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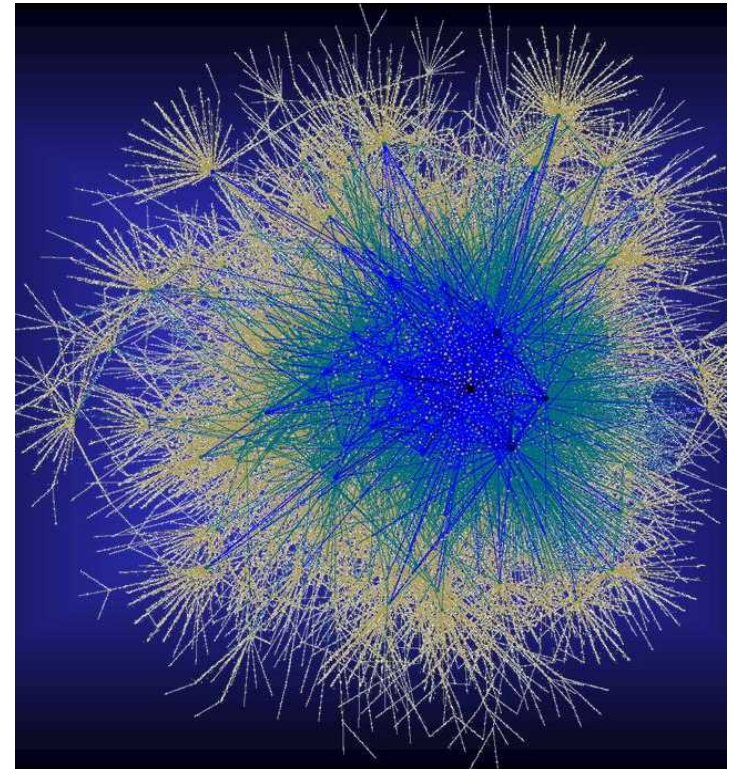
2D Partitioning for Scalable Matrix Computations on Scale-Free Graphs

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Introduction

- *Big data* is a challenge to HPC
 - Large graphs/networks are pervasive
 - E.g., WWW, social networks
 - Scale-free, small-world
 - Skewed degree distribution
 - Very different from PDE discretizations
- How to do efficient parallel computations?
 - Using distributed-memory computers
 - Data layout is important



BGP graph (credit: Richardson, Chung)
<http://math.ucsd.edu/~fan/graphs/gallery>

Key Message

- 2D (edge-based) partitioning is important for data analytics.
 - Scale-free, small-world, skewed degree distribution.
 - Long predicted, but demonstrated only recently, and not yet widely appreciated.
- Computing a “good” 2D distribution is no harder than 1D!
 - Still active research area.

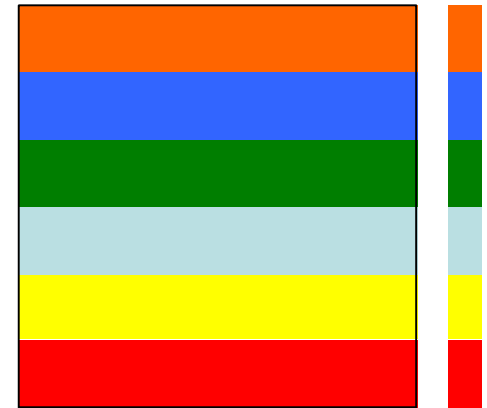
Sparse matvec (SpMV) important

- Linear algebra is a useful analysis tool for graphs
 - Both adjacency matrix and graph Laplacian are of interest
 - Spectral analysis using extreme eigenpairs
 - SpMV is core kernel in iterative methods
- SpMV is bottleneck for scale-free graphs on large distributed-memory computers
 - Example: For a social network (orkut) on 64 cores
 - SpMV took 95% of the compute time in an eigensolver
 - Maximum #messages, over all processes, is typically $p-1$
 - Due to some very high-degree vertices

