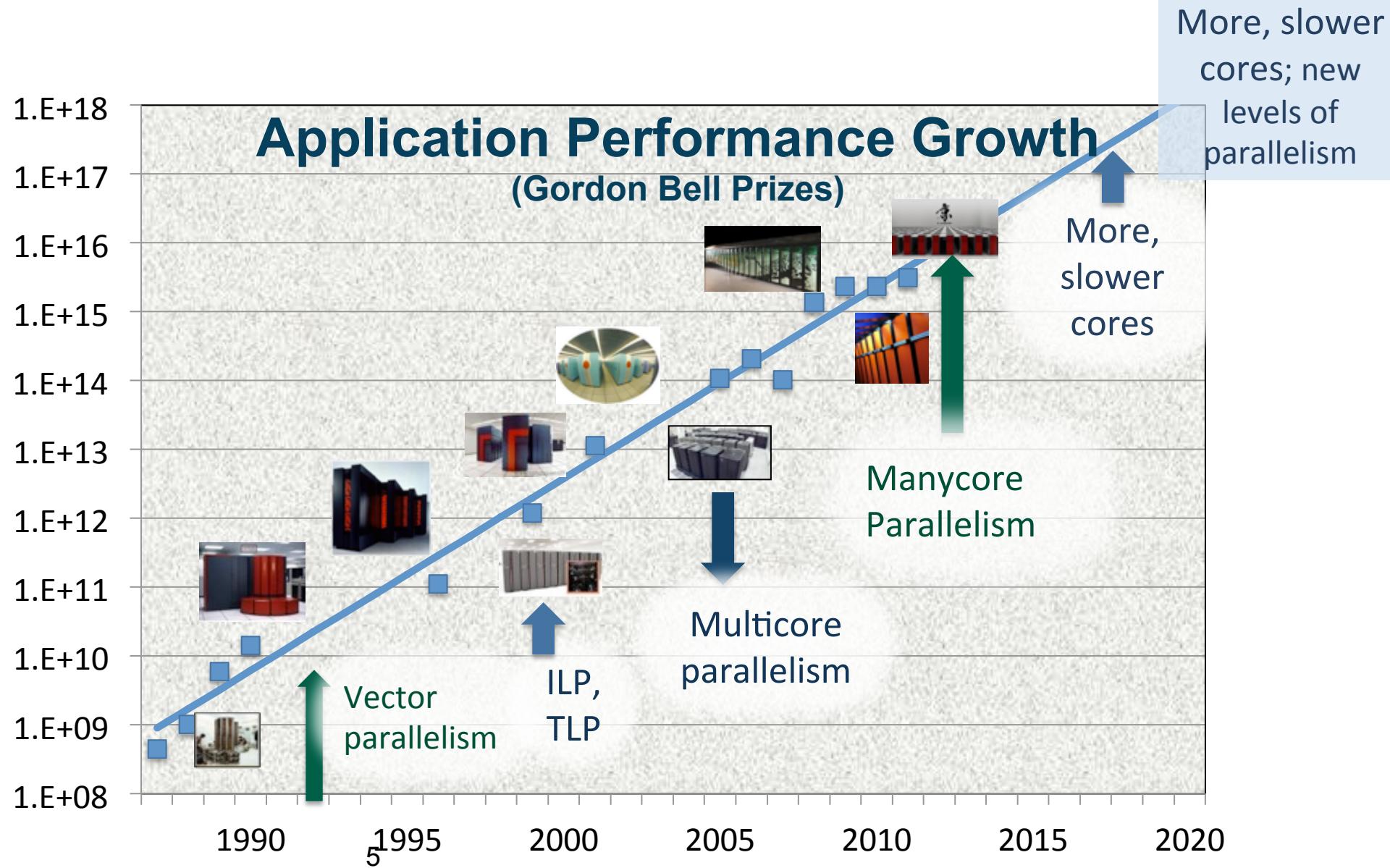
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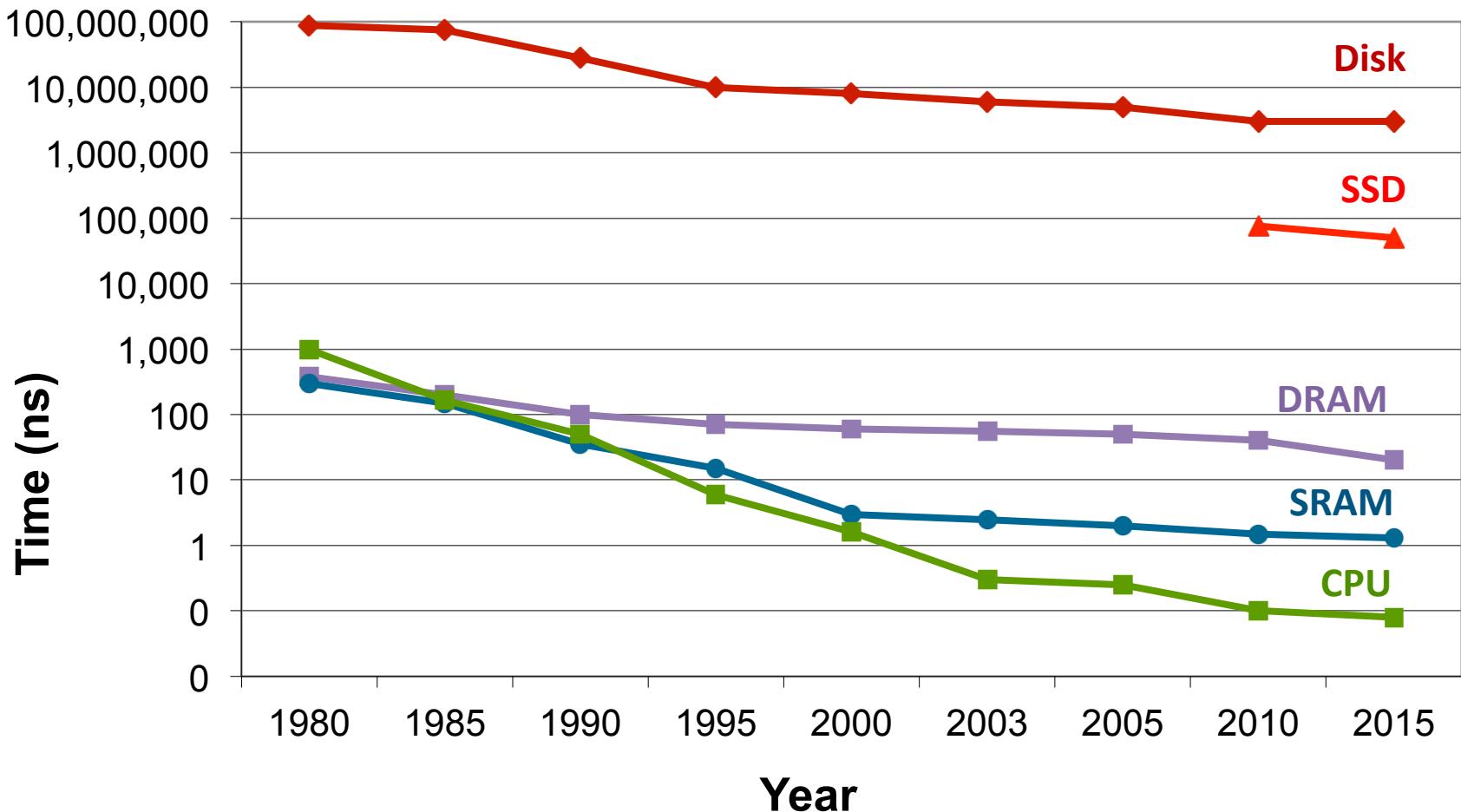
All the above will use Adaptive Mesh Refinement  
(AMR) mathematics and software,  
a method pioneered at Berkeley Lab

# Hardware Trend: More Parallelism at Lower Levels



# Data Movement is Expensive

## CPU cycle time vs memory access time

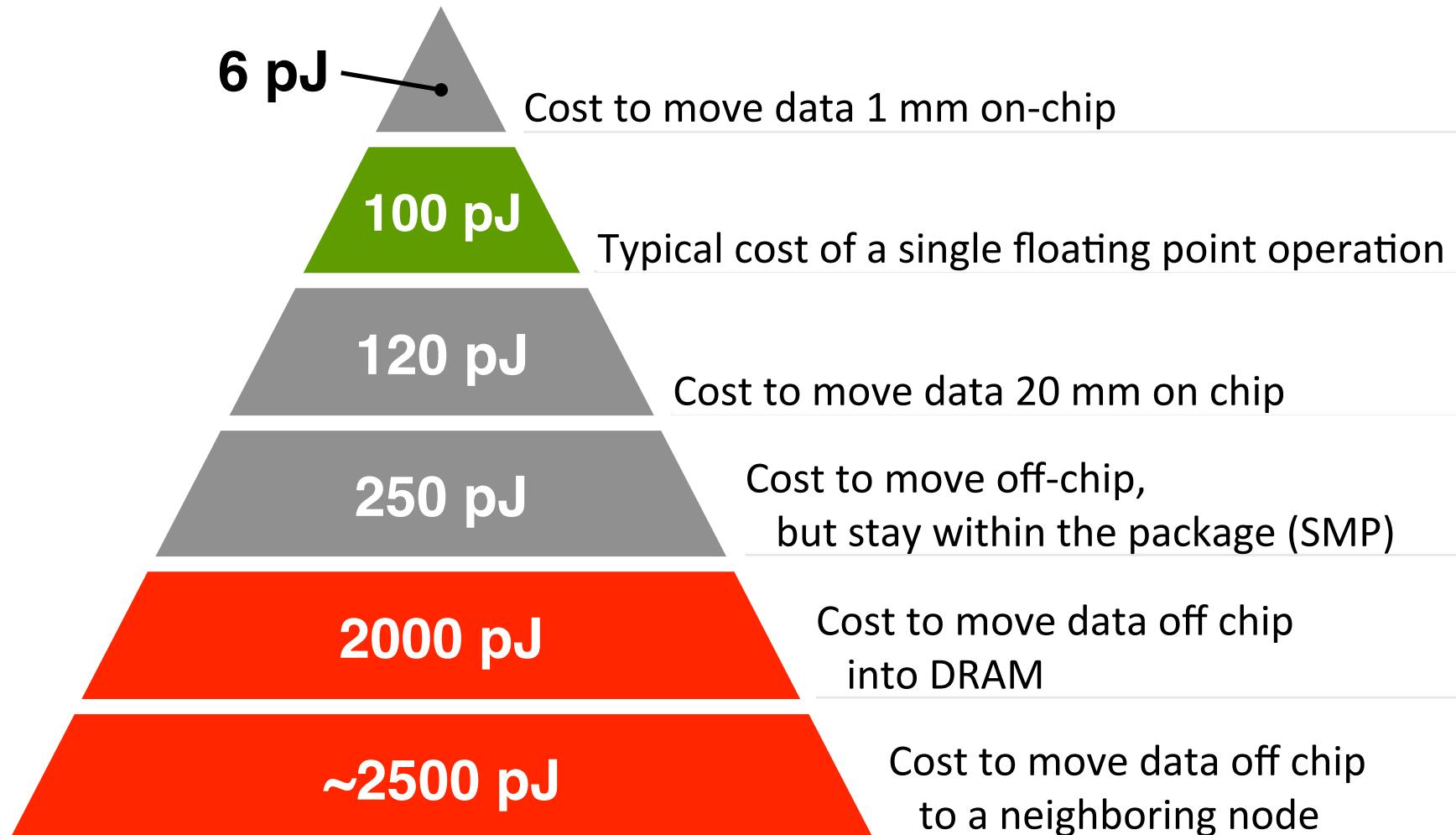


Sources:

<http://csapp.cs.cmu.edu/2e/figures.html>, <http://csapp.cs.cmu.edu/3e/figures.html>

# Data Movement is Expensive

## Hierarchical energy costs.



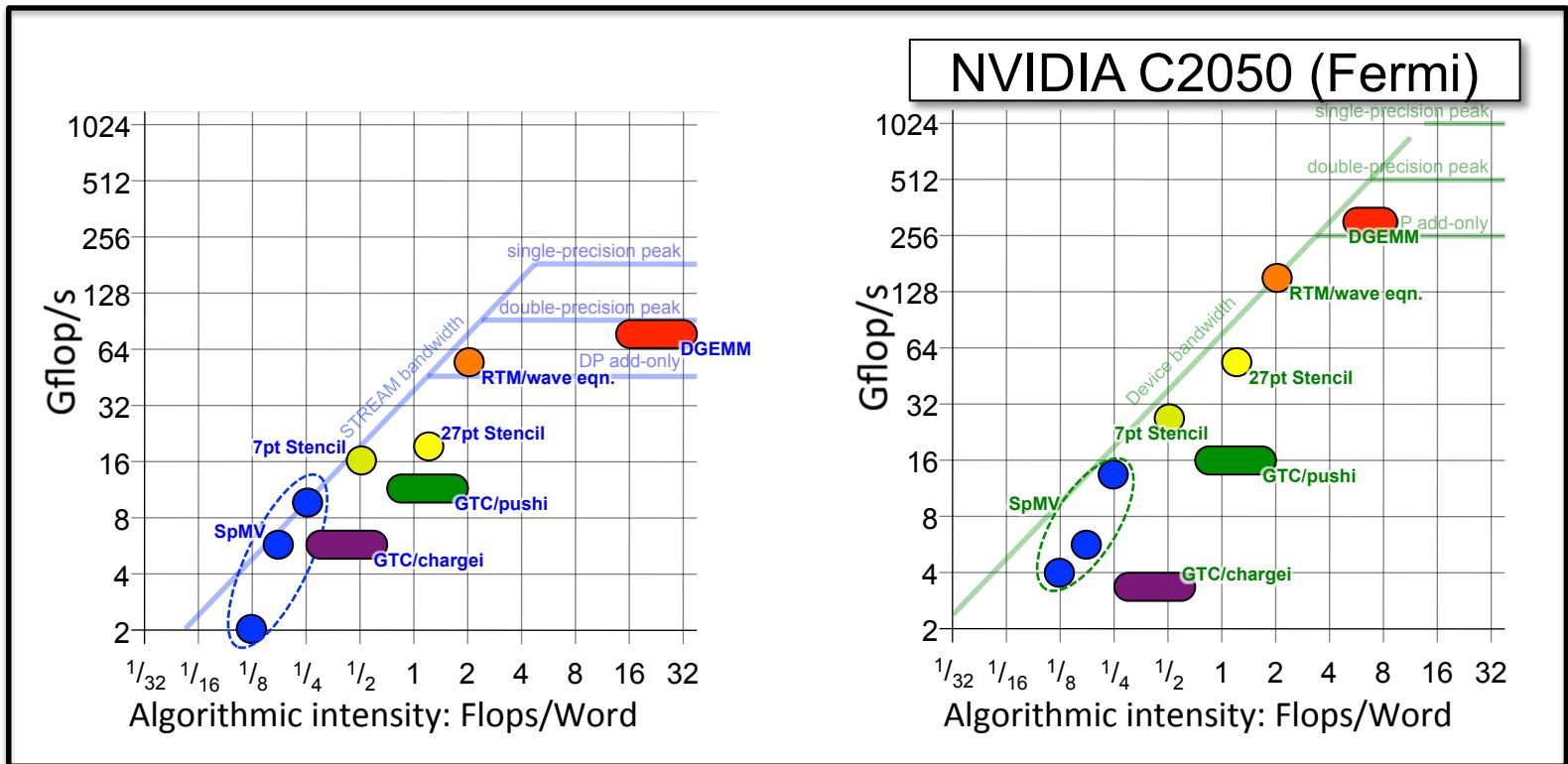
# Summary

- More parallelism
- Diversity of processors / accelerators
- Communication (data movement) is expensive

