



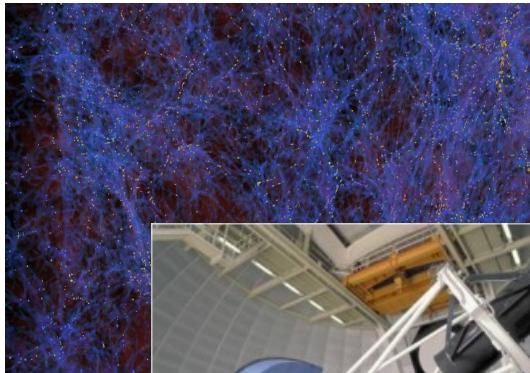
# Avoiding, Hiding and Managing Communication at the Exascale

Kathy Yelick

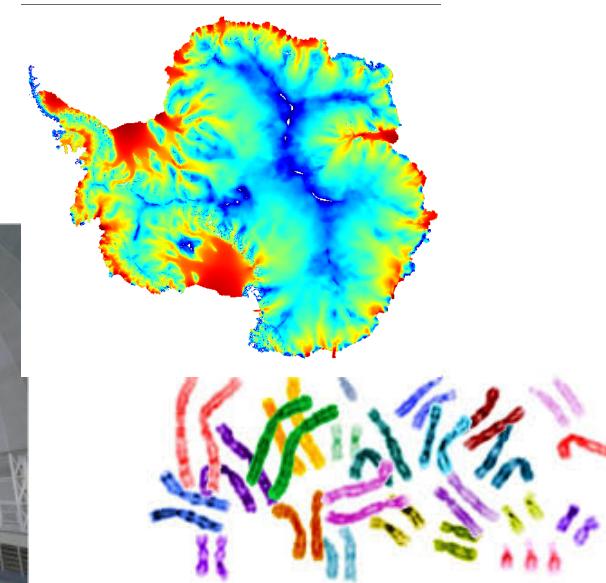
Associate Laboratory Director  
Computing Sciences  
Lawrence Berkeley National Lab

Professor of Electrical Engineering  
and Computer Sciences  
UC Berkeley

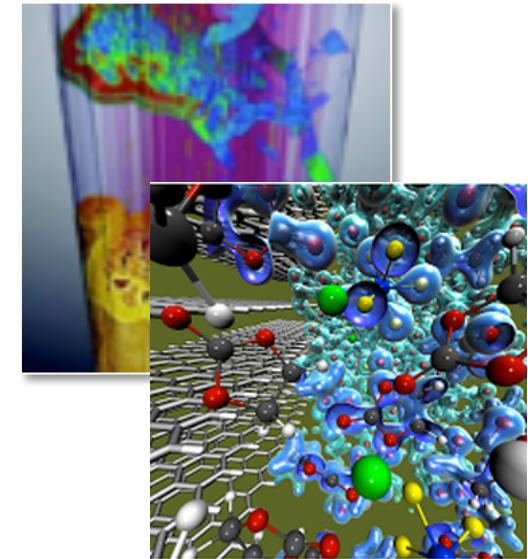
# Science at the Boundary of Simulation and Observation



*Cosmology*



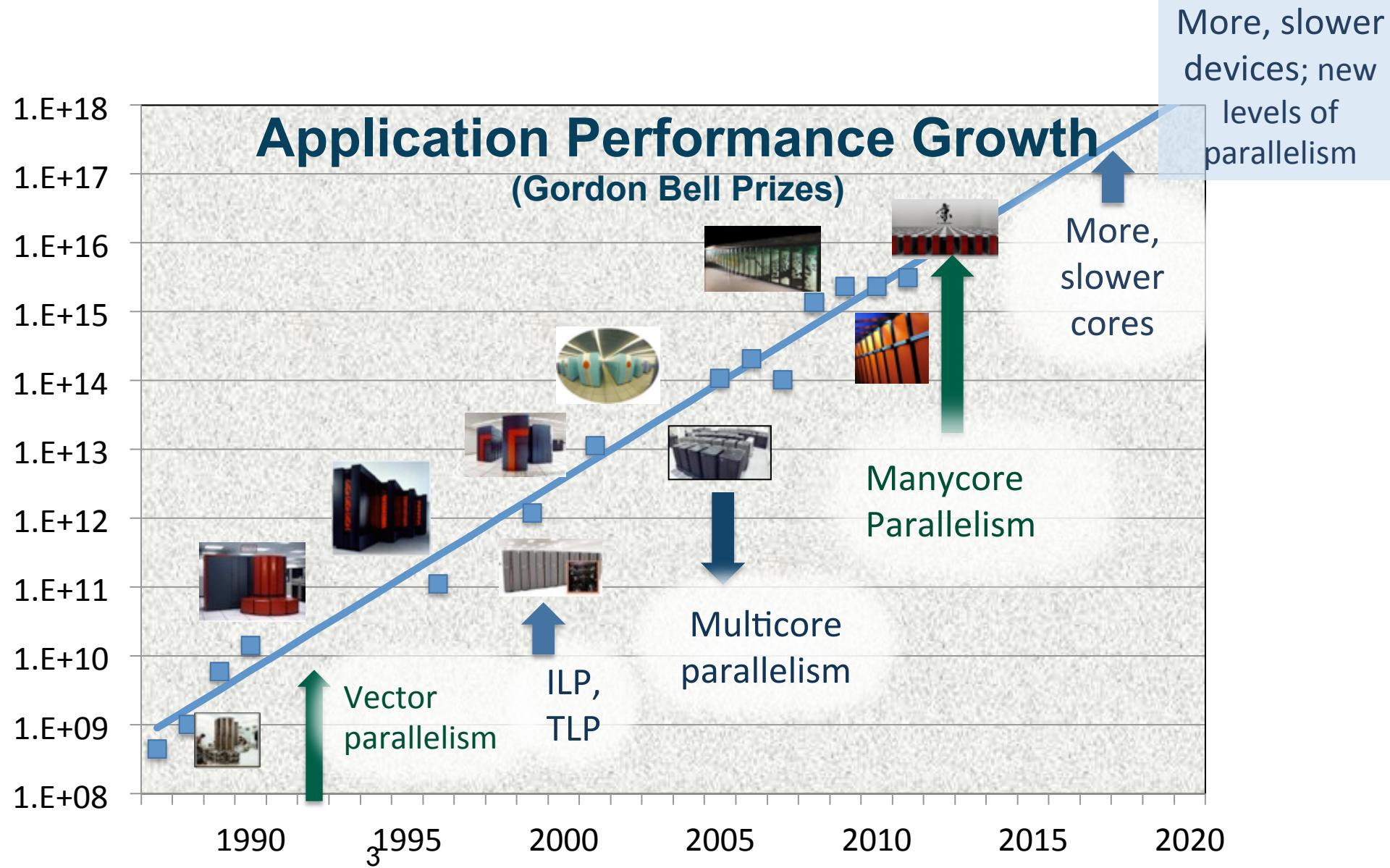
*Environment*



*Materials*

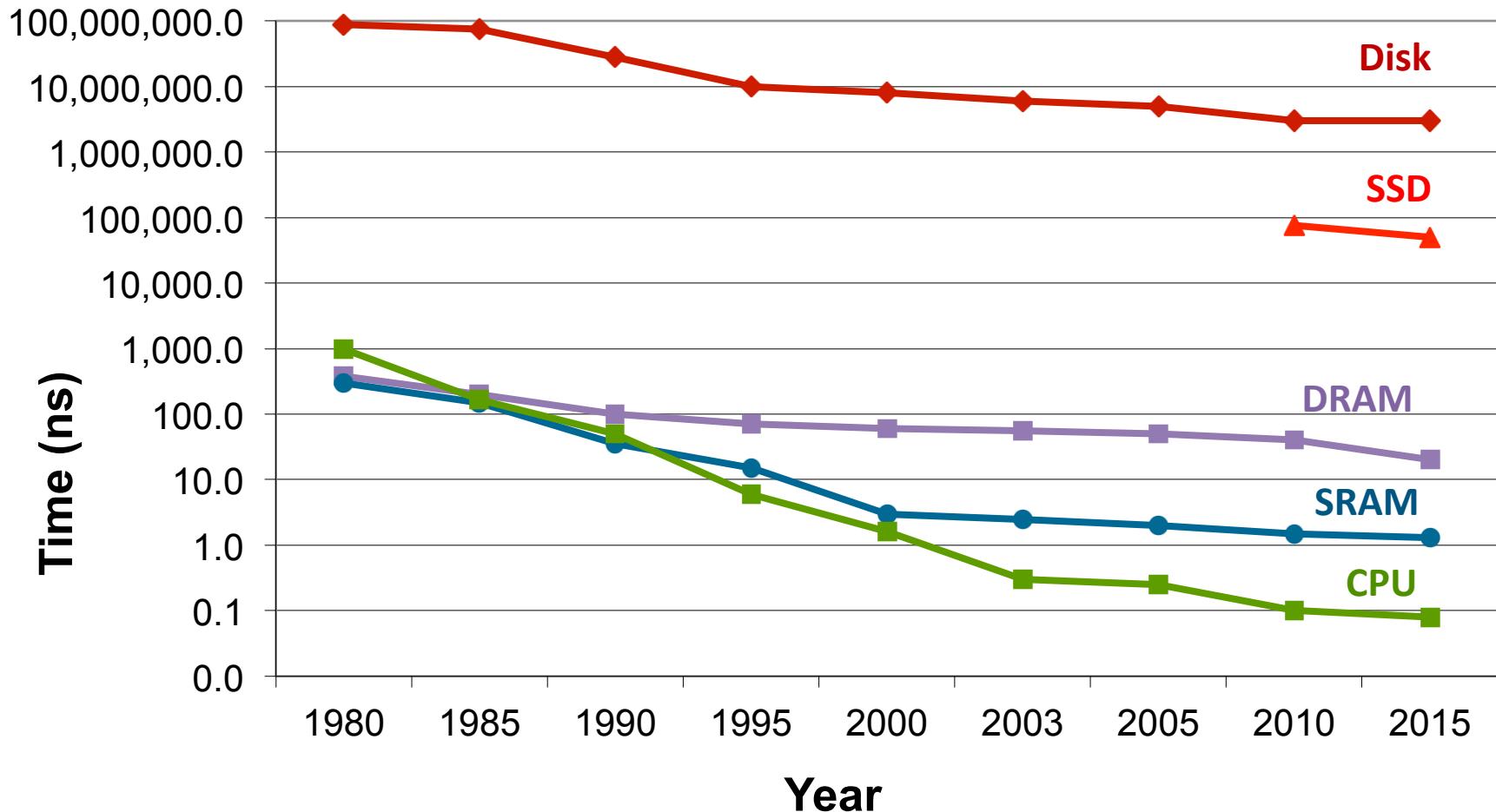
In many areas, there are opportunities to combine simulation and observation for new discoveries. These will require increased computing capabilities.