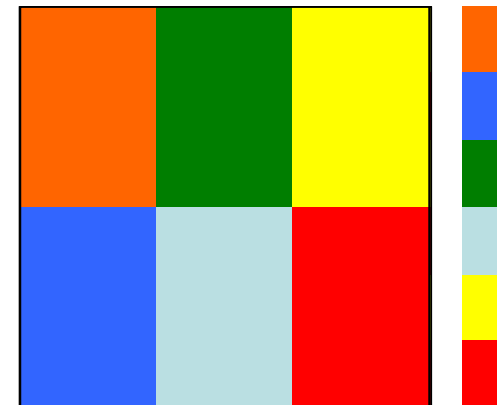
1D and 2D Matrix Distributions

We view graphs as sparse matrices.

- 1D (vertex) distribution:
 - Entire rows (or columns) of matrix assigned to a processor
 - Required in most software
- 2D (edge) distribution:
 - Cartesian methods: Each process owns intersection of some rows & columns
 - Processes are *logically* arranged in a 2D grid
 - This limits #messages per process to $O(\sqrt{p})$
 - Long used in parallel dense solvers (ScaLapack)
 - Beneficial also for sparse matrices (Fox et al. '88, Lewis & van de Geijn '93, Hendrickson et al. '95)
 - Yoo et al. (SC'11) demonstrated benefit over 1D layouts for eigensolves on scale-free graphs



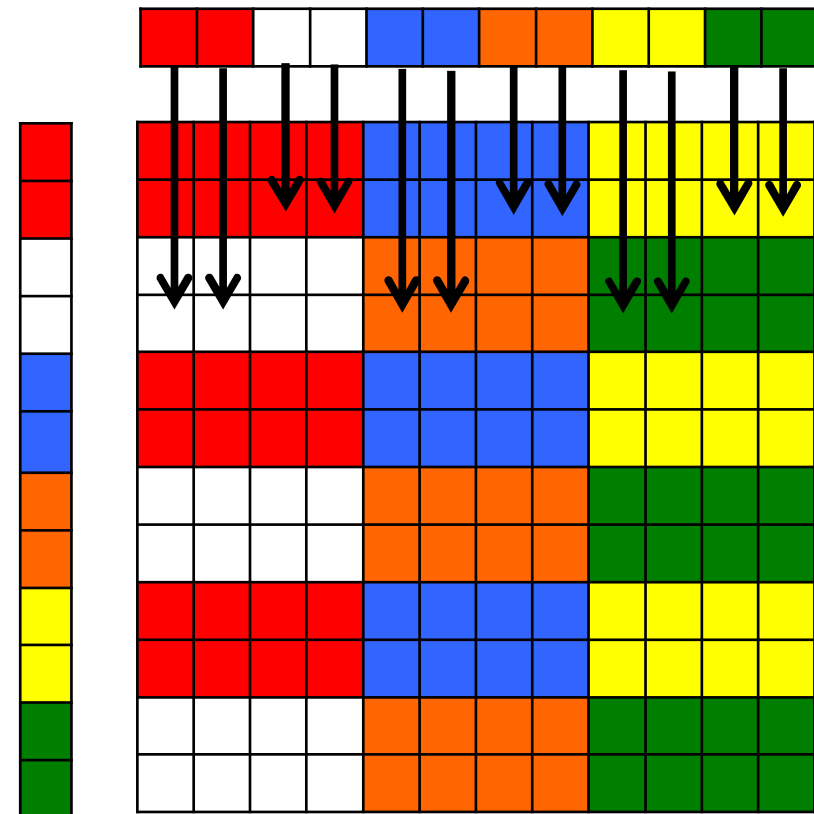
1D row-wise matrix distribution; 6 processes