# **Compilers and Autotuning**

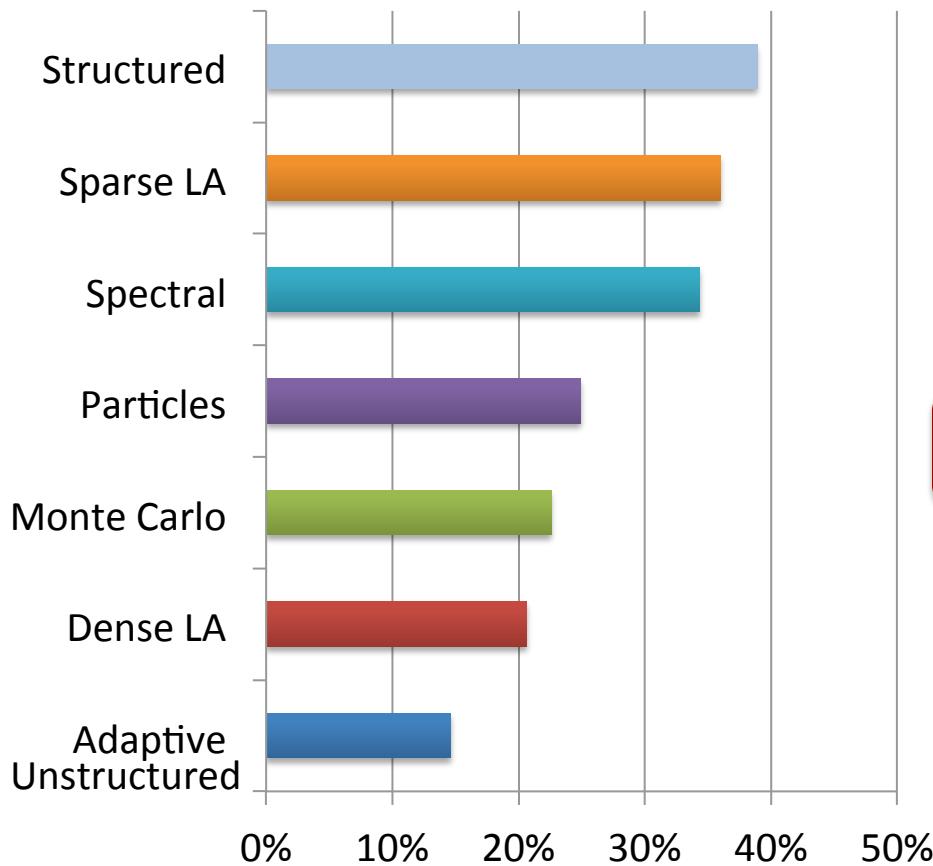
# Autotuning: Write Code Generators

- Two “unsolved” compiler problems:
  - dependence analysis and Domain-Specific Languages help with this
  - ✓ accurate performance models Autotuning avoids this problem
- Autotuners are code generators plus search



# Libraries vs. DSLs (domain-specific languages)

NERSC survey: what motifs do they use?



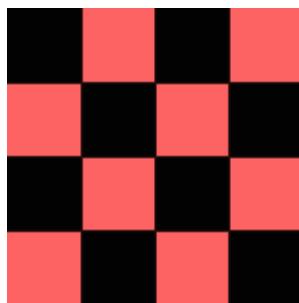
What code generators do we have?

<b>Dense Linear Algebra</b>	<b>Atlas</b>
Spectral Algorithms	FFTW, Spiral
Sparse Linear Algebra	OSKI
Structured Grids	TBD
Unstructured Grids	
Particle Methods	
Monte Carlo	

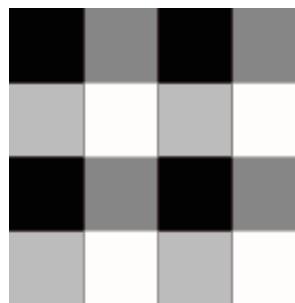
Stencils are both the most important motifs and a gap in our tools

# Approach: Small Compiler for Small Language

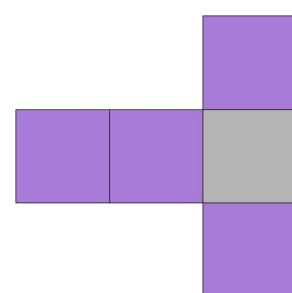
- **Snowflake: A DSL for Science Stencils**
  - Domain calculus inspired by Titanium, UPC++, and AMR in general



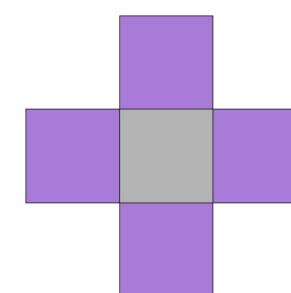
(a) Red-Black tiling



(b) 4-color tiling



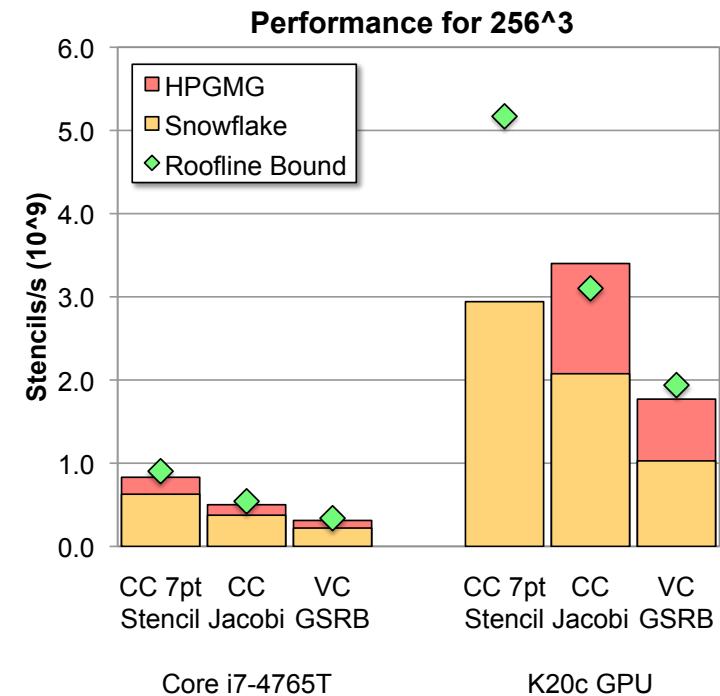
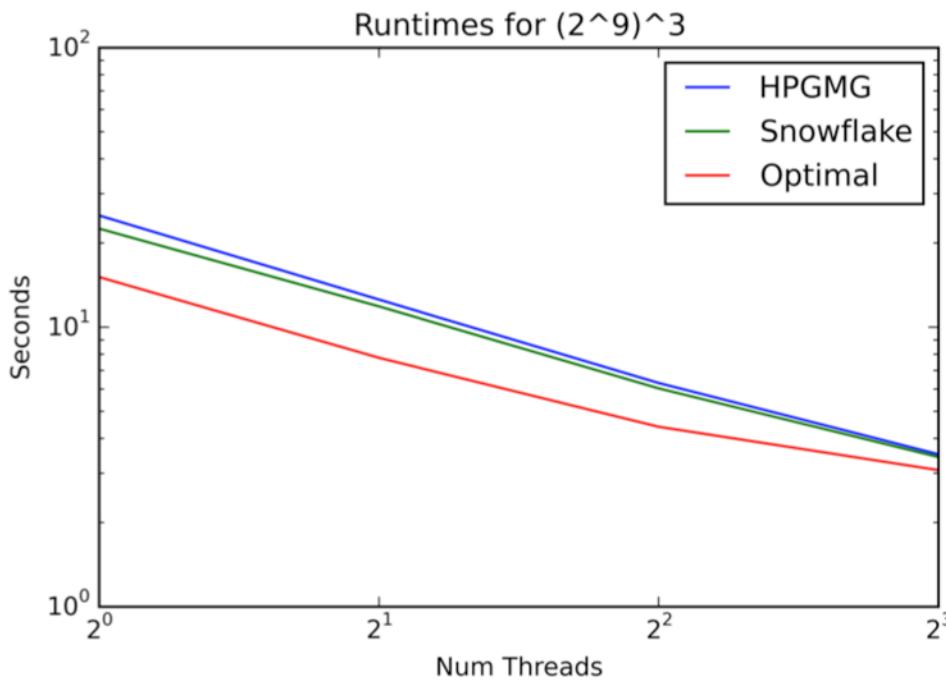
(c) Asymmetric stencil used  
near mesh boundary



(d) 5-point Jacobi stencil

- **Complex stencils: red/black, asymmetric**
- **Update-in-place while preserving provable parallelism**
- **Complex boundary conditions**

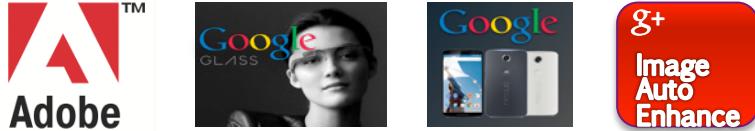
# Snowflake Performance



- **Performance on the HPGMG application benchmark using all the features of Snowflake**
- **Competitive with hand-optimized performance**
- **Within 2x of optimal roofline**

# DSLs popular outside scientific computing

## Developed for Image Processing



- 10+ FTEs developing Halide
- 50+ FTEs use it; > 20 kLOC

## HPGMG (Multigrid on Halide)

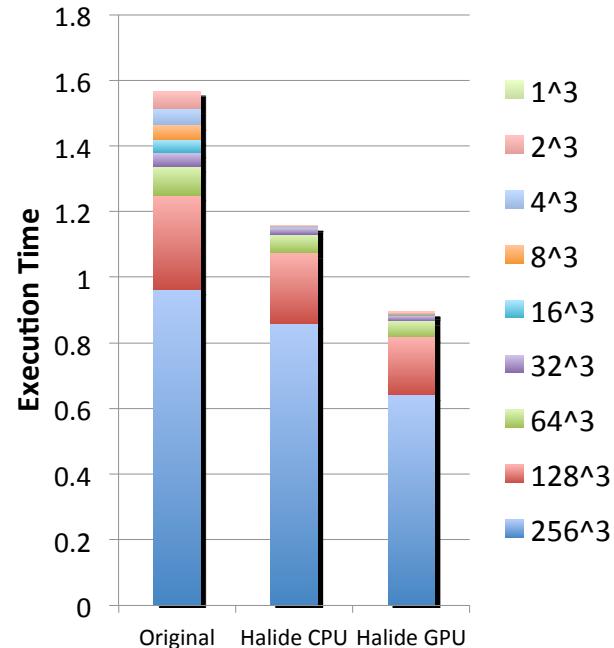
- **Halide Algorithm by domain expert**

```
Func Ax_n("Ax_n", lambda("lambda"), chebyshev("chebyshev"));
Var i("i"), j("j"), k("k");
Ax_n(i,j,k) = lambda(i,j,k)*x_n(i,j,k) - b*h2inv*(  
    beta_i(i,j,k)*valid(i-1,j,k)*x_n(i,j,k) + x_n(i-1,j,k)) - 2.0*x_n(i,j,k));  
    + beta_j(i,j,k)*valid(i,j-1,k)*x_n(i,j,k) + x_n(i,j-1,k)) - 2.0*x_n(i,j,k));  
    + beta_k(i,j,k)*valid(i,j,k-1)*x_n(i,j,k) + x_n(i,j,k-1)) - 2.0*x_n(i,j,k));  
    + beta_i(i+1,j,k)*valid(i+1,j,k)*x_n(i,j,k) + x_n(i+1,j,k)) - 2.0*x_n(i,j,k));  
    + beta_j(i,j+1,k)*valid(i,j+1,k)*x_n(i,j,k) + x_n(i,j+1,k)) - 2.0*x_n(i,j,k));  
    + beta_k(i,j,k+1)*valid(i,j,k+1)*(x_n(i,j,k) + x_n(i,j,k+1)) - 2.0*x_n(i,j,k));  
lambda(i,j,k) = 1.0 / (*alpha(i,j,k) - b*h2inv(*  
beta_i(i,j,k)*valid(i-1,j,k) - 2.0f)  
+ beta_j(i,j,k)*valid(i,j-1,k) - 2.0f)  
+ beta_k(i,j,k)*valid(i,j,k-1) - 2.0f)  
+ beta_i(i+1,j,k)*valid(i+1,j,k) - 2.0f)  
+ beta_j(i,j+1,k)*valid(i,j+1,k) - 2.0f));  
+ beta_k(i,j,k+1)*valid(i,j,k+1) - 2.0f));
chebyshev(i,j,k) = x_n(i,j,k) + c1*(x_n(i,j,k)-Ax_n(i,j,k))+  
c2*lambda(i,j,k)*(rhs(i,j,k)-Ax_n(i,j,k));
```

- **Halide Schedule either**
  - Auto-generated by autotuning with opentuner
  - Or hand created by an optimization expert

## Halide performance

- Autogenerated schedule for CPU
- Hand created schedule for GPU
- No change to the algorithm



# **Algorithms for the Hardware**

# 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

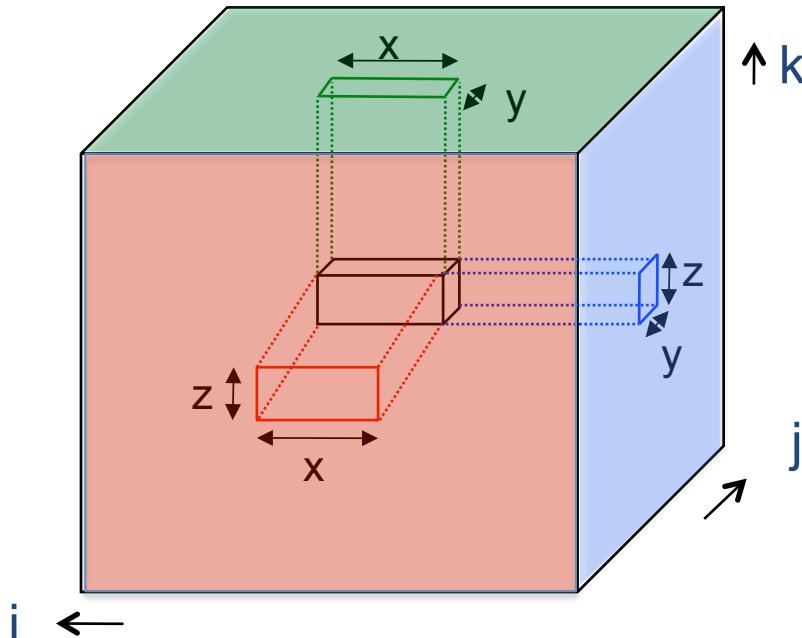
**Lesson: Never waste fast memory**

**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] ...

# Using .5D ideas on N-body

- **n particles, k-way interaction.**
  - Molecules, stars in galaxies, etc.
- **Most common: 2-way N-body**

```
for t timesteps
```

```
    forall i1, ..., ik
```

```
        force[i1] += interact(particle[i1], ..., particle[ik])
```

```
    forall i
```

```
        move(particle[i], force[i])
```

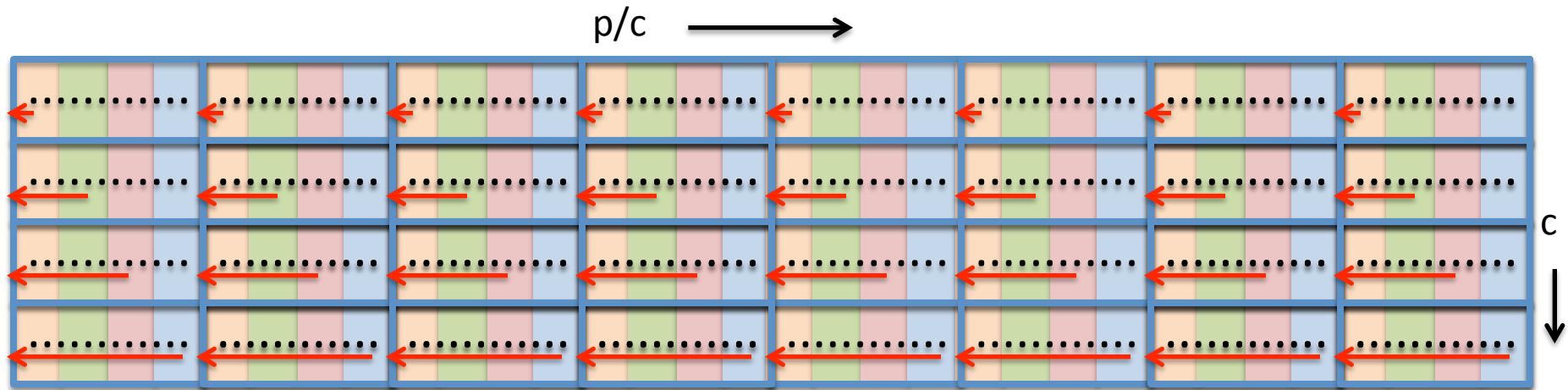
$O(n^k)$ .