# Hardware Trend: More Parallelism at Lower Levels



# Data Movement is Expensive

## CPU cycle time vs memory access time



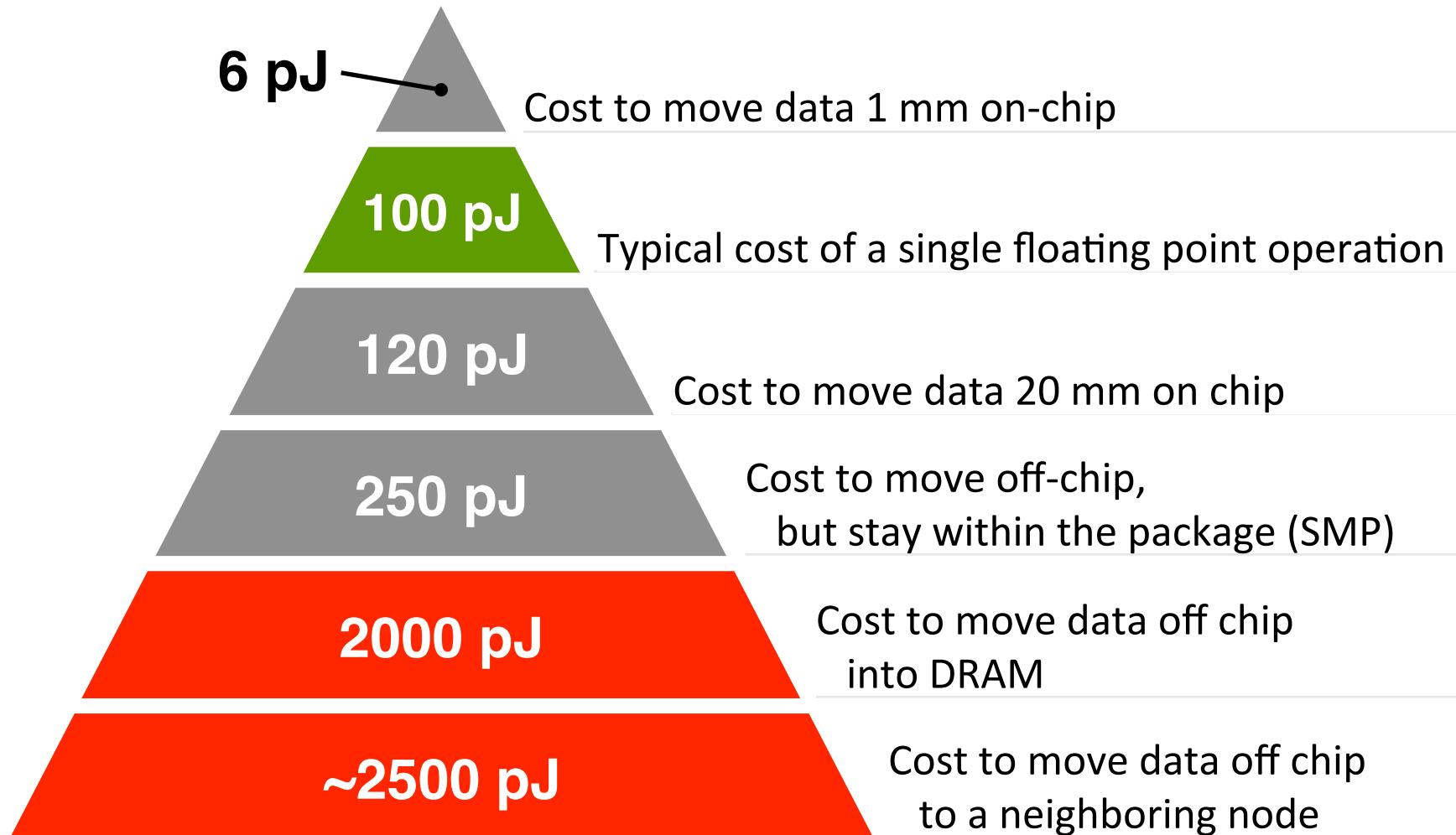
Sources:

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

Communication Avoidance for Algorithms with Sparse All-to-all Interactions

# Data Movement is Expensive

## Hierarchical energy costs.



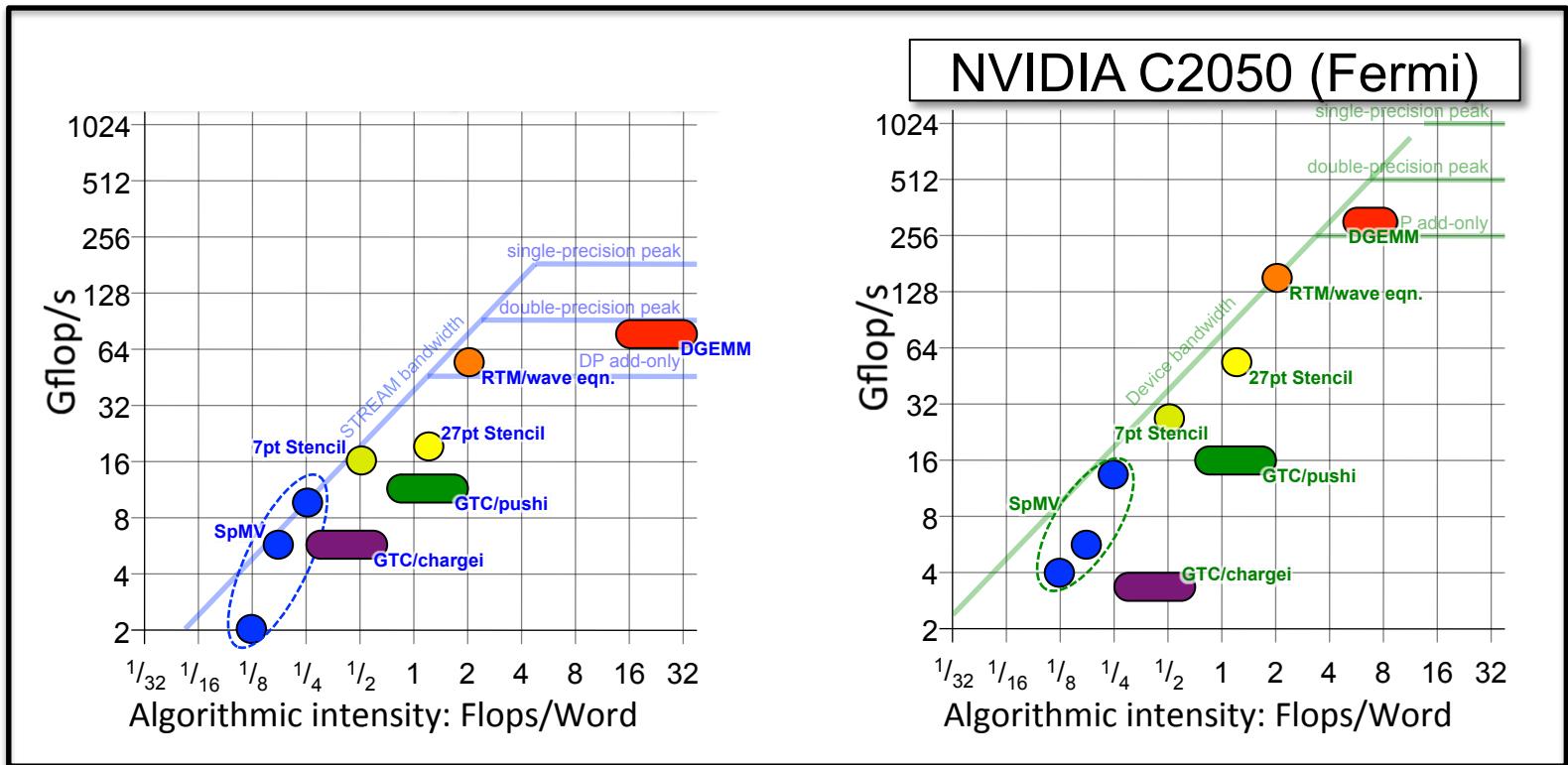
# Summary

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

# **Compilers and Autotuning**

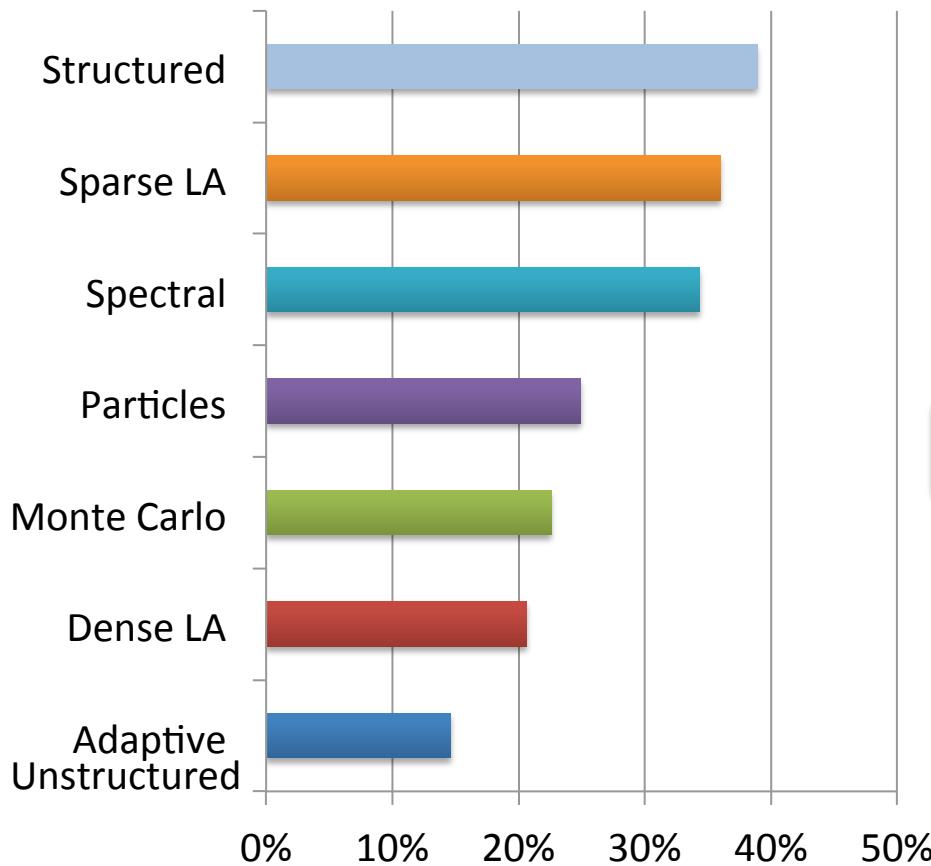
# Autotuning: Write Code Generators

- 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?