2D matrix distribution; 6 processes

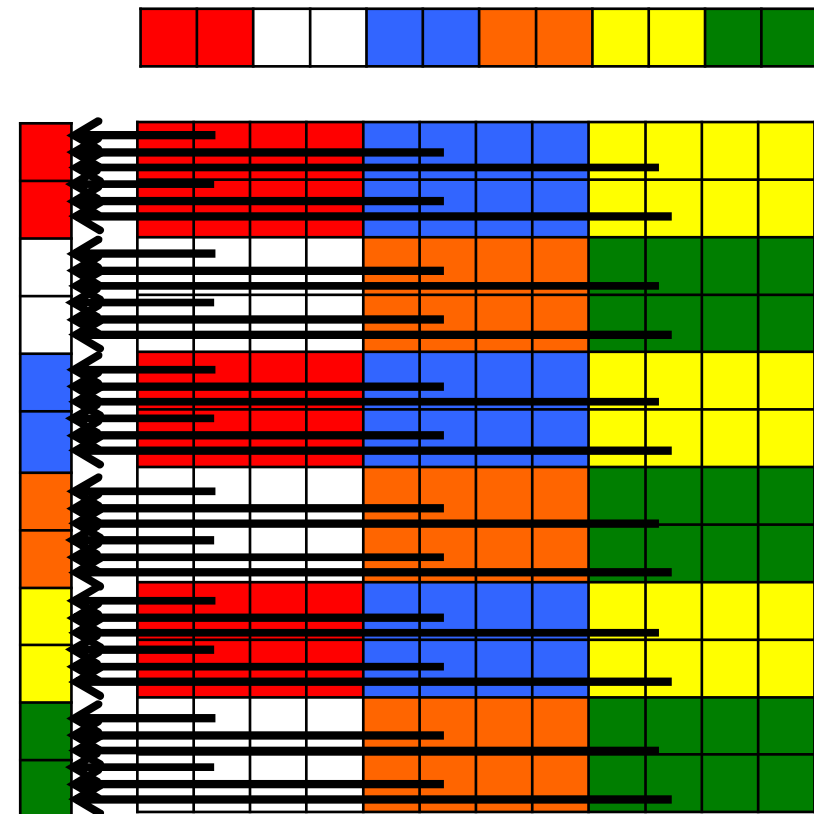
Benefit of 2D Matrix Distribution

- During matrix-vector multiplication ($y=Ax$), communication occurs only along rows or columns of processors.
 - Expand (vertical):
Vector entries x_j sent to column processors to compute local product $y^p = A^p x$
 - Fold (horizontal):
Local products y^p summed along row processors; $y = \sum y^p$
- In 1D, fold is not needed, but expand may be all-to-all.



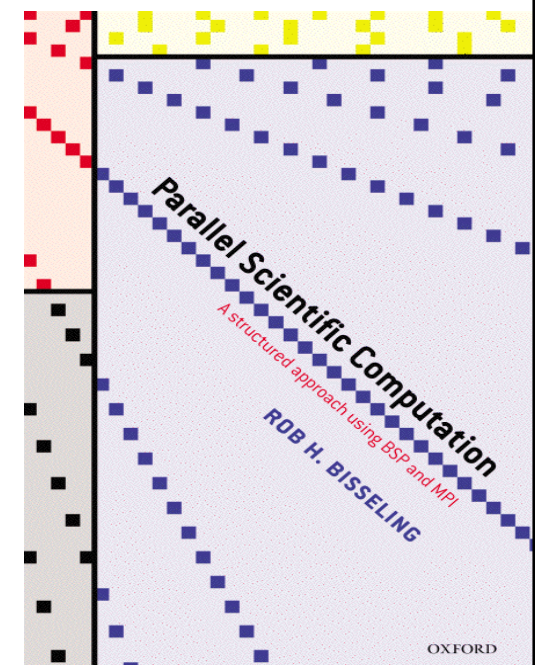
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2D Partitioning Methods

- Cartesian 2D block (Fox et al. '88)
 - Simple/fast to compute but ignores the structure of the graph.
 - Low #messages, but communication volume may be high.
- Coarse-grain hypergraph (Catalyurek & Aykanat '01)
 - Cartesian product, but uses the matrix structure.
 - Requires multiconstraint hypergraph partitioning.
- Fine-grain hypergraph (Catalyurek & Ayk. '01)
 - Assign each nonzero separately
 - Not Cartesian, high #messages
 - Larger hypergraph, impractical for big problems
- Mondriaan (Vastenhouw & Bisseling '05)
 - Recursive bisection, hypergraph partitioning
 - Not Cartesian, no bound on #messages