- **Best algorithm is to divide n particles into p groups??**



No!

# Communication Avoiding 2-way N-body (using a “1.5D” decomposition)

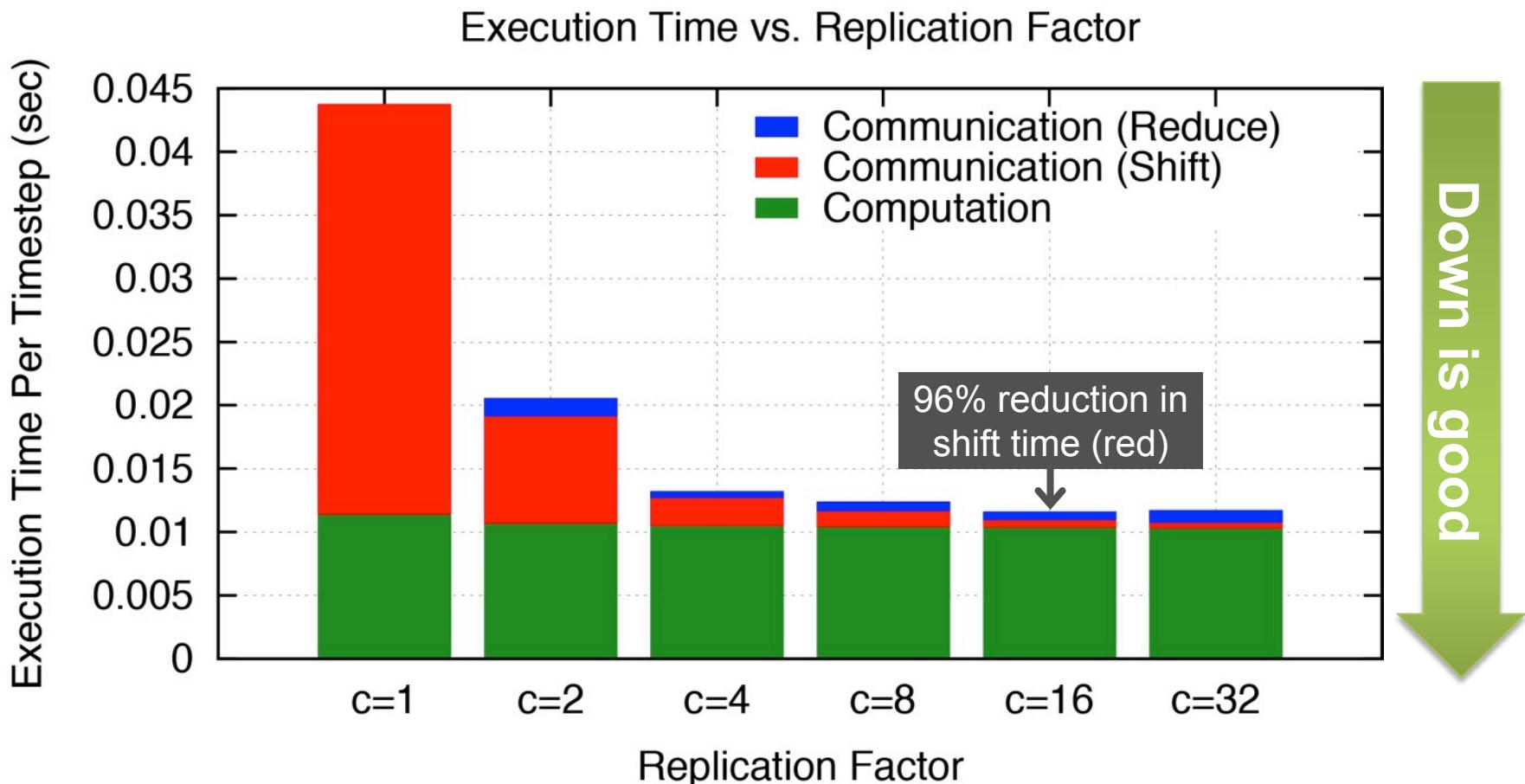


- Divide  $p$  into  $c$  groups
- Replicate particles across groups
- Repeat: shift copy of  $n/(p*c)$  particles to the left within a group
- Reduce across  $c$  to produce final value for each particle

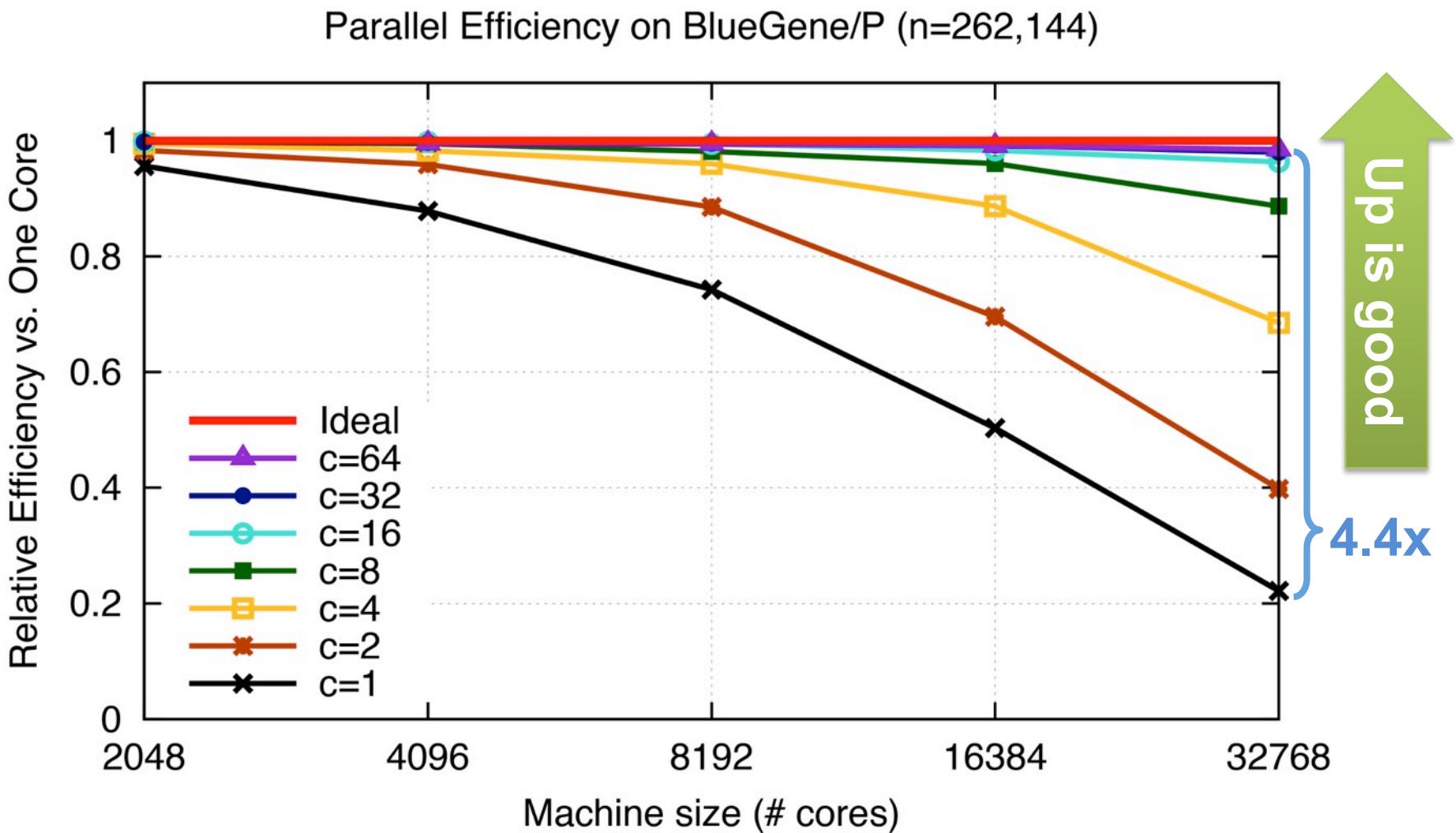
Total Communication:  $O(\log(p/c) + \log c)$  messages,  
 $O(n*(c/p+1/c))$  words

# Less Communication..

- Cray XE-6; n=24K particles, p=6K cores

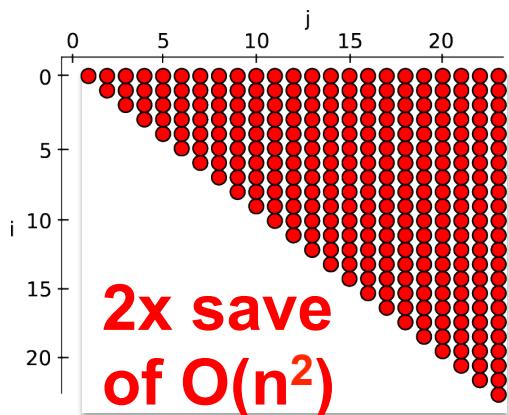


# Strong Scaling

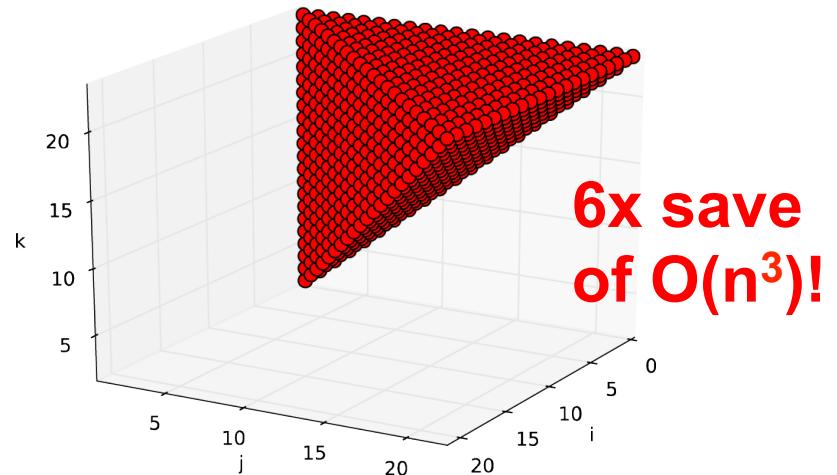


# Challenge: Symmetry & Load Balance

- Force symmetry ( $f_{ij} = -f_{ji}$ ) saves computation
- 2-body force matrix vs 3-body force cube



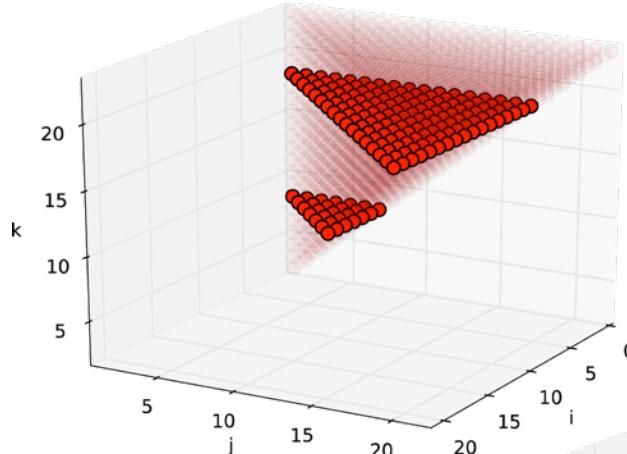
2x save  
of  $O(n^2)$



6x save  
of  $O(n^3)$ !

- How to divide work equally?

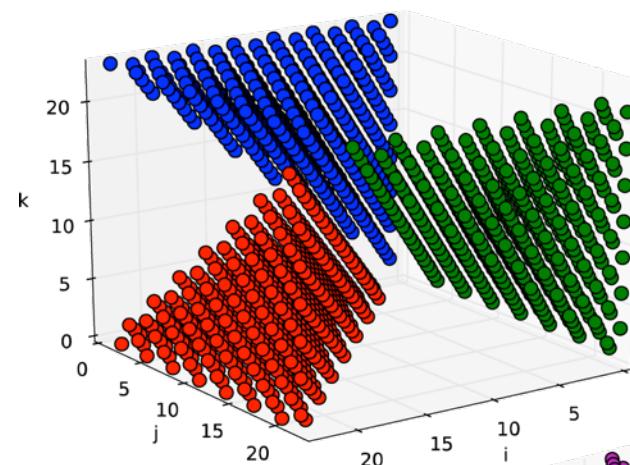
# All-triplets 3-body: Challenges



[Li et al. 2006]  
[Li et al. 2008]

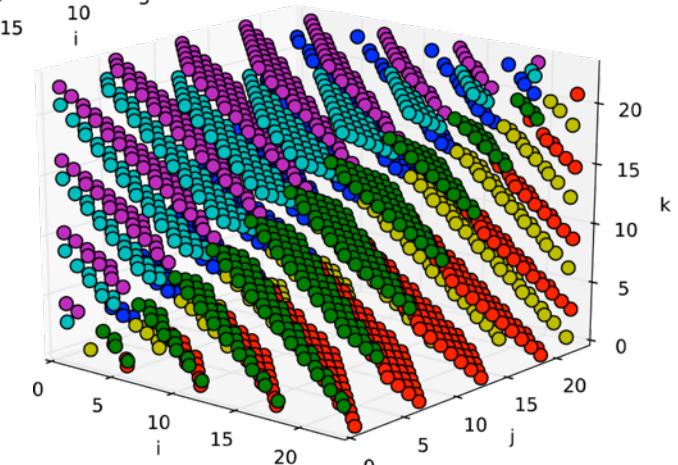
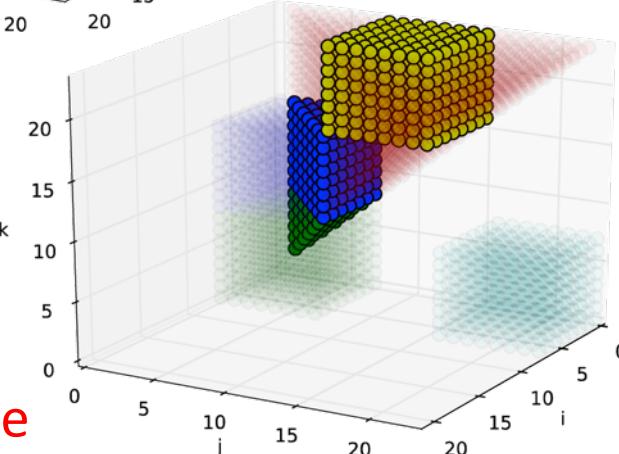
Symmetry  
Load balance

Communication?



[Sumanth et al. 2007]

Symmetry  
Load balance  
Communication?