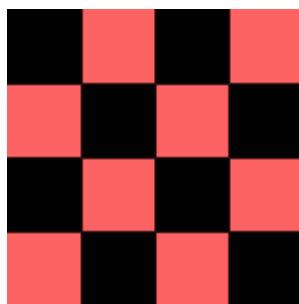
What code generators do we have?

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

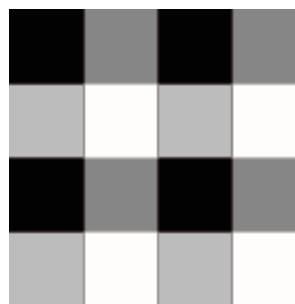
Stencils are both the most important motifs and a gap in our tools

# Approach: Small Compiler for Small Language

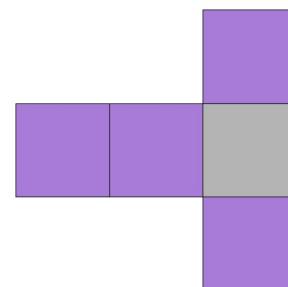
- **Snowflake: A DSL for Science Stencils**
  - Domain calculus inspired by Titanium, UPC++, and AMR in general



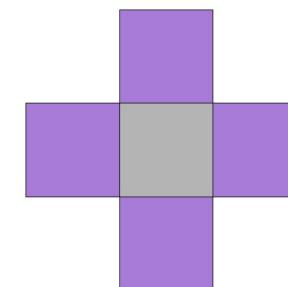
(a) Red-Black tiling



(b) 4-color tiling



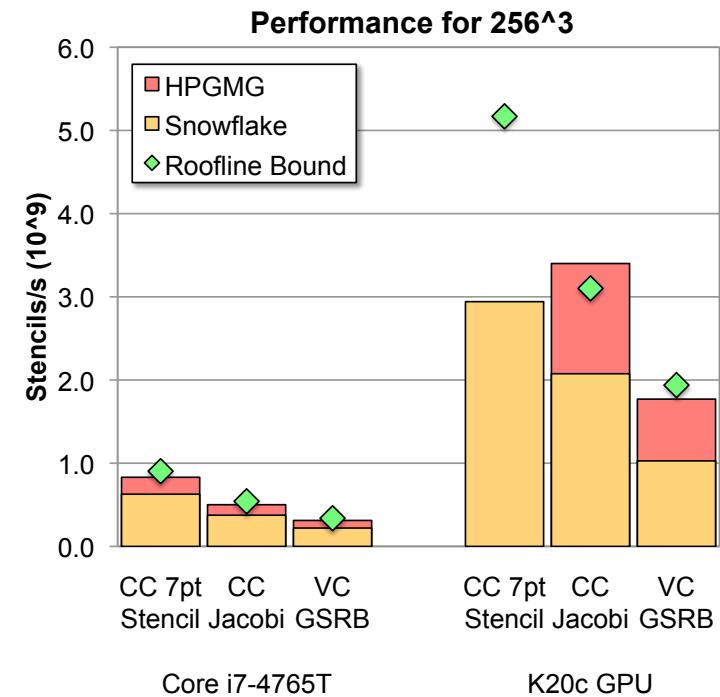
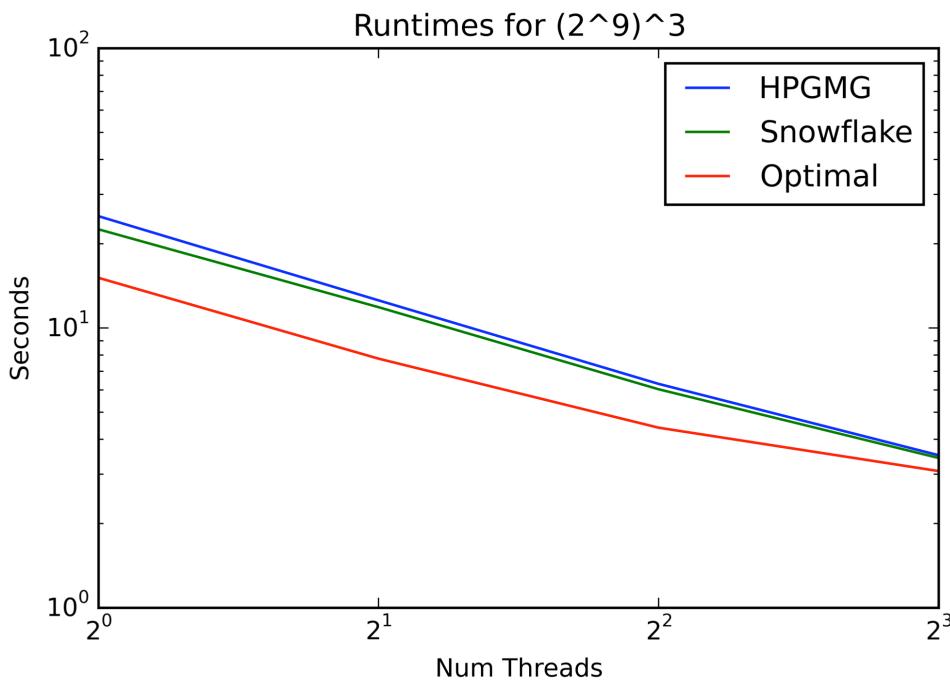
(c) Asymmetric stencil used  
near mesh boundary



(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)

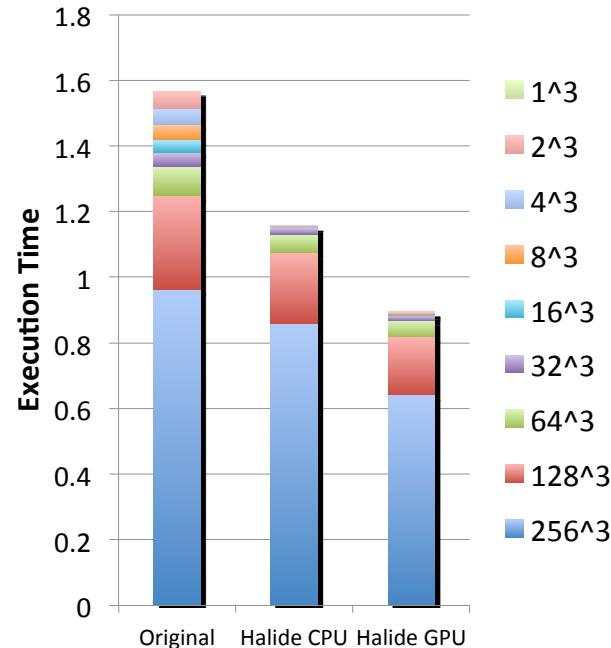
- **Halide Algorithm by domain expert**

```
Func Ax_n("Ax_n", lambda("lambda"), chebyshev("chebyshev"));
Var i("i^3"), j("j^3"), k("k^3");
Ax_n(i,j,k) = lambda(i,j,k)*x_n(i,j,k) - b^2*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) * valid(i-1,j,k) - 2.0f)  
    + beta_i(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

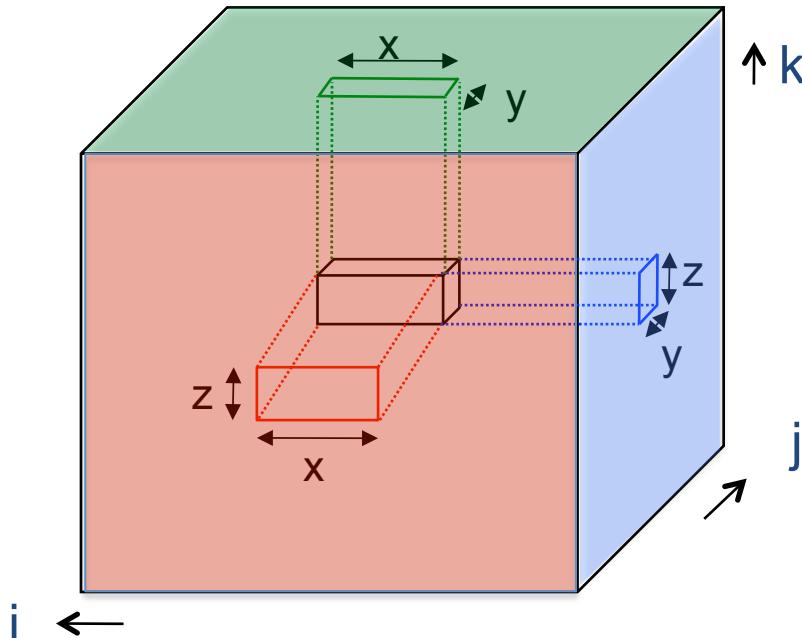
**Lesson: Never waste fast memory**

**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])
```

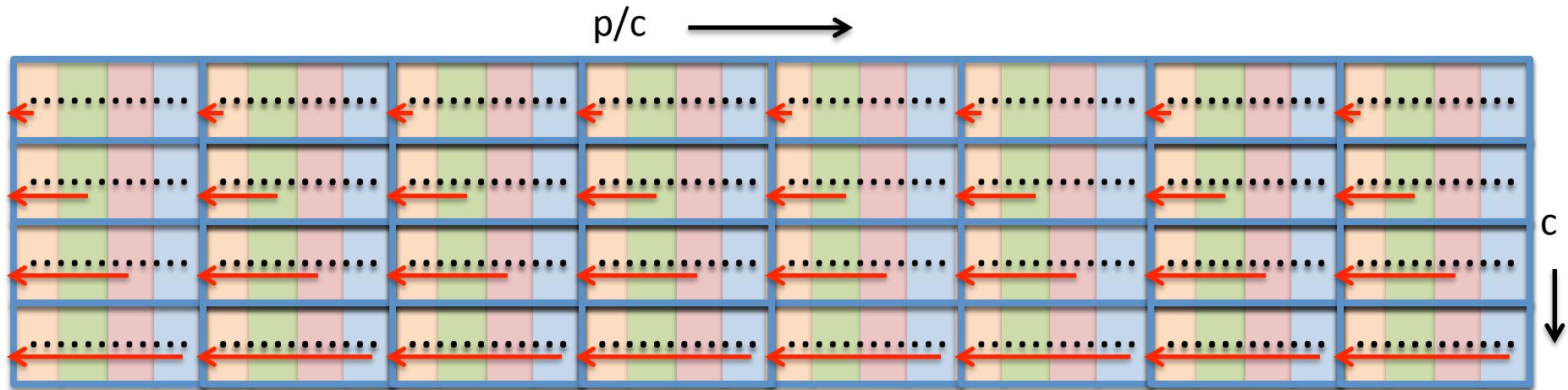
$O(n^k)$ .

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



No!