Trilinos Computational Science Toolkit



- Collection of ~60 packages
 - Heroux et al., Sandia
- Trilinos Capabilities:
 - Scalable Linear & Eigen Solvers
 - Discretizations, Meshes & Load Balancing
 - Nonlinear & Optimization Solvers
 - Software Engineering Technologies & Integration
- In this project, we used
 - Distributed Matrix/Vector classes *Epetra*
 - Partitioning package *Zoltan*
 - Eigensolver package *Anasazi*

Petra Object Model

- Maps describe the distribution of global IDs for rows/columns/vector entries to processors.
- Four maps needed in most general case:
 - Row map for matrix
 - Column map for matrix
 - Range map for vector
 - Domain map for vector
- Implemented in *Epetra* (and *Tpetra*) packages
- Allows 2D distributions!

Load-Balancing by Randomization

- Simple “block” partitioning balances rows but not nonzeros
- Randomization is a simple but powerful technique
- On input, randomly permute matrix rows/columns
 - Eliminates any inherent structure in input file (e.g., high degree nodes first)
 - Gives better balance in number of nonzeros per processor for 1D and 2D
 - But can drastically increase communication volume

liveJournal matrix (4M rows; 73M nonzeros) on 1024 processes				
Method	Imbalance in nonzeros (Max/Avg per proc)	Max # Messages per SpMV	Comm. Vol. per SpMV (doubles)	100 SpMV time (secs)
1D-Block	12.8	1023	34.5M	2.14
1D-Random	1.3	1023	55.3M	1.52
2D-Block	11.4	62	43.4M	0.95
2D-Random	1.0	62	64.2M	0.43

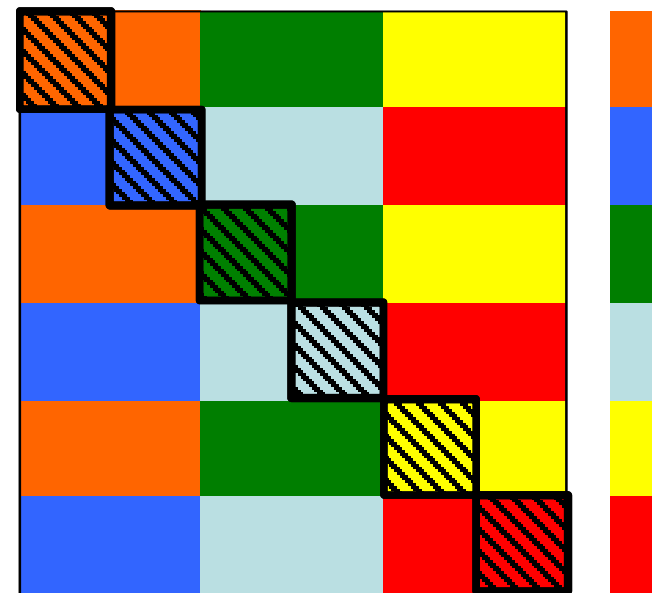
New Method: Graph Partitioning + 2D

- Our idea: Apply (hyper)graph partitioning and 2D distribution together
 - Compute vertex-based partition of graph using ParMETIS or Zoltan
 - Apply 2D distribution to the resulting permuted graph/matrix
- Advantages:
 - Balance the number of nonzeros per process
 - Exploit structure in the graph to reduce communication volume
 - Reduce the number of messages via 2D distribution
- Don't optimize a single objective but try do fairly well in all

2D (Hyper-)Graph Partitioning (GP/HP)

- Partition vertices of original graph into p parts
 - Using standard (hyper)graph partitioner
- Implicitly, let $A_{\text{perm}} = PAP^T$
 - Where P is permutation from partitioning above
- Assign A_{perm} to processes using Cartesian block 2D layout