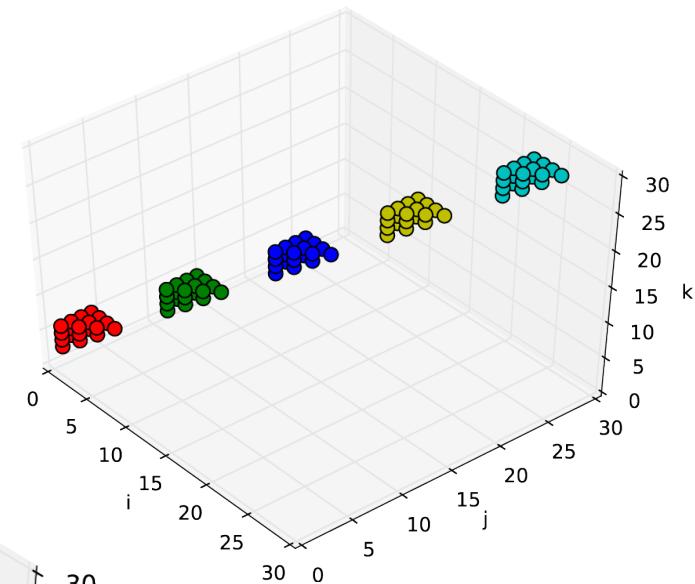
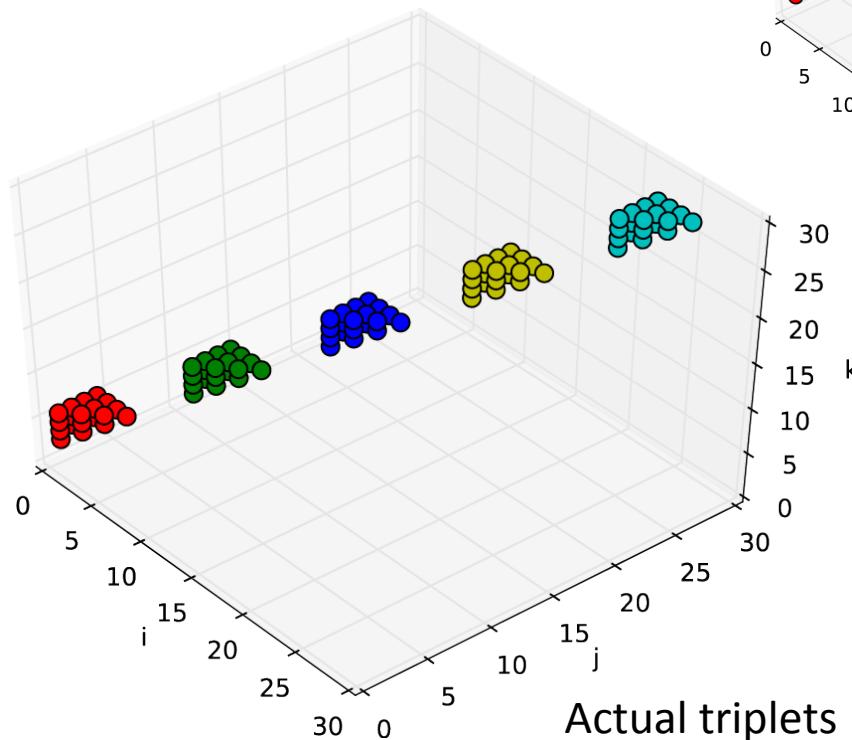


\* Colors have no special meaning --  
for illustration purpose only.

# CA 3-body

[Koanantakool and Yellick 2014]

- **p=5 (in colors)**
- **6 particles per processor**
- **5x5 subcubes**

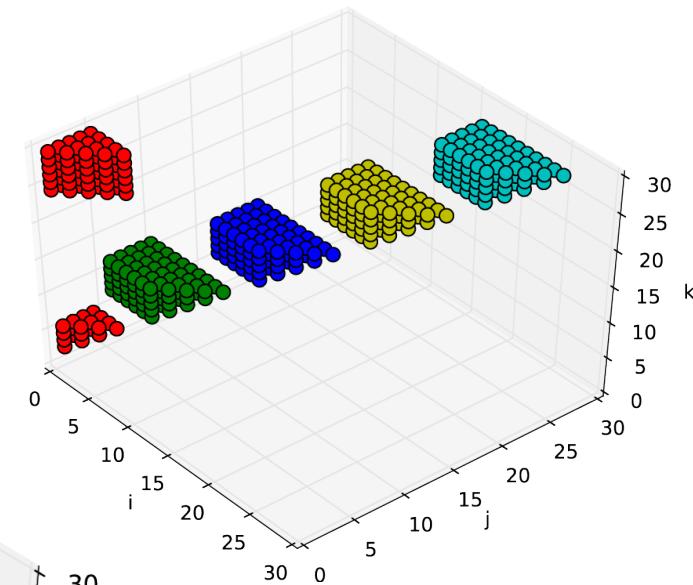
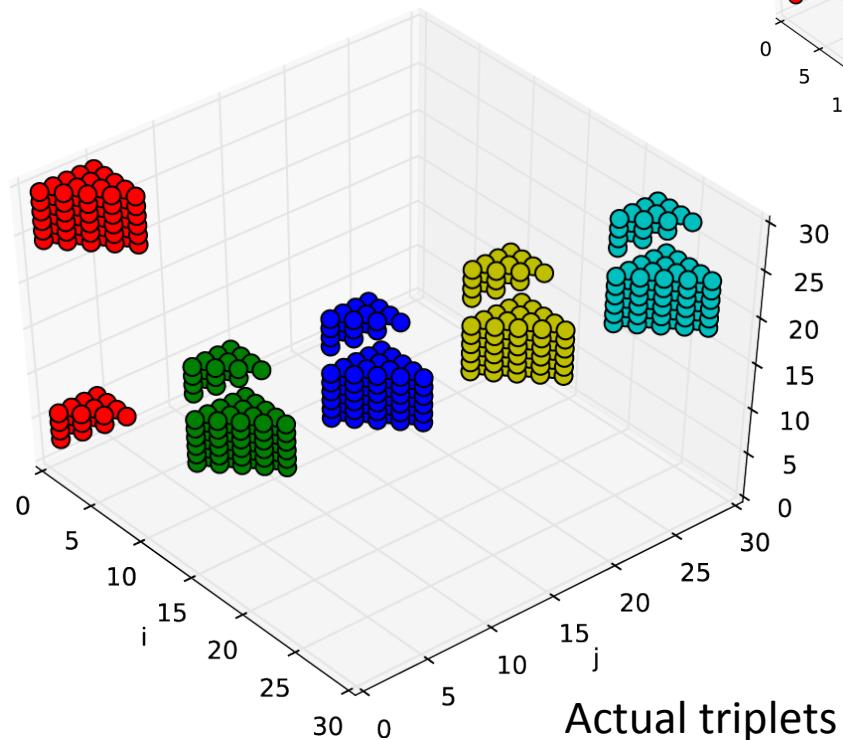


Equivalent triplets in  
the big tetrahedron

# CA 3-body

[Koanantakool and Yellick 2014]

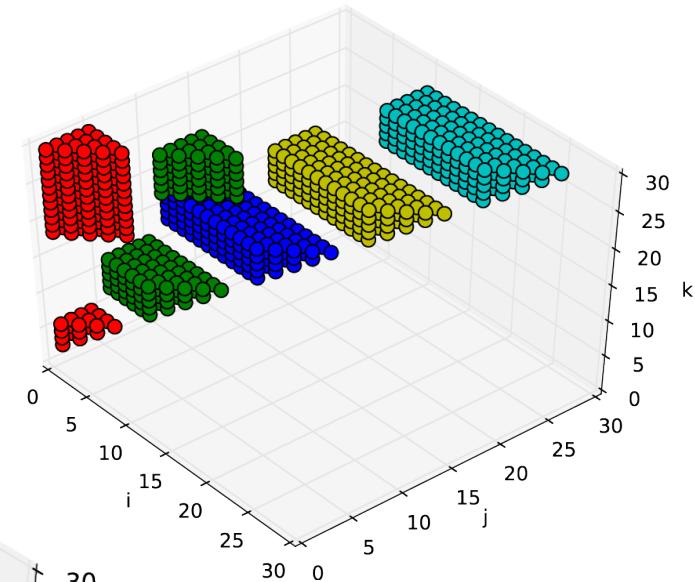
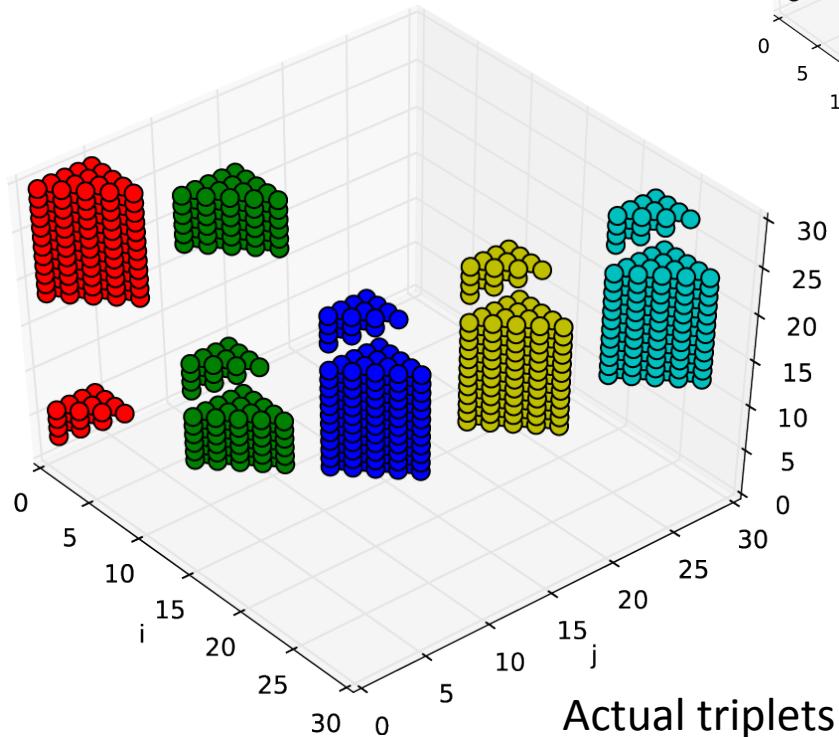
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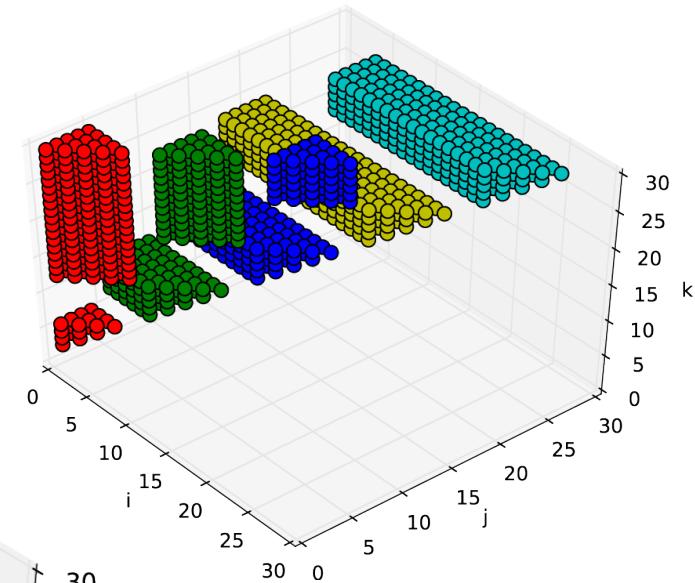
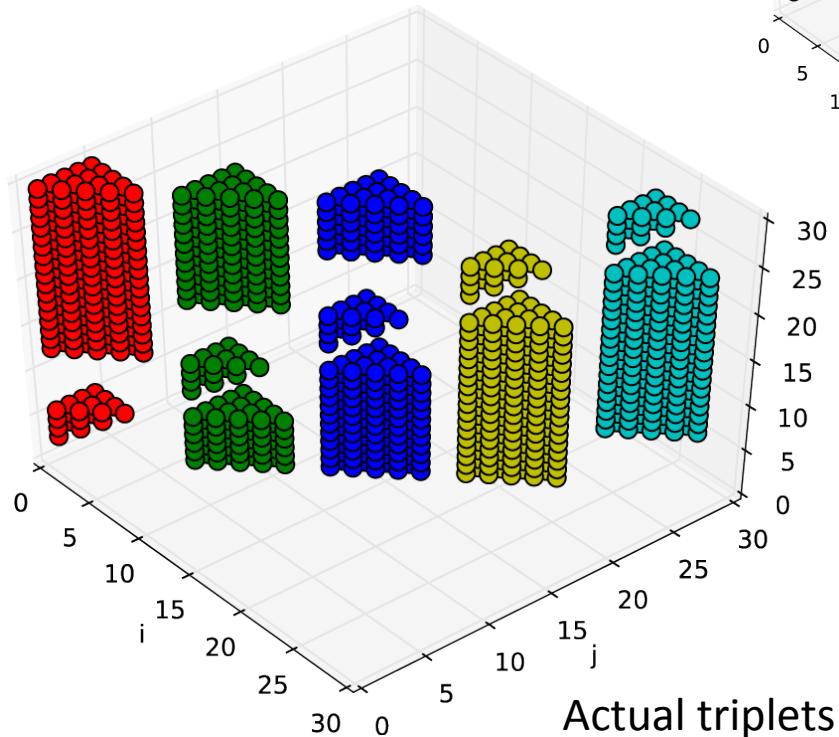


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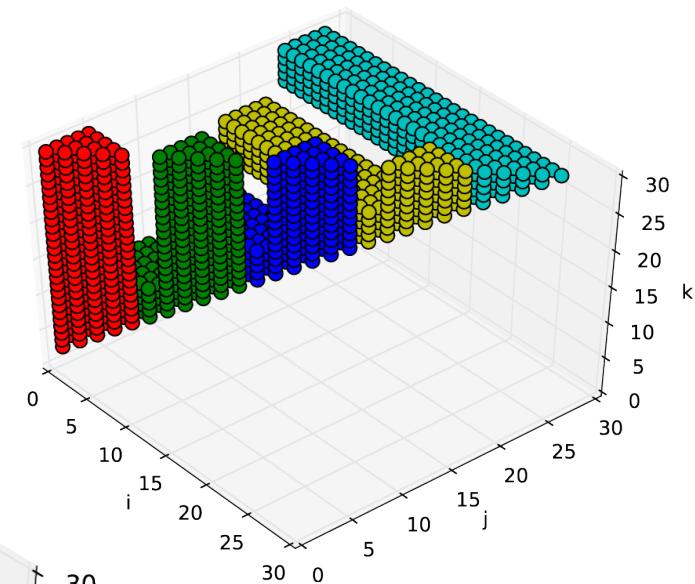
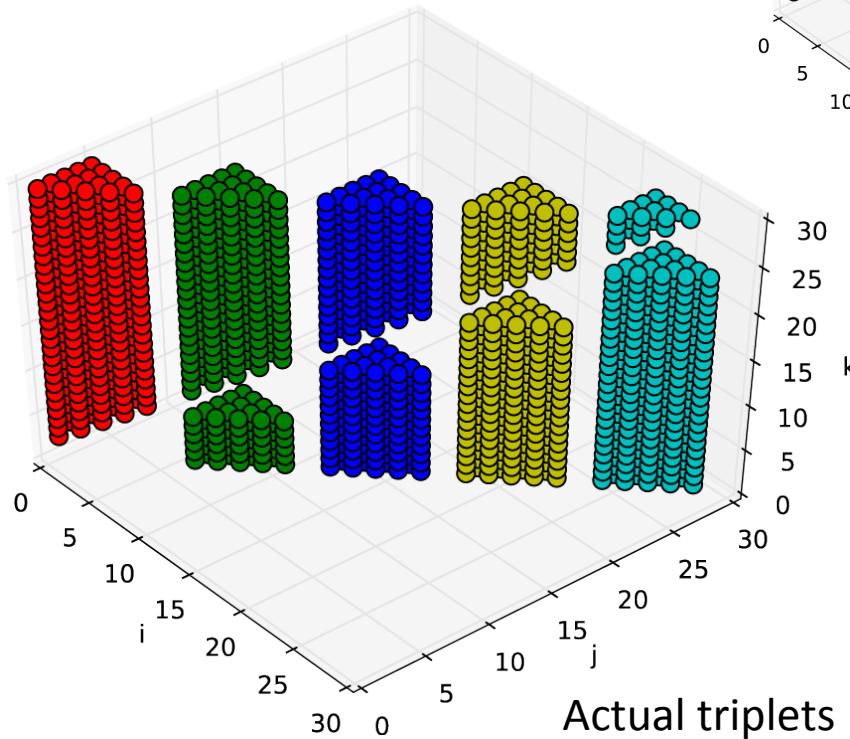
- **p=5 (in colors)**
- **6 particles per processor**
- **5x5 subcubes**