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

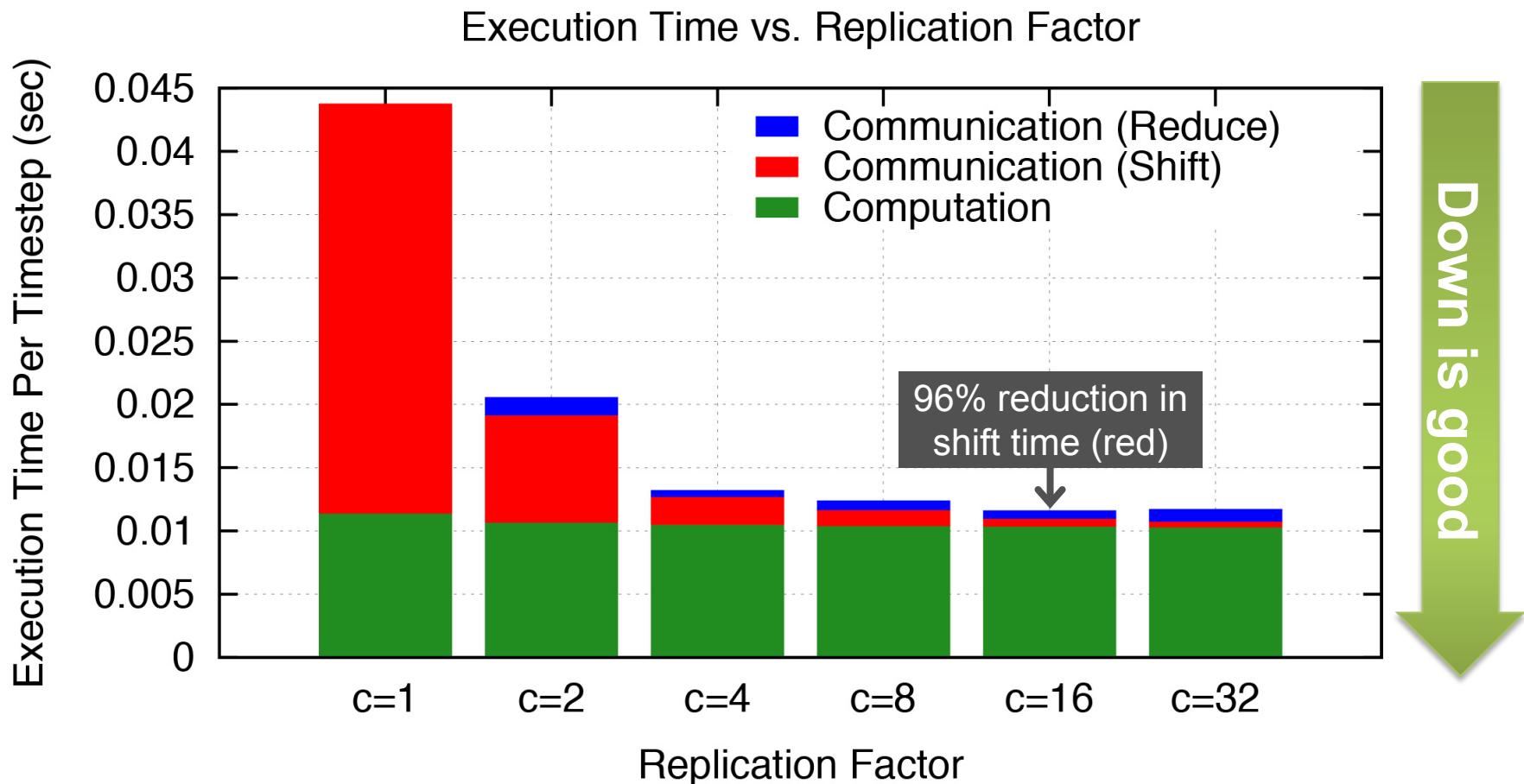


- Divide  $p$  into  $c$  groups
- Replicate particles across groups
- Repeat: shift copy of  $n/(p*c)$  particles to the left within a group
- Reduce across  $c$  to produce final value for each particle

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

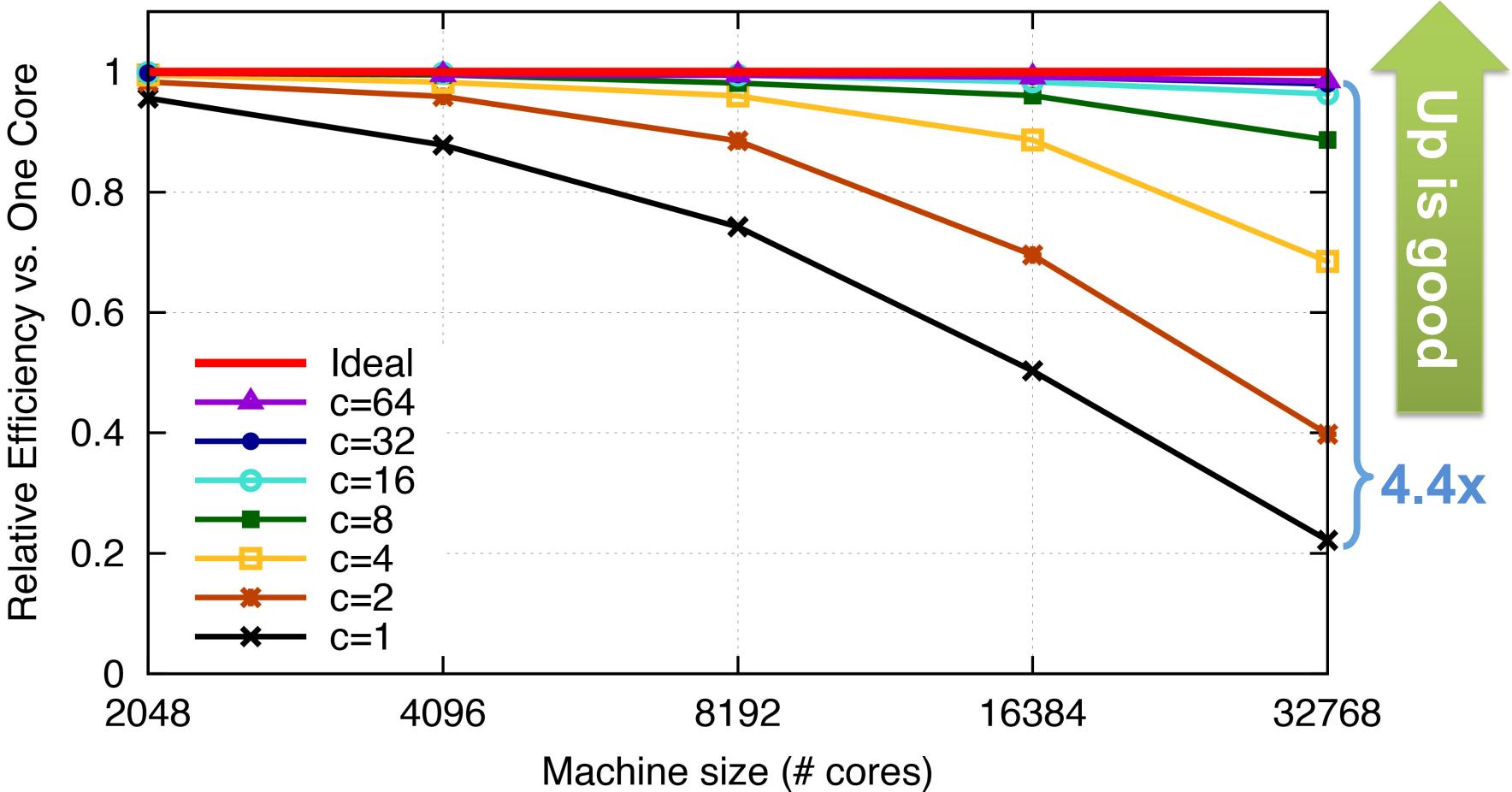
# Less Communication..

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



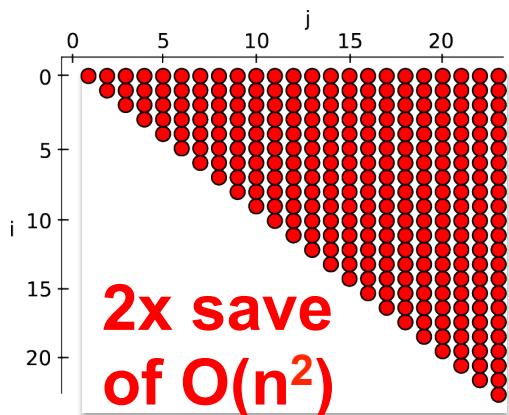
# Strong Scaling

Parallel Efficiency on BlueGene/P (n=262,144)