# CA 3-body

[Koanantakool and Yellick 2014]

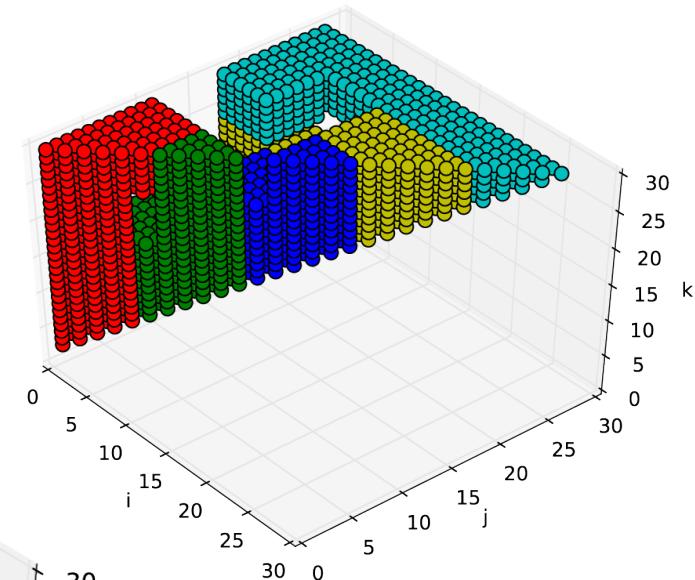
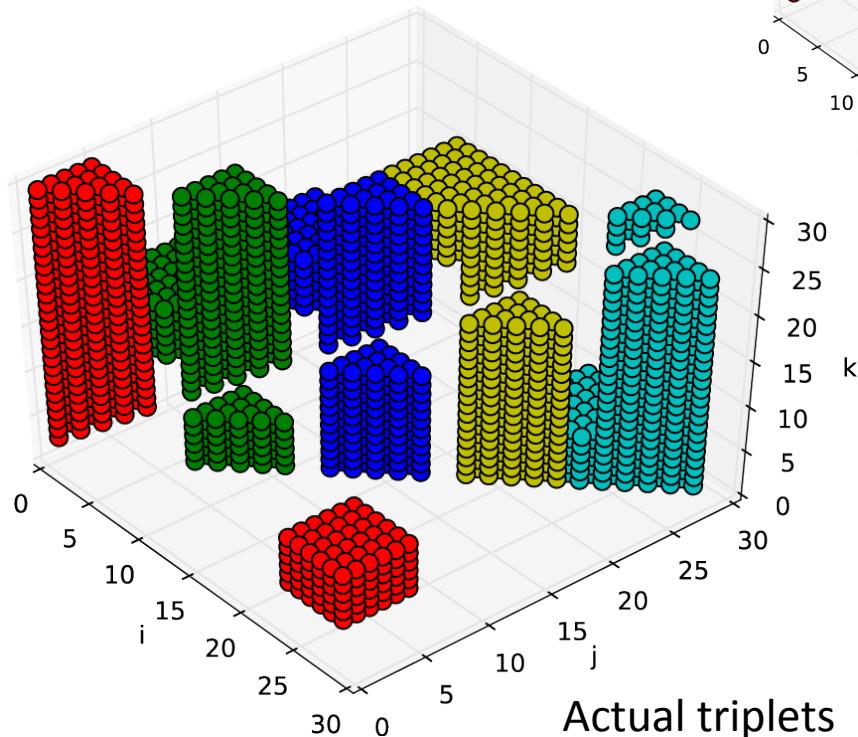
- **p=5 (in colors)**
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# CA 3-body

[Koanantakool and Yellick 2014]

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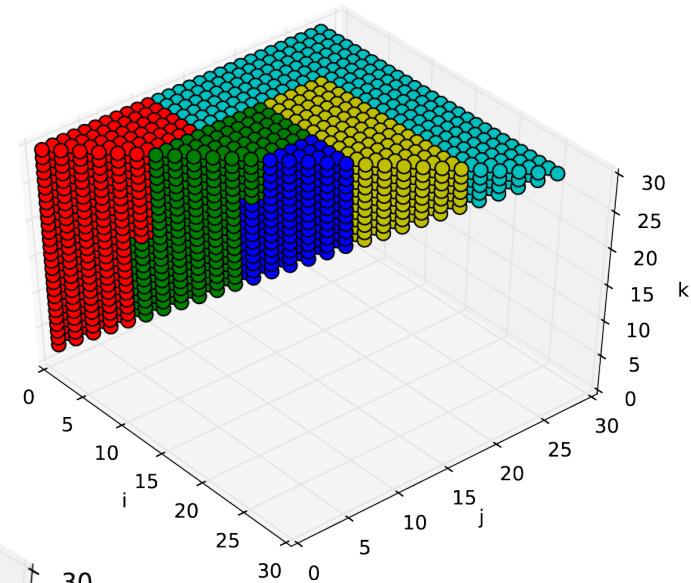
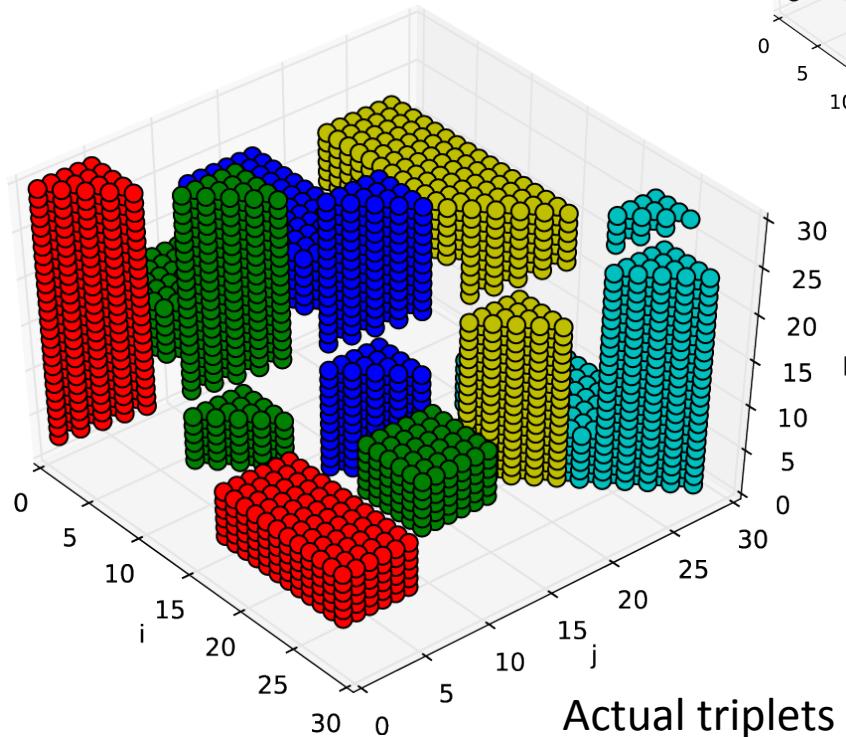


Equivalent triplets in  
the big tetrahedron

# CA 3-body

[Koanantakool and Yellick 2014]

- $p=5$  (in colors)
- 6 particles per processor
- $5 \times 5$  subcubes

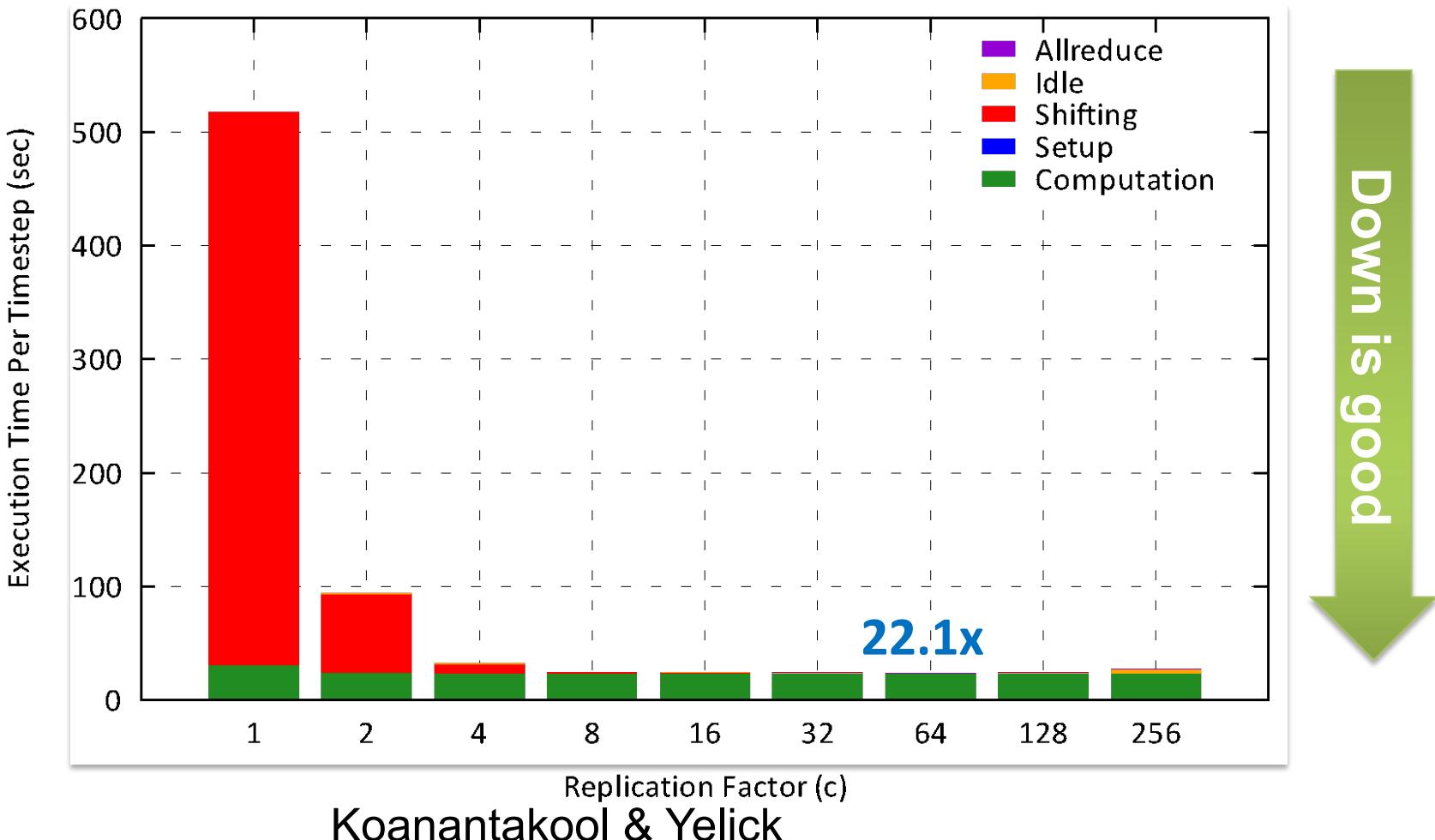


Equivalent triplets in  
the big tetrahedron

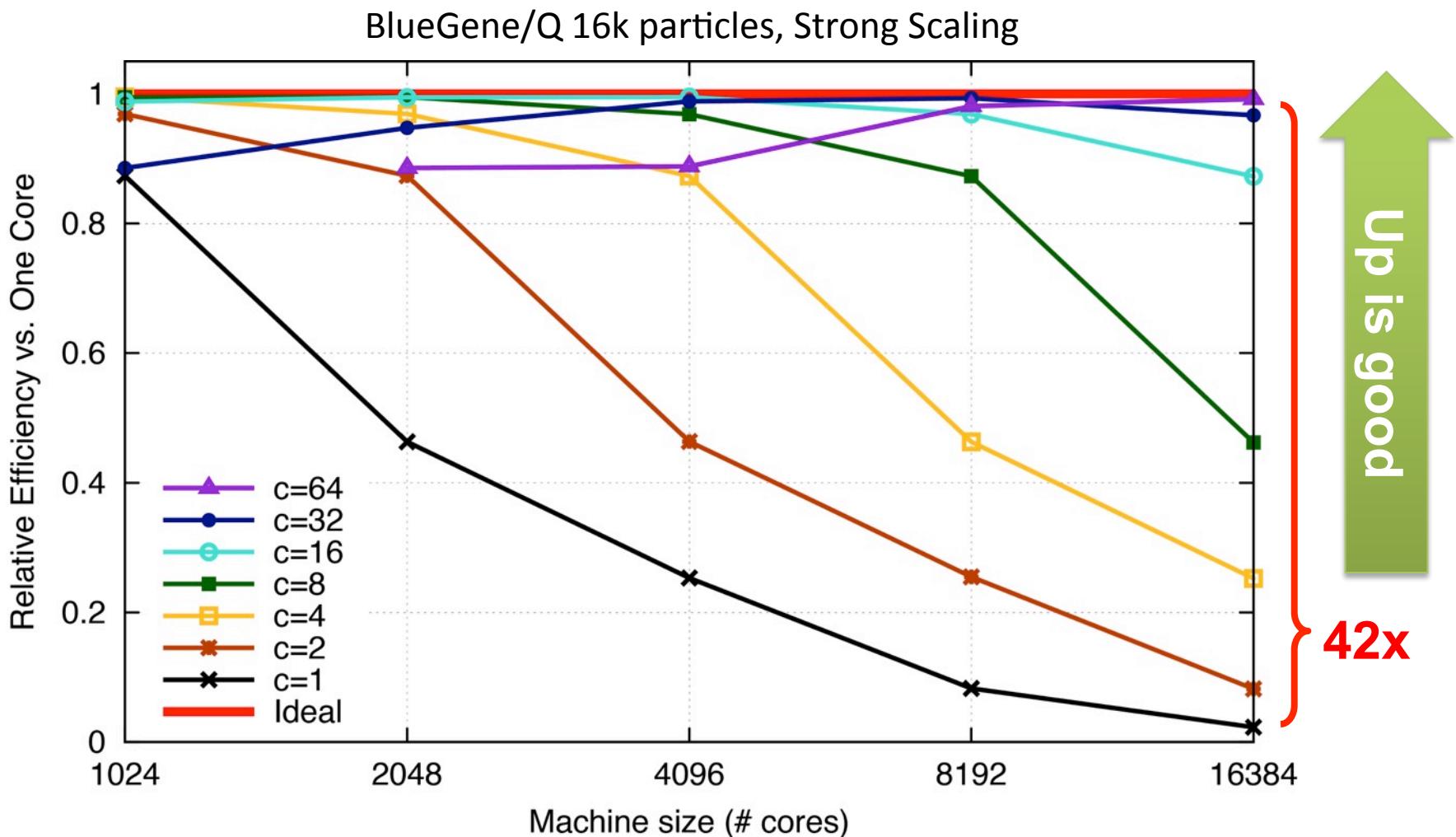
Communication optimal.  
Replication decreases  
#msgs and #words by  
factors of  $c^3$  and  $c^2$ .

# 3-Way N-Body Speedup

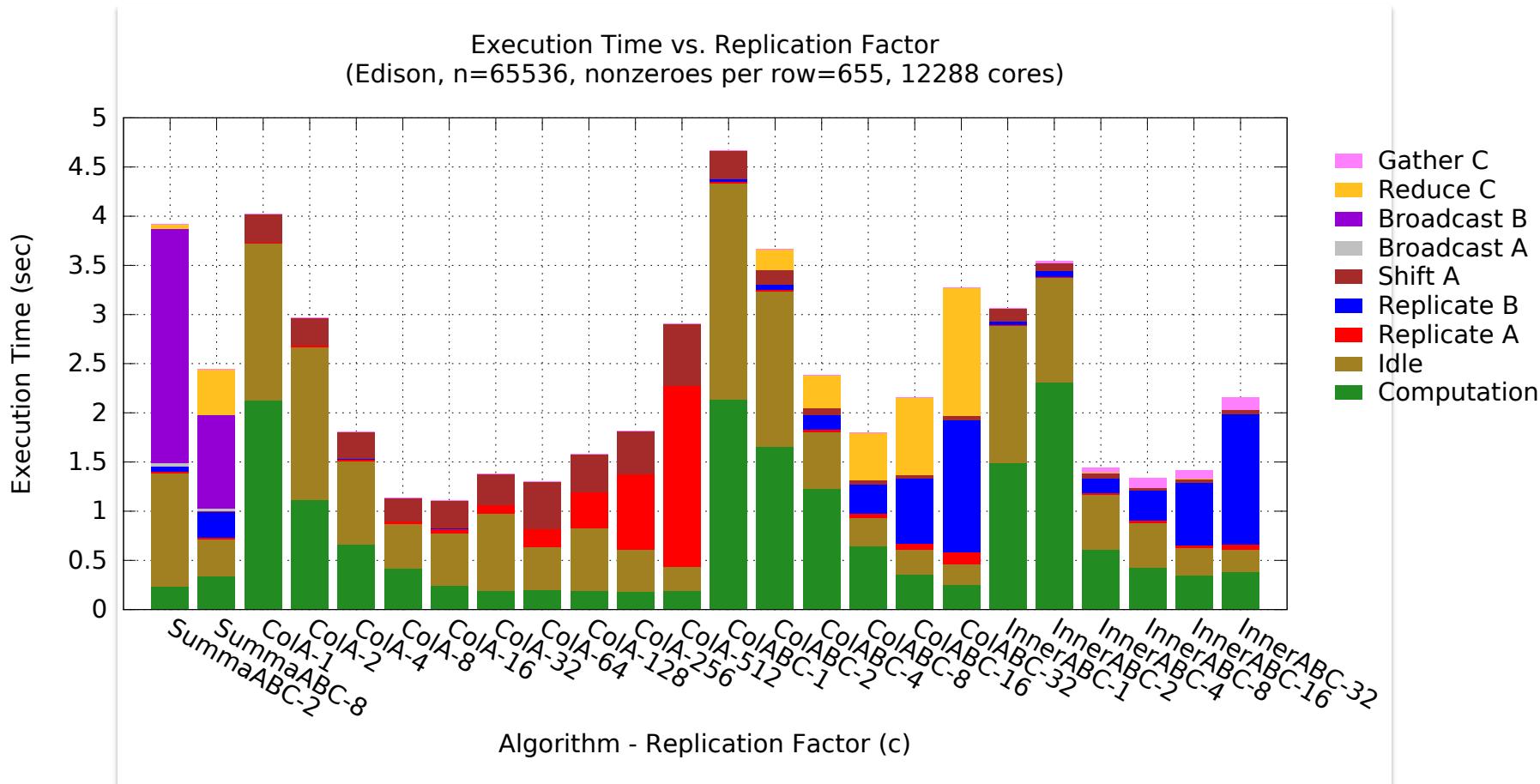
- Cray XC30, 24k cores, 24k particles



# Perfect Strong Scaling



# Sparse-Dense Matrix Multiply Too!

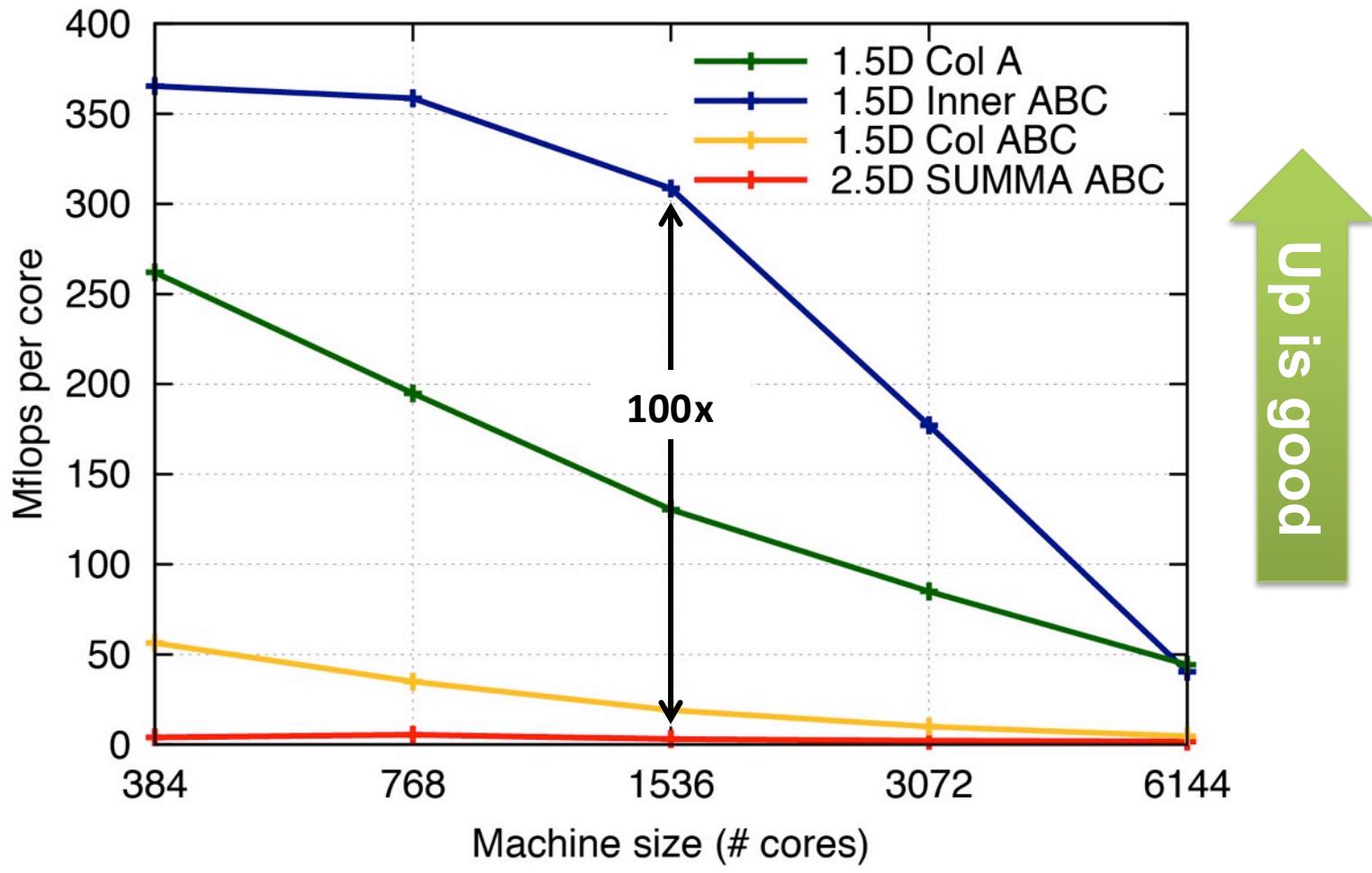


- Variety of algorithms that divide in 2 dimensions

Koanantakool et al

# 100x Improvement

- $A^{66k \times 172k}, B^{172k \times 66k}, 0.0038\% \text{ nnz}$ , Cray XC30



# Communication-Avoiding Algorithm Sample Speedups

- Up to 11.8x faster for direct N-body on 32K core IBM BG/P
- Up to 100x faster for sparse-dense matmul on Cray XC30
- Up to 12x faster for 2.5D matmul on 64K core IBM BG/P
- Up to 3x faster for tensor contractions on 2K core Cray XE/6
- Up to 6.2x faster for APSP on 24K core Cray CE6
- Up to 2.1x faster for 2.5D LU on 64K core IBM BG/P
- Up to 13x faster for TSQR on Tesla C2050 Fermi NVIDIA GPU
- Up to 6.7x faster for symeig (band A) on 10 core Intel Westmere
- Up to 2x faster for 2.5D Strassen on 38K core Cray XT4
- Up to 4.2x faster for MiniGMG benchmark bottom solver,  
using CA-BiCGStab (2.5x for overall solve)

Dense Linear algebra results by Demmel, Ballard, Solomonik, Grigori, et al

# Linear Algebra is important to Machine Learning too!

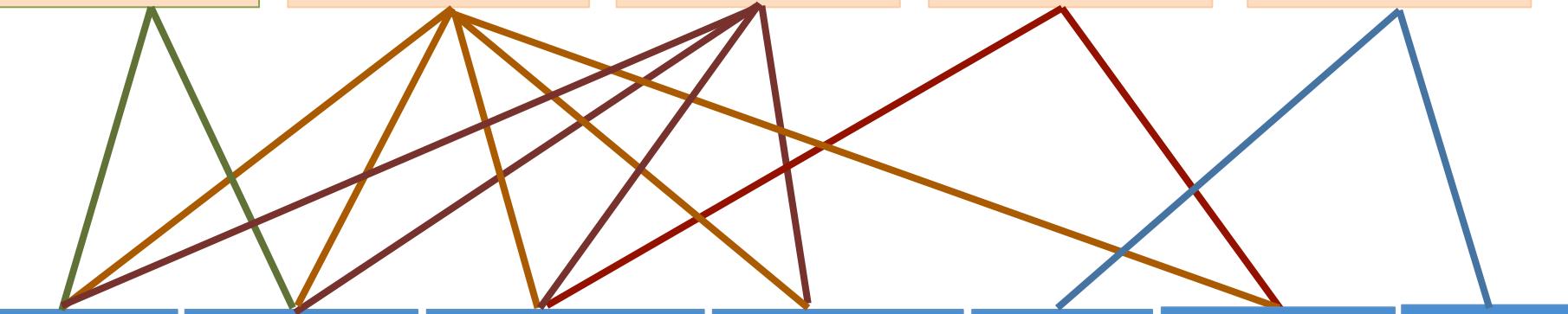
Logistic Regression,  
Support Vector  
Machines

Dimensionality  
Reduction (e.g.,  
NMF, CX/CUR,  
PCA)

Clustering  
(e.g., MCL,  
Spectral  
Clustering)

Graphical  
Model  
Structure  
Learning (e.g.,  
CONCORD)

Deep Learning  
(Convolutional  
Neural Nets)



Sparse  
Matrix-  
Sparse  
Vector  
(SpMSpV)

Sparse  
Matrix-  
Dense  
Vector  
(SpMV)

Sparse Matrix  
Times  
Multiple  
Dense Vectors  
(SpMM)

Sparse -  
Sparse  
Matrix  
Product  
(SpGEMM)

Dense  
Matrix  
Vector  
(BLAS2)

Sparse -  
Dense  
Matrix  
Product  
(SpDM<sup>3</sup>)

Dense  
Matrix  
Matrix  
(BLAS3)

Increasing arithmetic intensity

# **Overhead Can't be Tolerated**

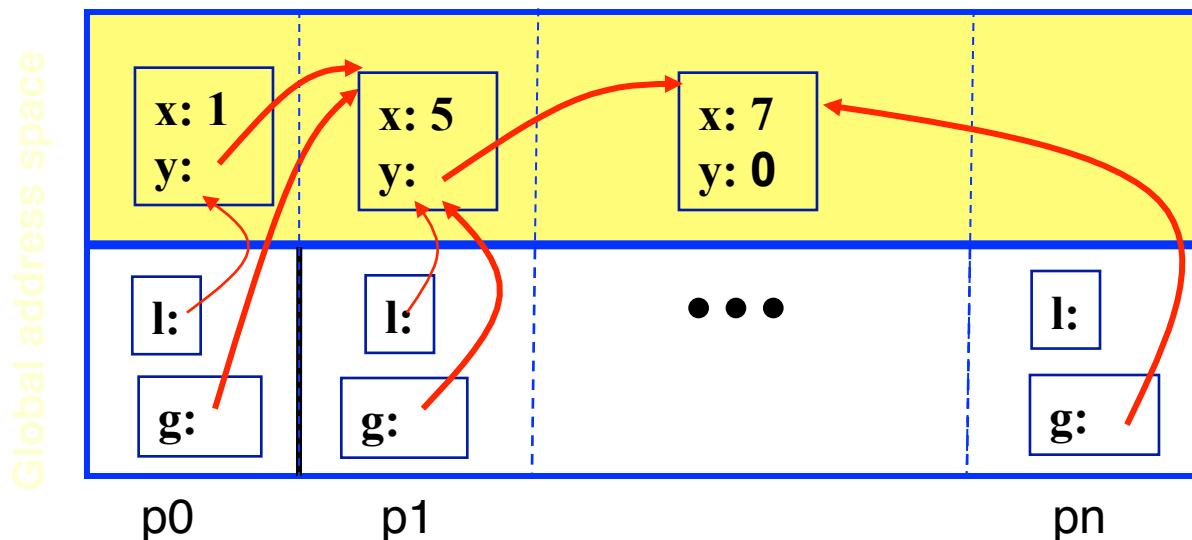
# PGAS: A programming model for exascale

- **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

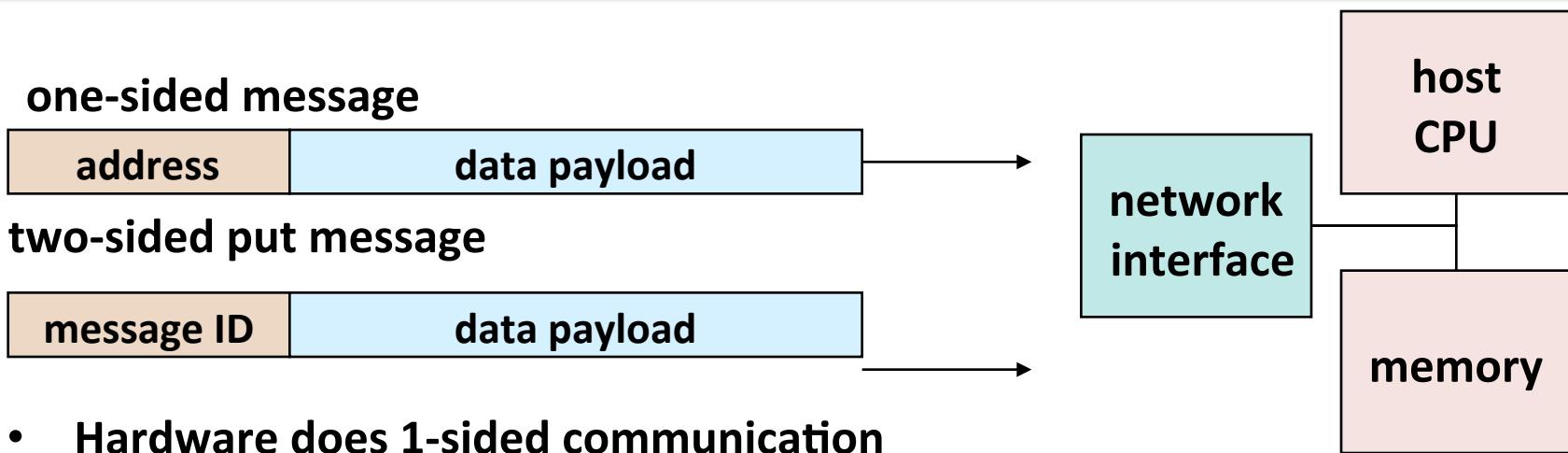
`shared int [ ] ga; and upc_malloc (...)`