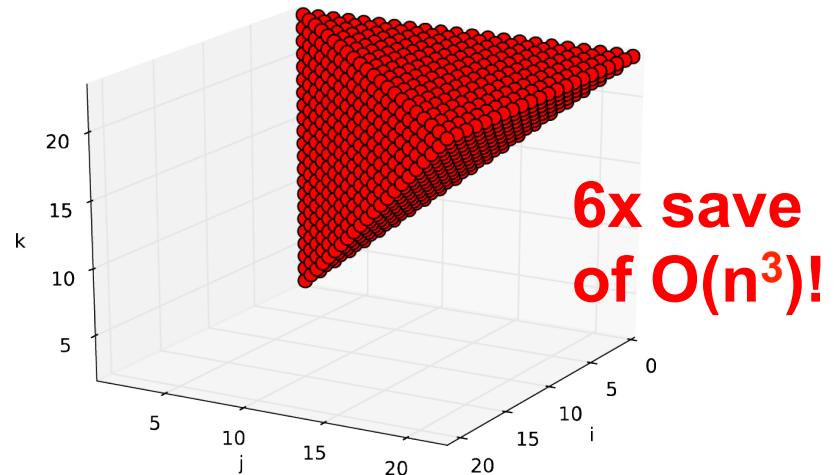


# Challenge: Symmetry & Load Balance

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



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

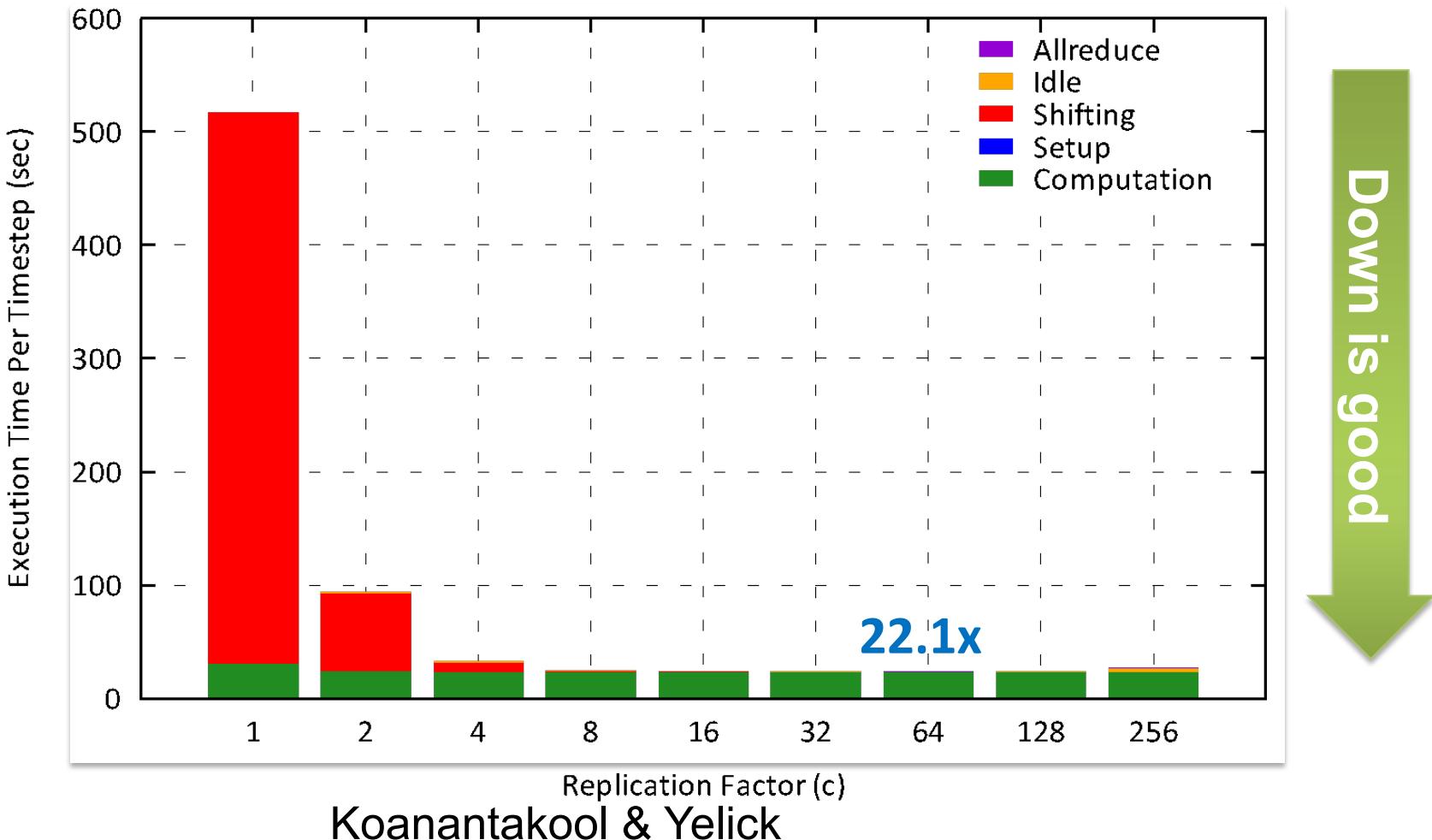


6x save  
of  $O(n^3)$ !

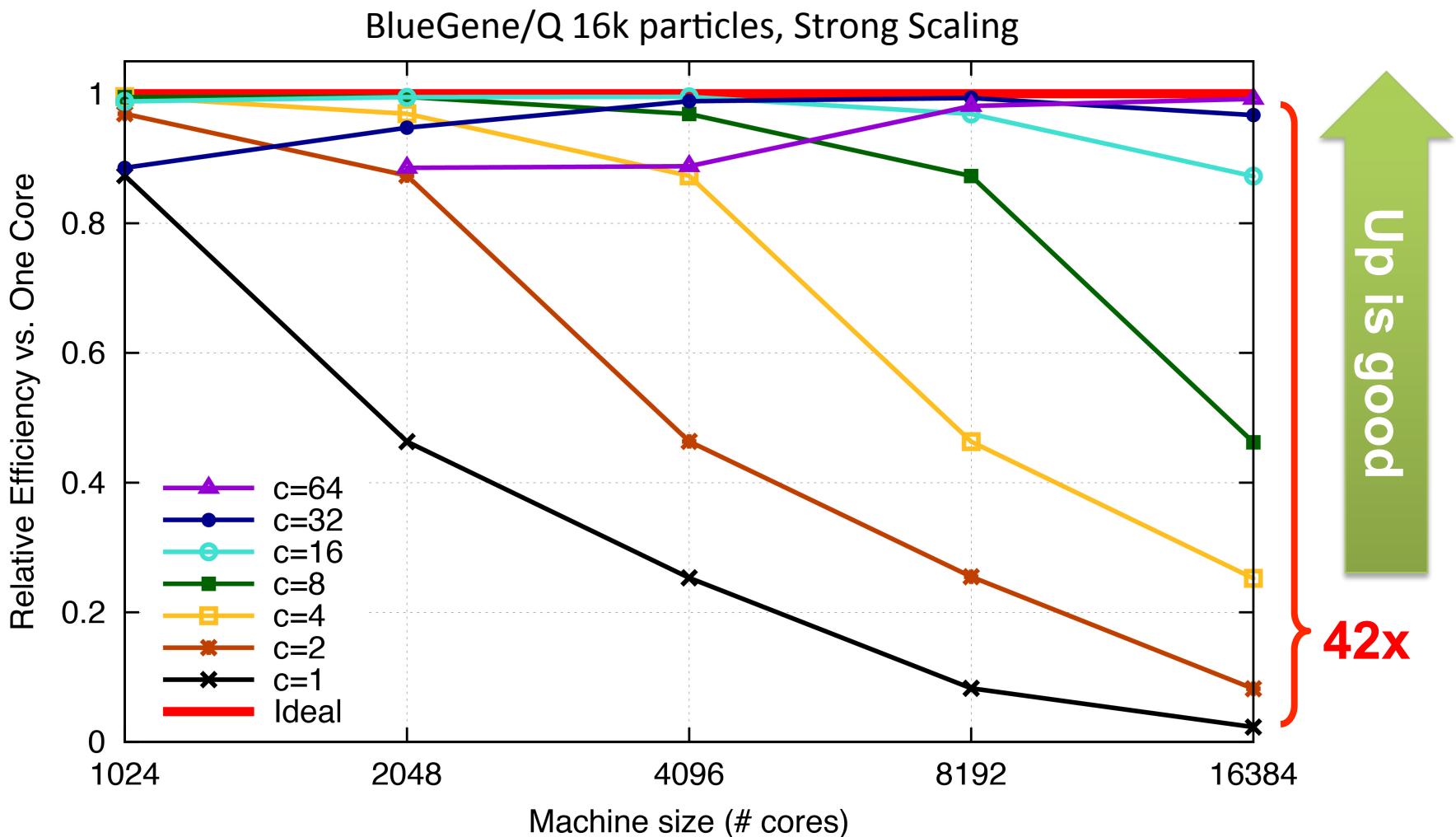
- How to divide work equally?