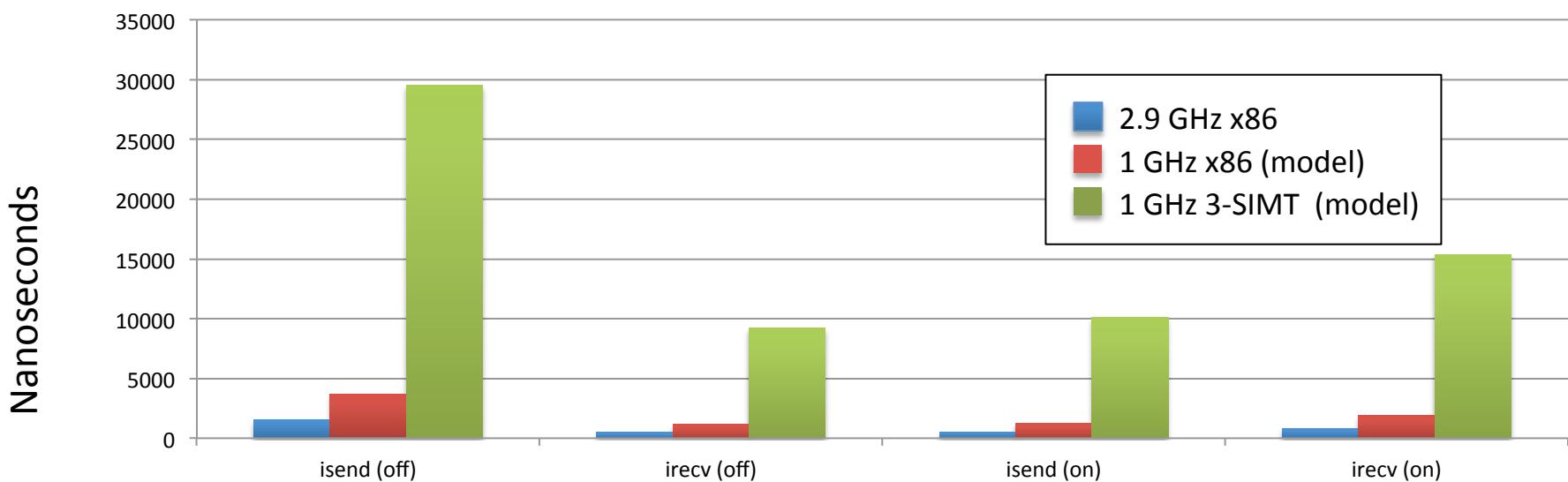
Examples:  
UPC  
UPC++

A programming model can influence how programmers think

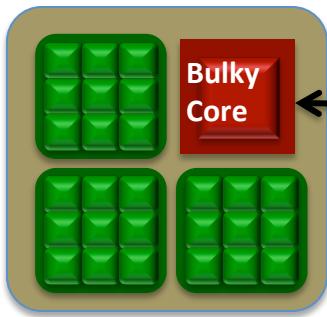
# One-Sided Communication is Closer to Hardware



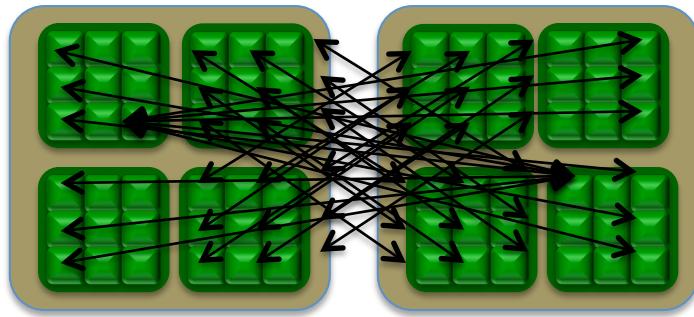
- **Hardware does 1-sided communication**
- **Overhead for send/receive messaging is worse at exascale**



# MPI+OpenMP: The Problem is the “+”

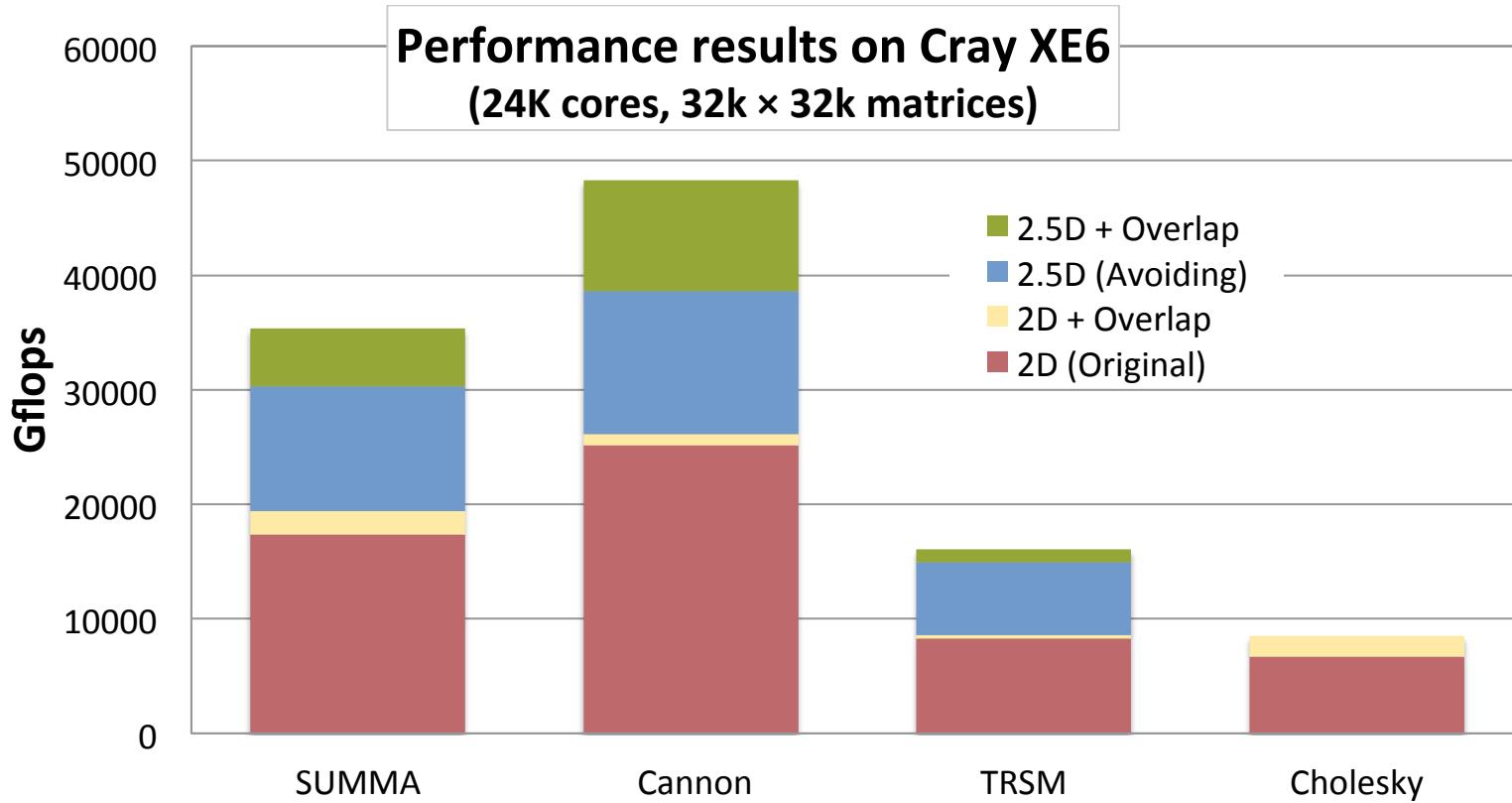


Funneled model does not scale



Addresses are better than tags

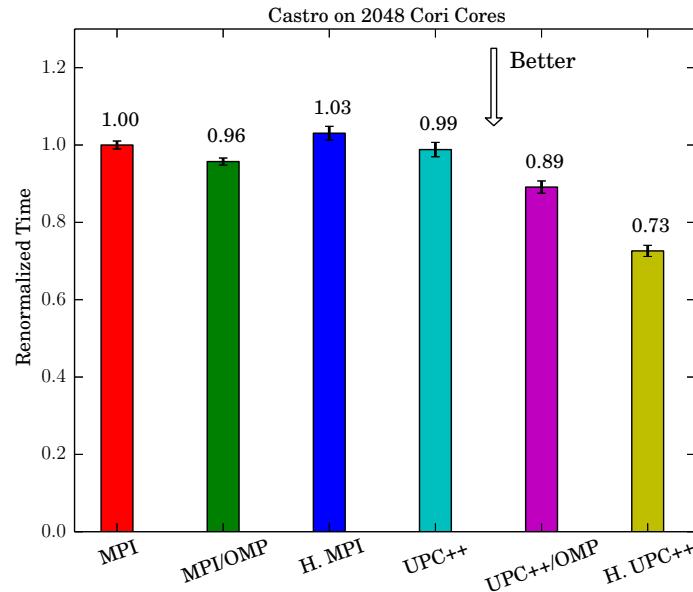
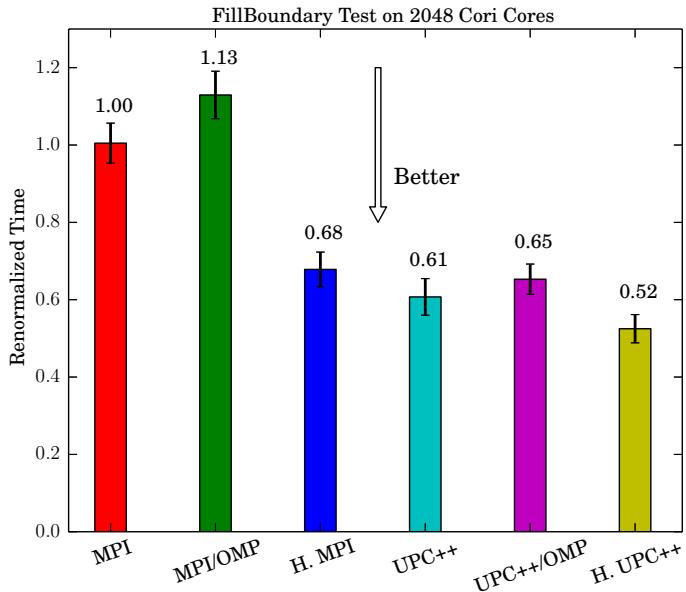
- **Avoid congestion at node interface:**
  - allow all cores to communicate
- **Coordinate on-node and between-node communication**
  - Performance is often lost between two software layers
- **Need small memory footprint per core**
  - Memory access, not heavyweight communication layer per core



**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)*

# 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)

# Avoid Unnecessary Synchronization

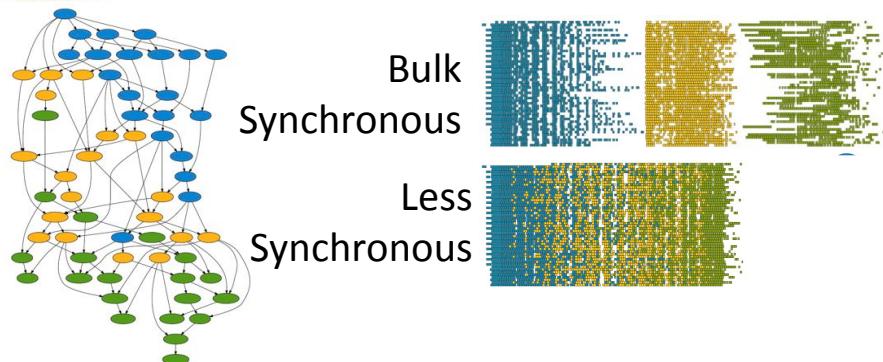
# Sources of Unnecessary Synchronization

## Loop Parallelism

```
!$OMP PARALLEL DO
DO I=2,N
  B(I) = (A(I) + A(I-1)) / 2.0
ENDDO
!$OMP END PARALLEL DO
```

“Simple” OpenMP parallelism implicitly synchronized between loops

## Abstraction



LAPACK: removing barriers ~2x faster (PLASMA)

## Libraries

Analysis	% barriers	Speedup
Auto	42%	13%
Guided	63%	14%

NWChem: most of barriers are unnecessary (Corvette)

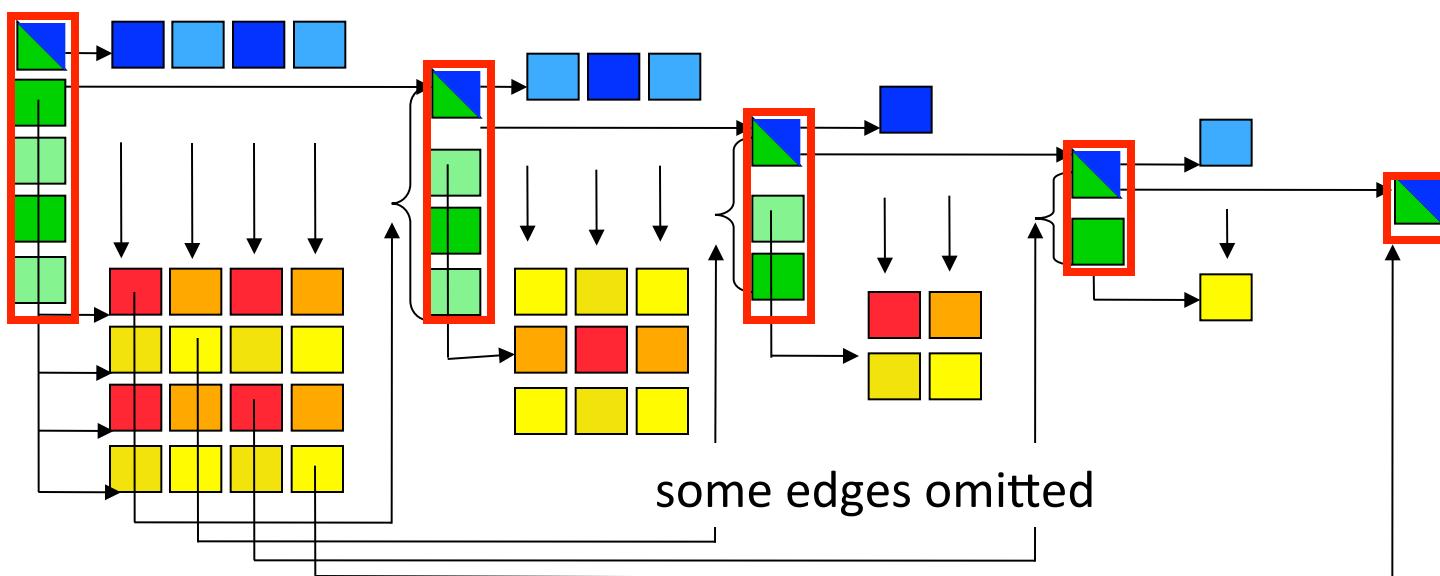
## Accelerator Offload

```
!$acc data copyin(cx1,c11,c12,c13,c14,r,b,uxyz,cell,rho,grad,index_max,index,&
!$acc& ciy,ciz,wet,np,streaming_sbuf1, &
!$acc& streaming_sbuf1,streaming_sbuf2,streaming_sbuf4,streaming_sbuf5,&
!$acc& streaming_sbuf7s,streaming_sbuf8s,streaming_sbuf9s,streaming_sbuf10s,&
!$acc& streaming_sbuf11n,streaming_sbuf12n,streaming_sbuf13s,streaming_sbuf14n,&
!$acc& streaming_sbuf7e,streaming_sbuf8w,streaming_sbuf9e,streaming_sbuf10e,&
!$acc& streaming_sbuf11w,streaming_sbuf12e,streaming_sbuf13w,streaming_sbuf14w,&
!$acc& streaming_rbuf1,streaming_rbuf2,streaming_rbuf4,streaming_rbuf5,&
!$acc& streaming_rbuf7n,streaming_rbuf8n,streaming_rbuf9s,streaming_rbuf10n,&
!$acc& streaming_rbuf11s,streaming_rbuf12s,streaming_rbuf13n,streaming_rbuf14s,&
!$acc& streaming_rbuf7w,streaming_rbuf8e,streaming_rbuf9w,streaming_rbuf10w,&
!$acc& streaming_rbuf11e,streaming_rbuf12w,streaming_rbuf13e,streaming_rbuf14e, &
!$acc& send_e,send_w,send_n,send_s,recv_e,recv_w,recv_n,recv_s)
```

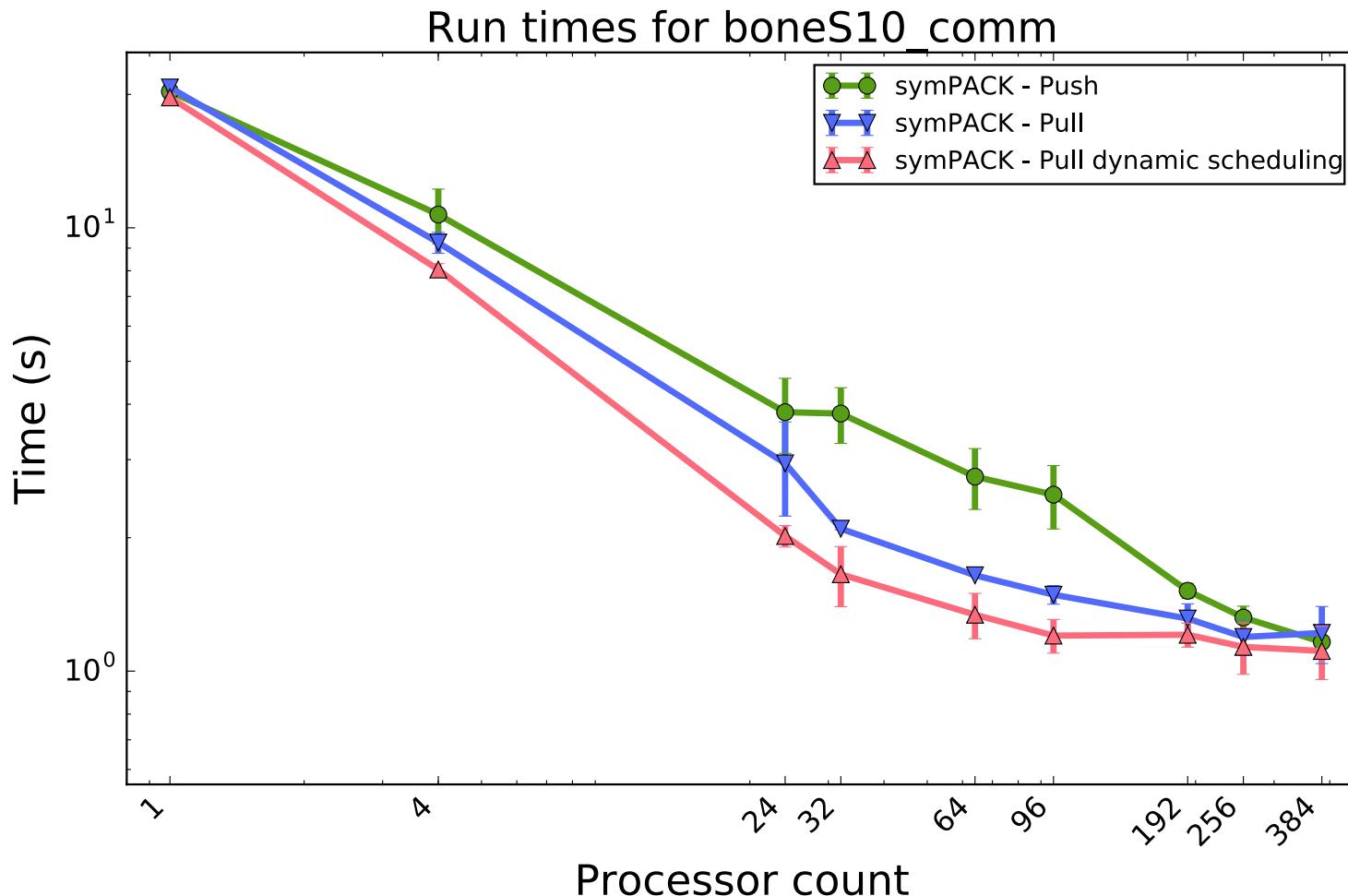
The transfer between host and GPU can be slow and cumbersome, and may (if not careful) get synchronized

# Event Driven LU in UPC

- Assignment of work is static; schedule is dynamic
- Ordering needs to be imposed on the schedule
  - Critical path operation: Panel Factorization
- General issue: dynamic scheduling in partitioned memory
  - Can deadlock in memory allocation
  - “memory constrained” lookahead



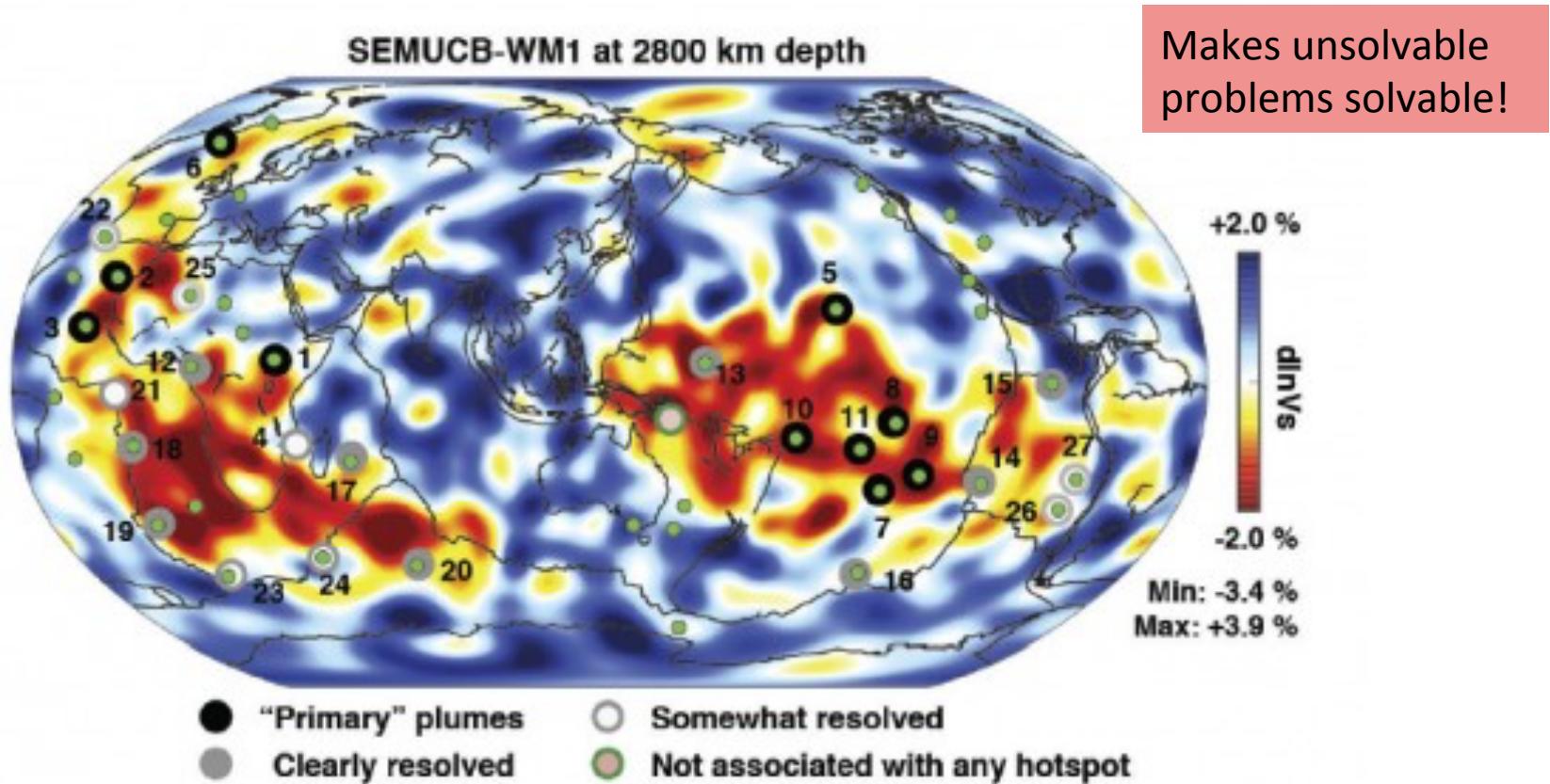
# Asynchronous Sparse Cholesky in UPC++



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

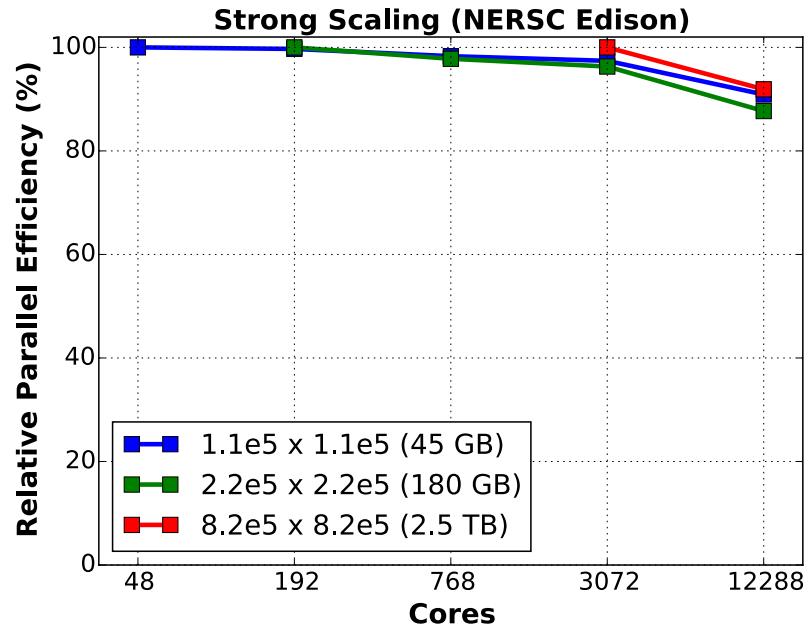
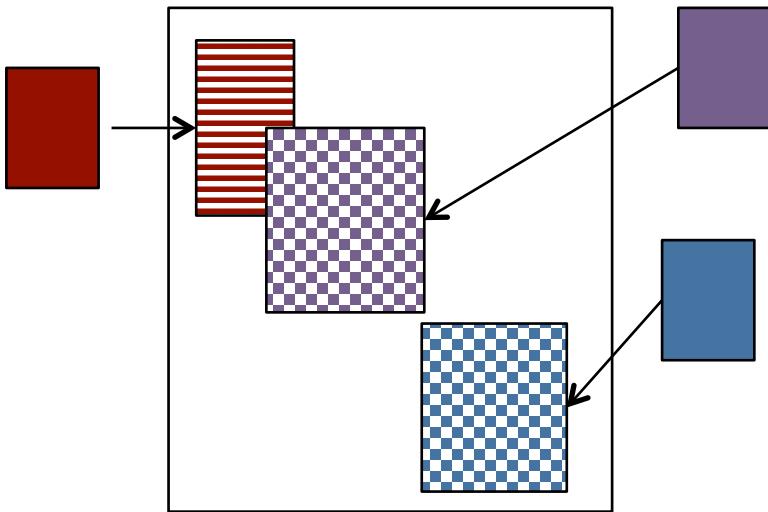
# Whole-Mantle Seismic Model Using UPC++

- 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

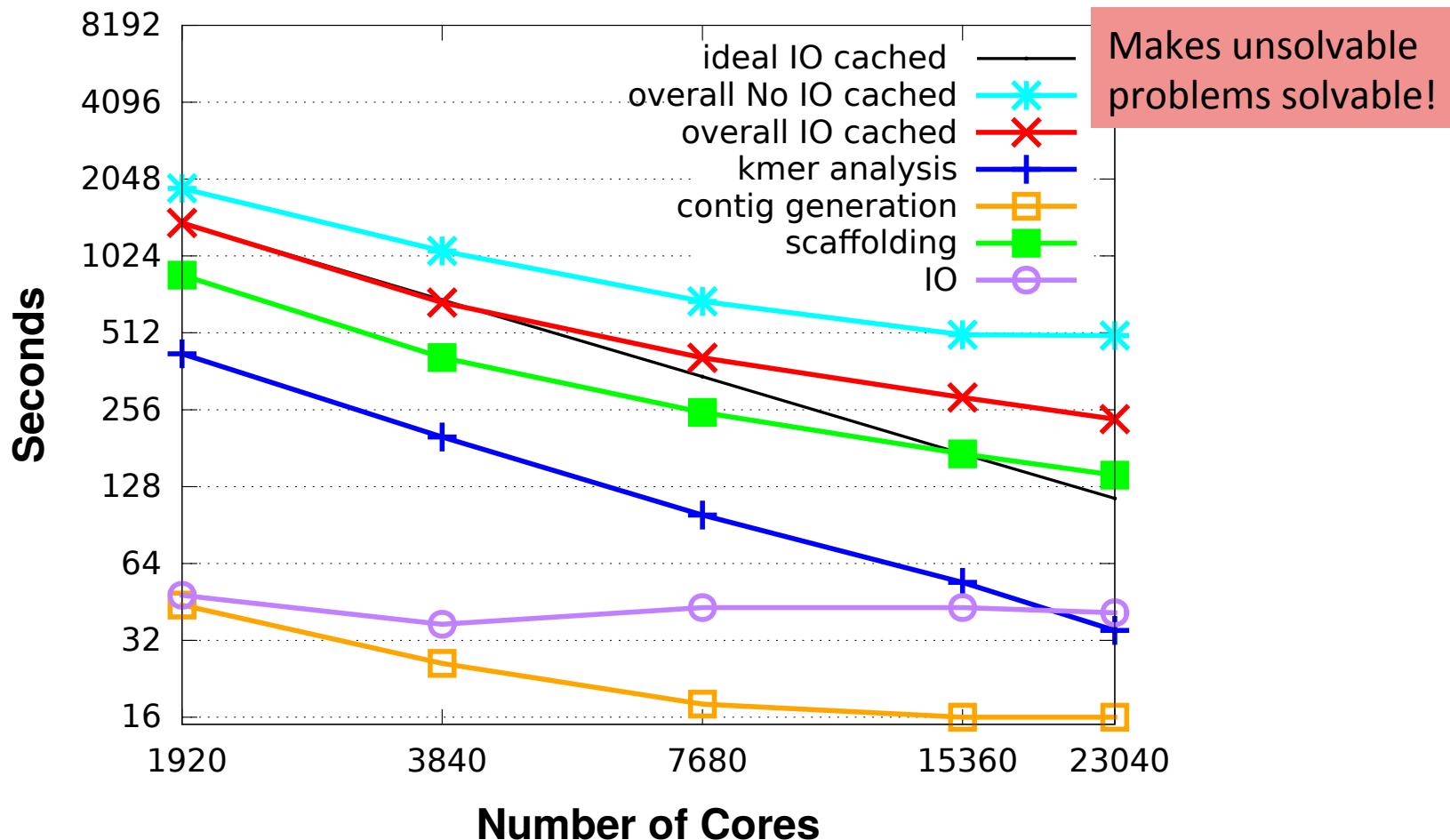
Scott French, Y. Zheng, B. Romanowicz, K. Yelick

# *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



# 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)

# Summary

- **Communication is the most expensive thing computers do**
  - Memory
  - Network
- **Compilers**
  - Domain specific languages simplify program analysis
  - Autotuning helps identify optimizations
- **PGAS and one-sided communication can help lower costs**
  - Better match to RDMA hardware
  - Also supports applications with irregular accesses
- **Algorithms can avoid communication**
  - Both data volume (bandwidth) and number of messages (latency)
  - Probably optimal and faster (more scalable) in practice



# Thank you!



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