# 3-Way N-Body Speedup

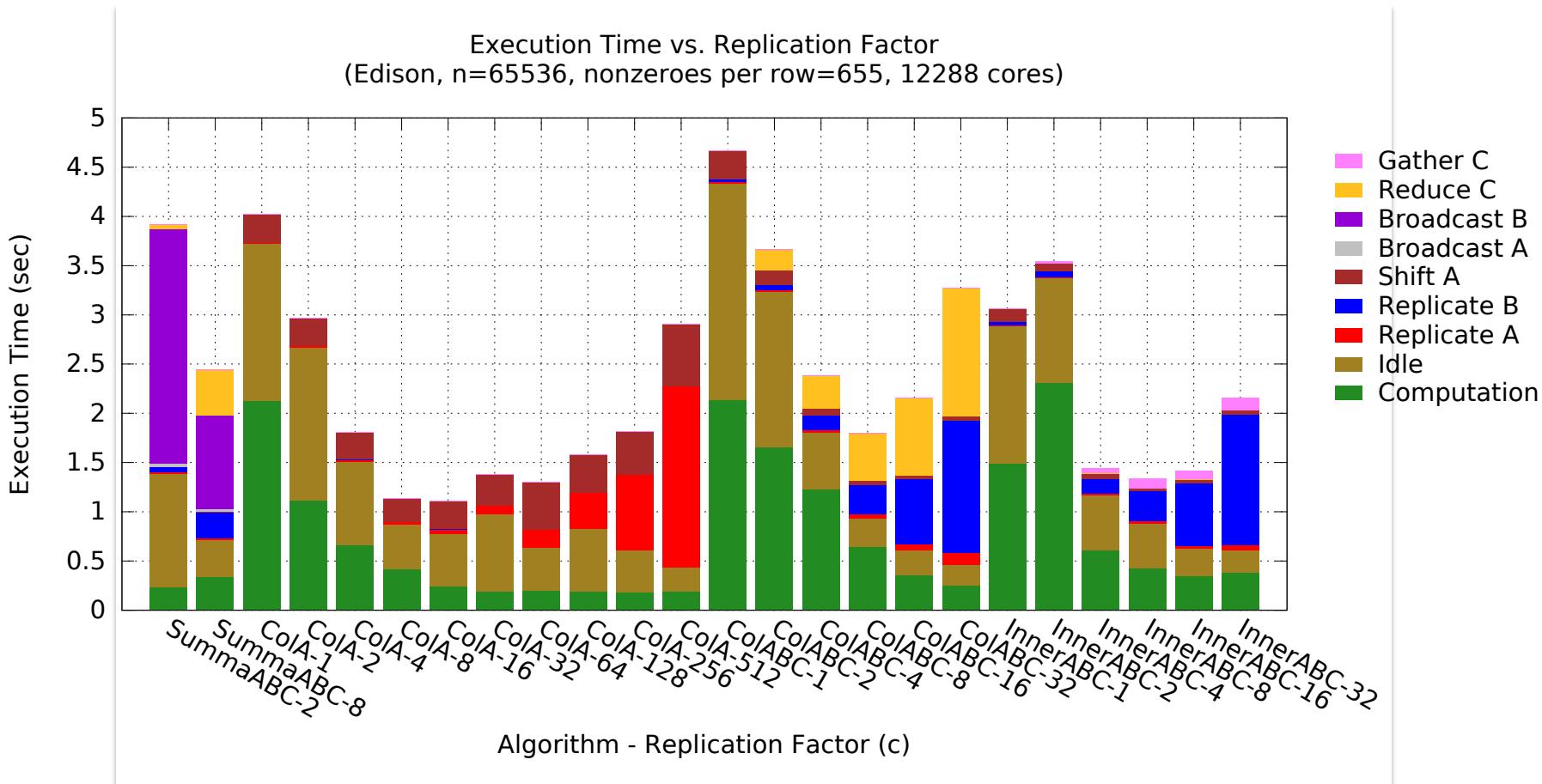
- Cray XC30, 24k cores, 24k particles



# Perfect Strong Scaling



# Sparse-Dense Matrix Multiply Too!

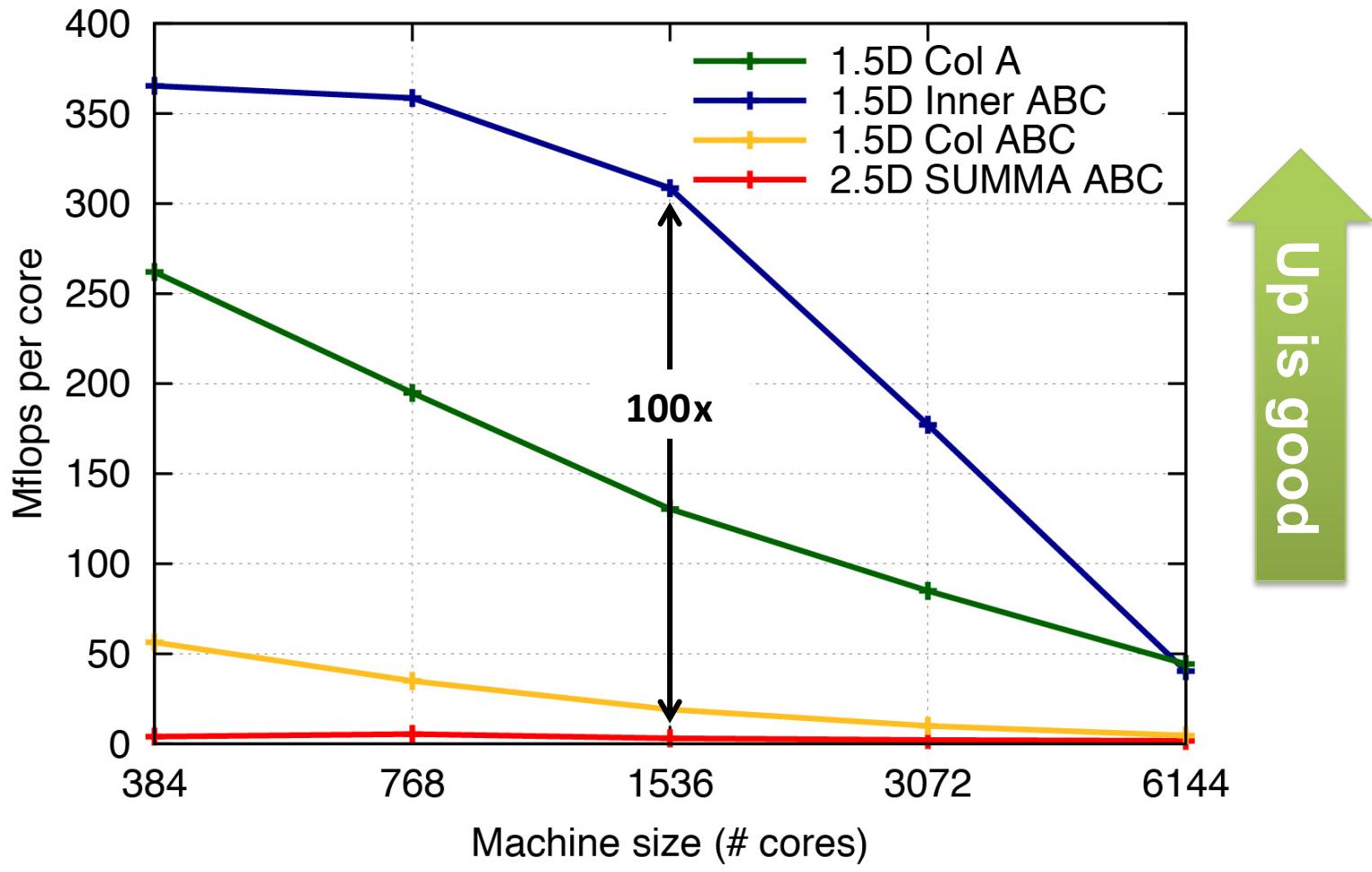


- Variety of algorithms that divide in or 2 dimensions

Koanantakool & Yelick

# 100x Improvement

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



# Linear Algebra is important to Machine Learning too!

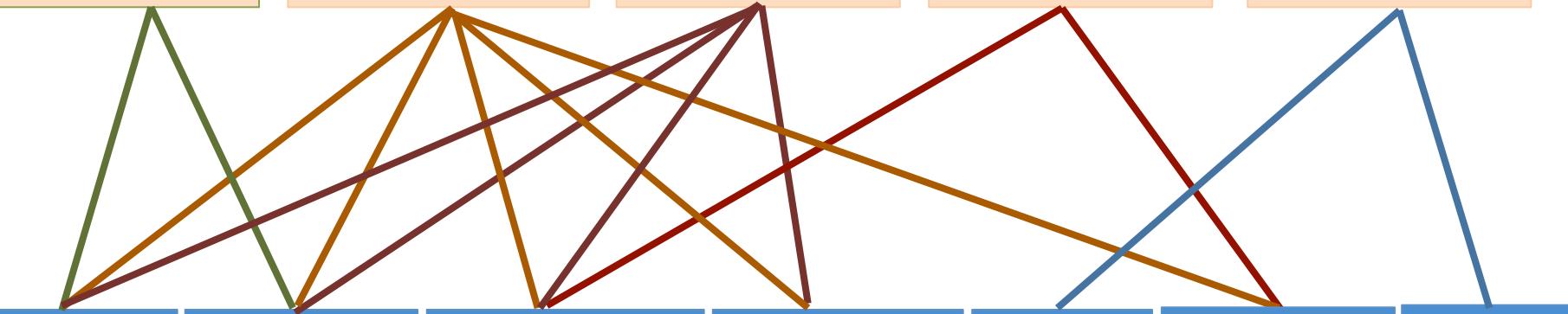
Logistic Regression,  
Support Vector  
Machines

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**

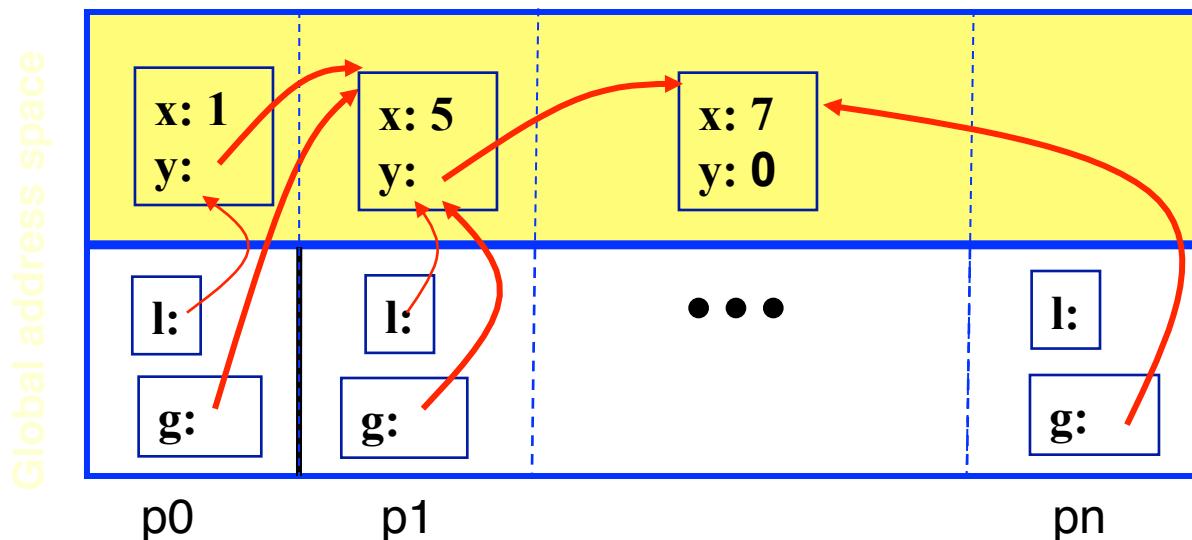
# PGAS: A programming model for exascale

- **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

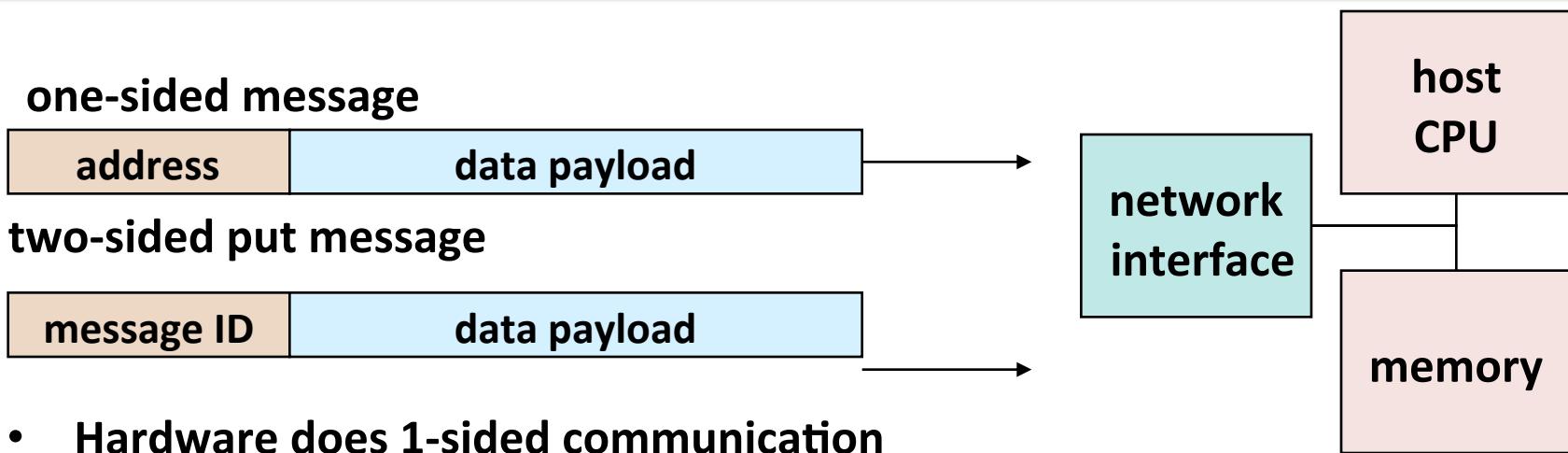
`shared int [ ] ga; and upc_malloc (...)`



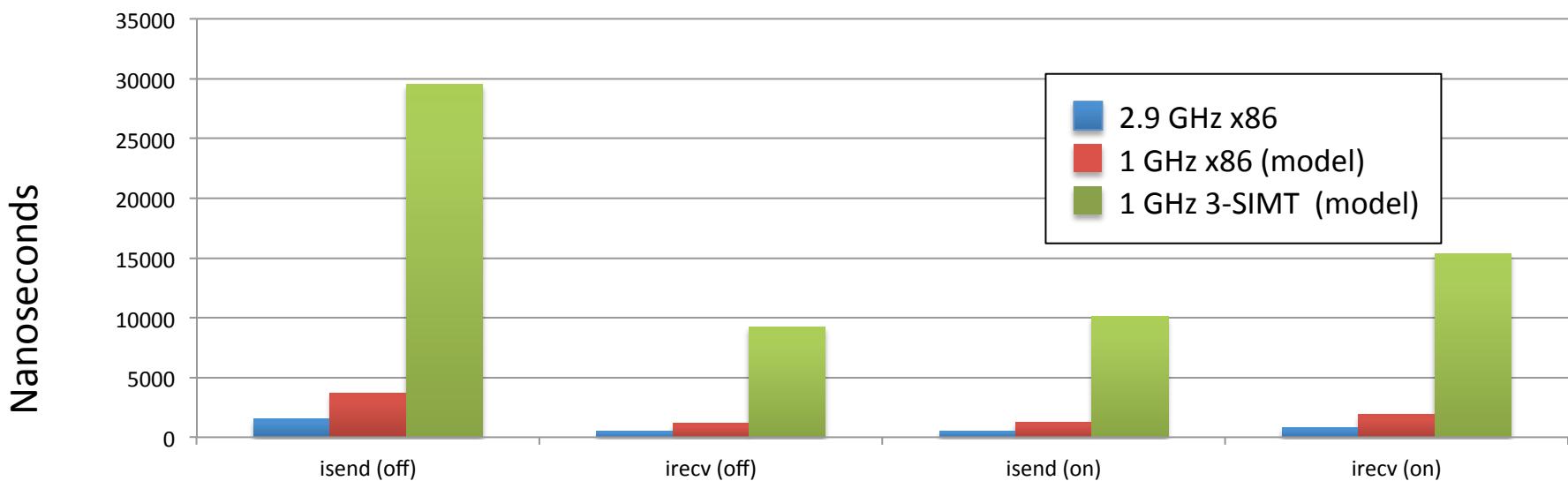
Examples:  
UPC  
UPC++

A programming model can influence how programmers think

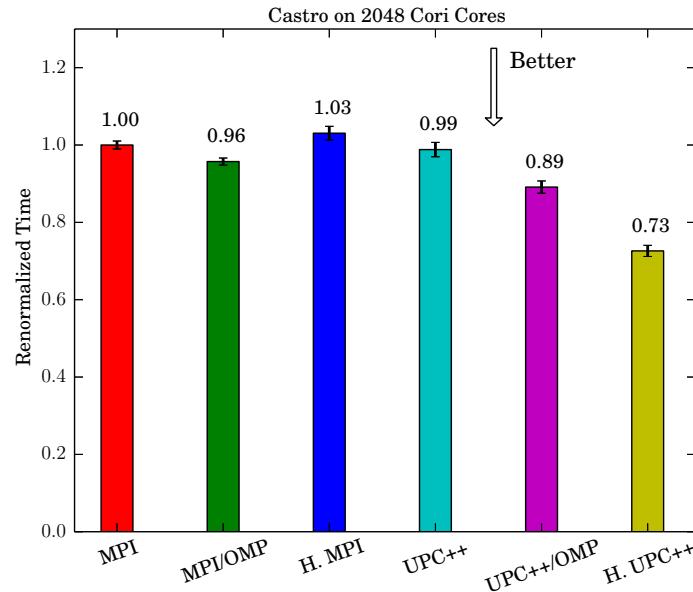
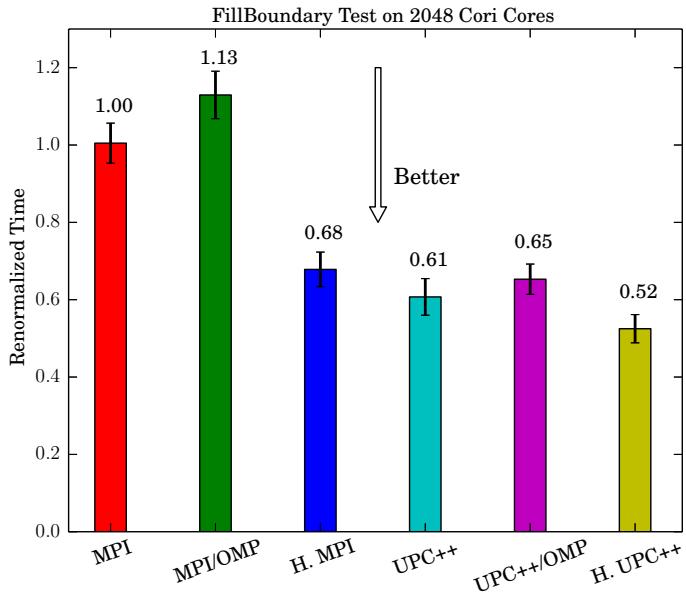
# One-Sided Communication is Closer to Hardware



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



# 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

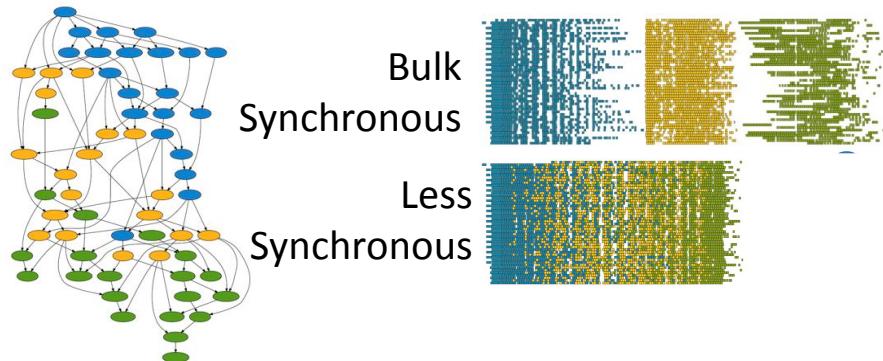
# Sources of Unnecessary Synchronization

## 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%

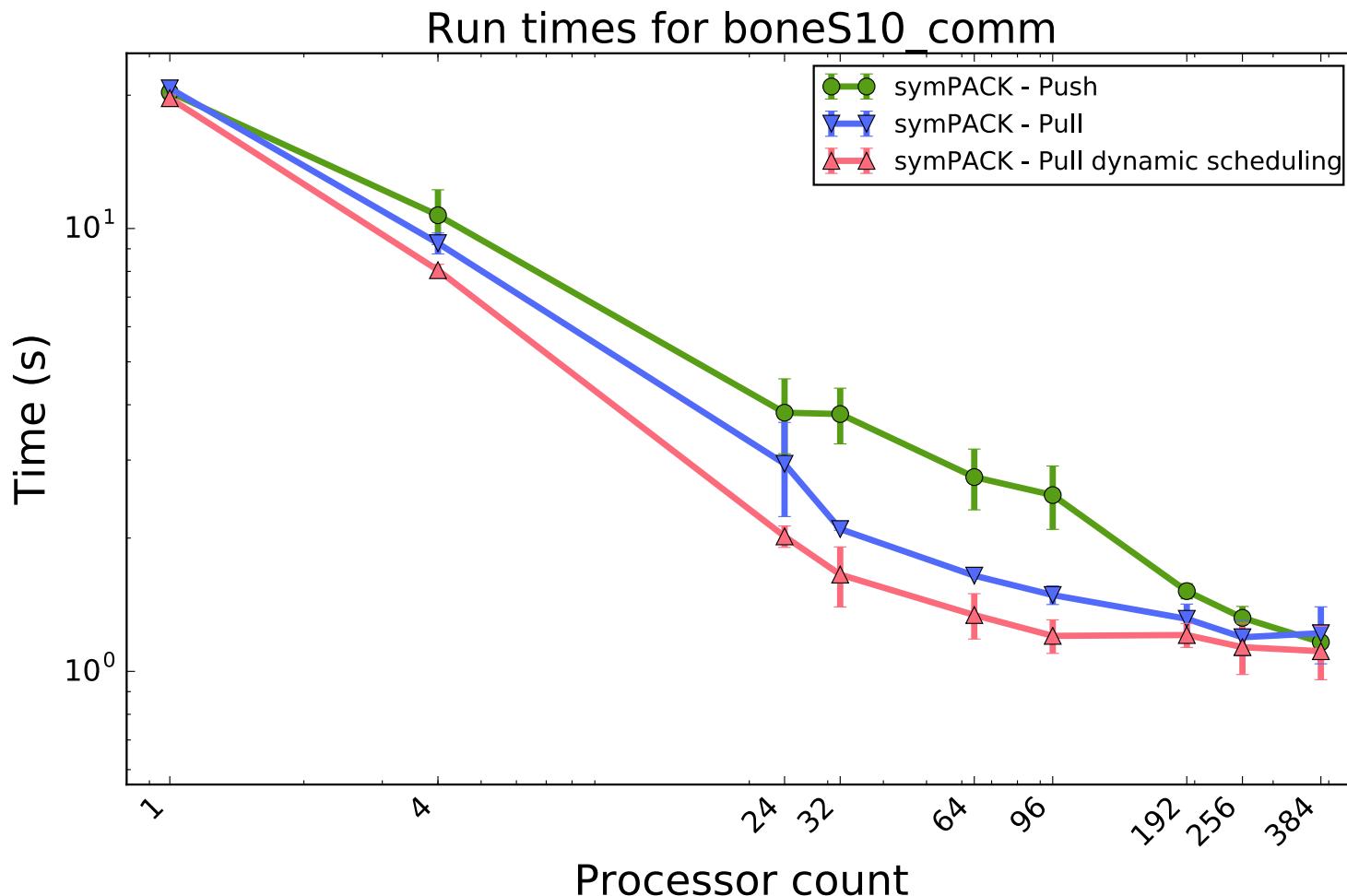
NWChem: most of barriers are unnecessary (Corvette)

## Accelerator Offload

```
!$acc data copyin(cx,cx1,cx2,cx3,cx4,cx5,cx6,cx7,cx8,cx9,cx10,cx11,&
!$acc& cx12,cx13,cx14,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,&
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!$acc& streaming_rbuf1,streaming_rbuf2,streaming_rbuf4,streaming_rbuf5,&
!$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

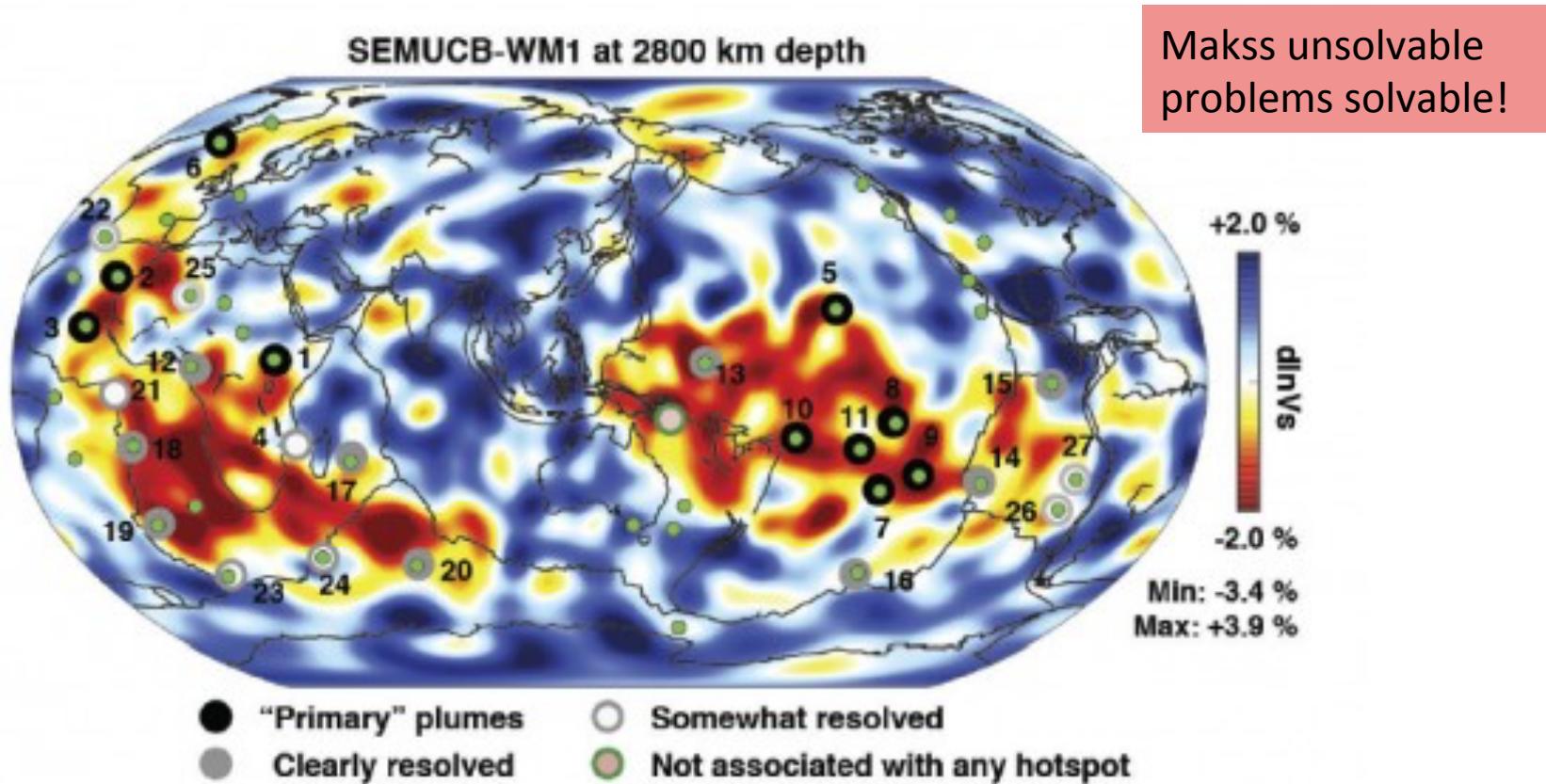
# Asynchronous Sparse Cholesky in UPC++



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

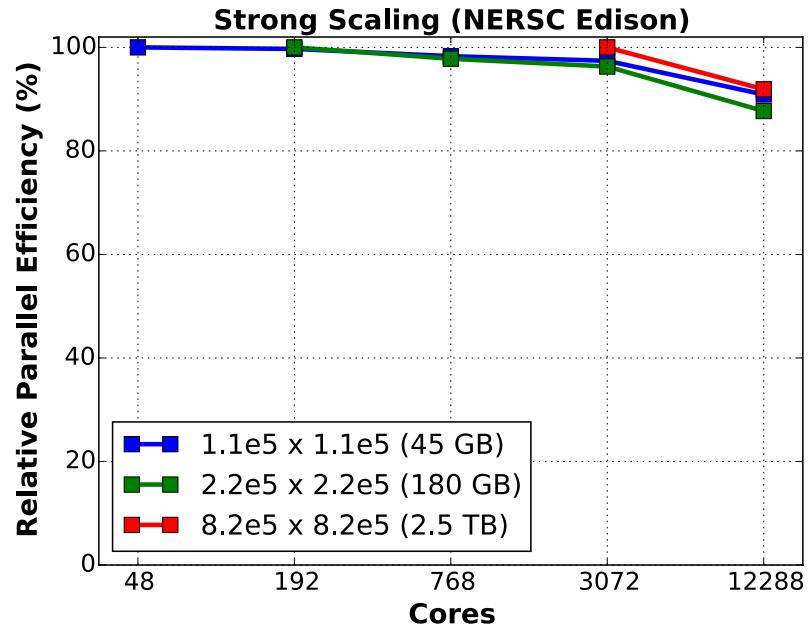
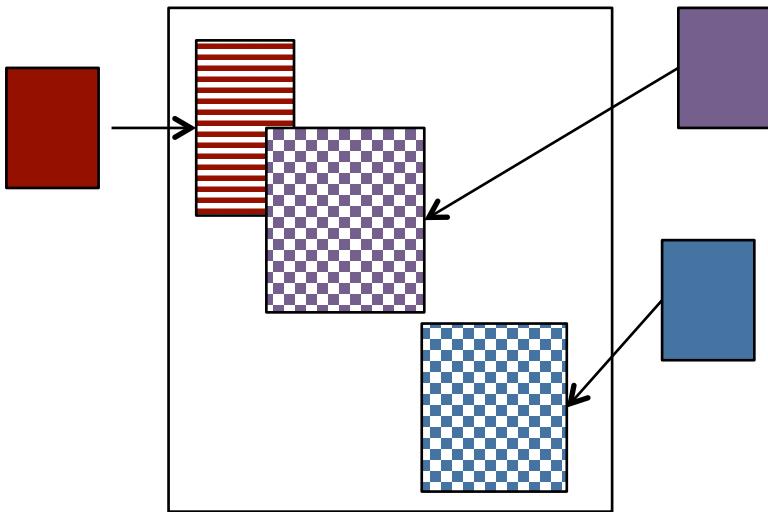
# Whole-Mantle Seismic Model Using UPC++

- 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

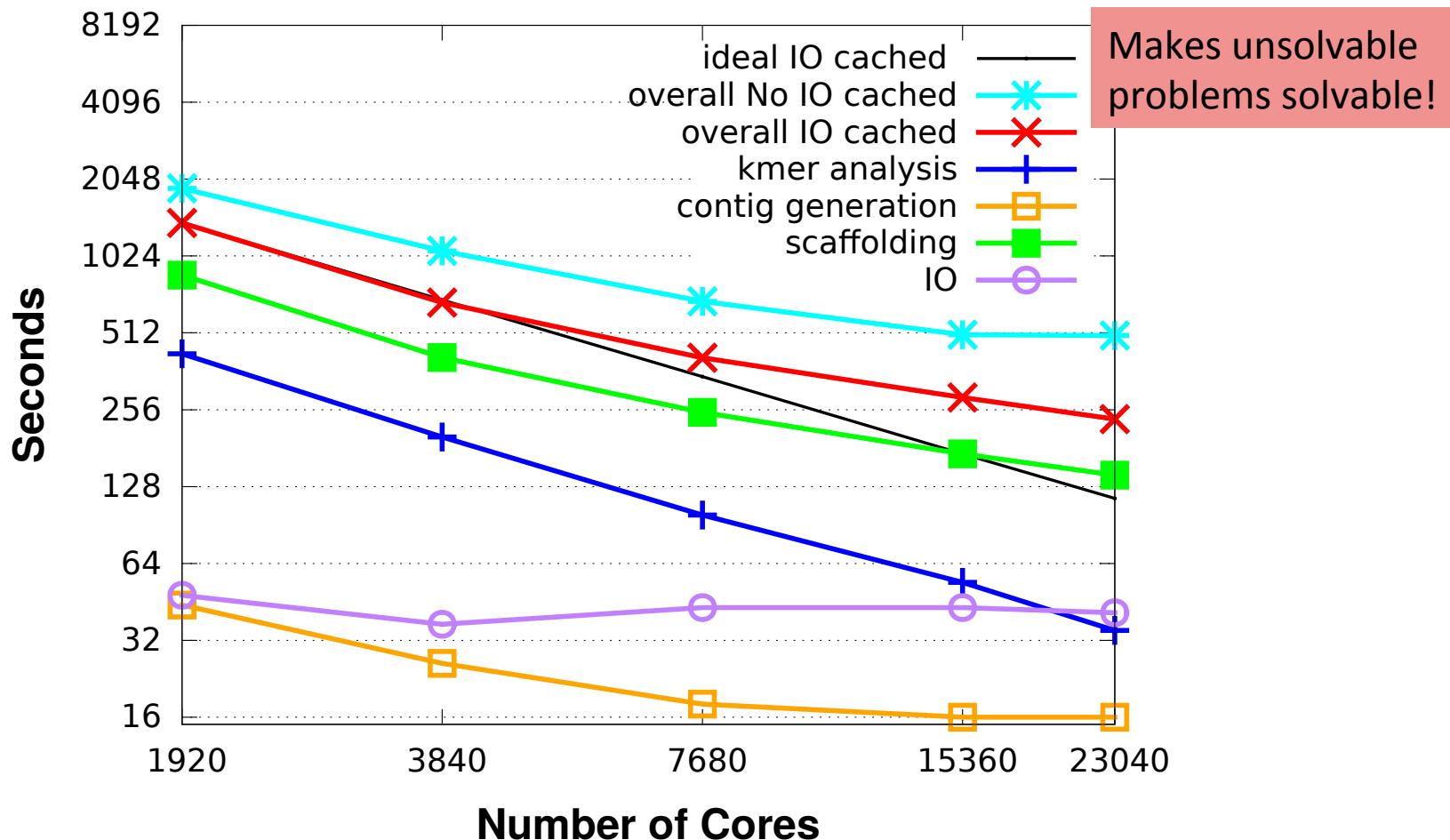
Scott French, Y. Zheng, B. Romanowicz, K. Yelick

# *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!