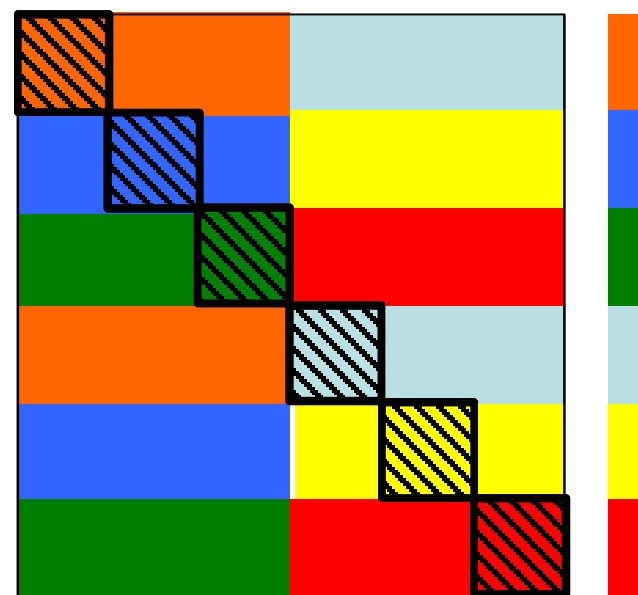
Due to partitioning, diagonal blocks of A_{perm} will be denser:



Observations

- We first partition into p parts
 - NOT \sqrt{p}
- Many choices for Cartesian 2d layout in second step
 - Fast method: just use (i,j) indices, ignore structure
 - Future: Pick “best” option based on structure

Example: Another possible 2d layout



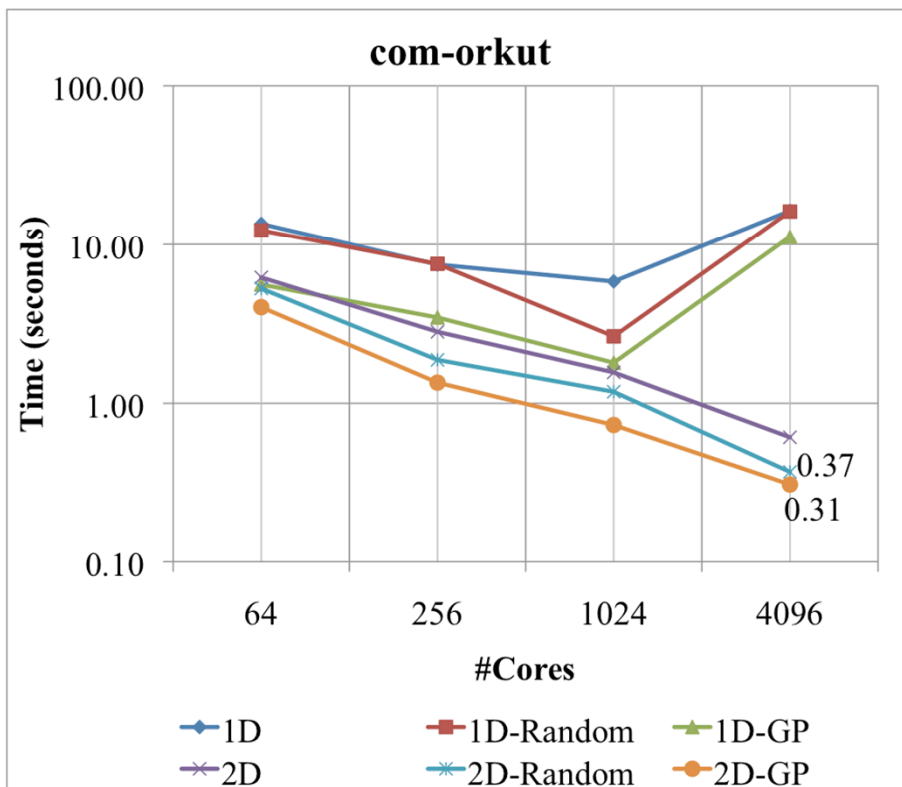
Results 1D vs 2D (Block, Random, GP)

Platform: cab cluster at LLNL (1200 Intel Xeon E5 16-core nodes operating at 2.6 GHz, 32 GB memory/node, Infiniband)

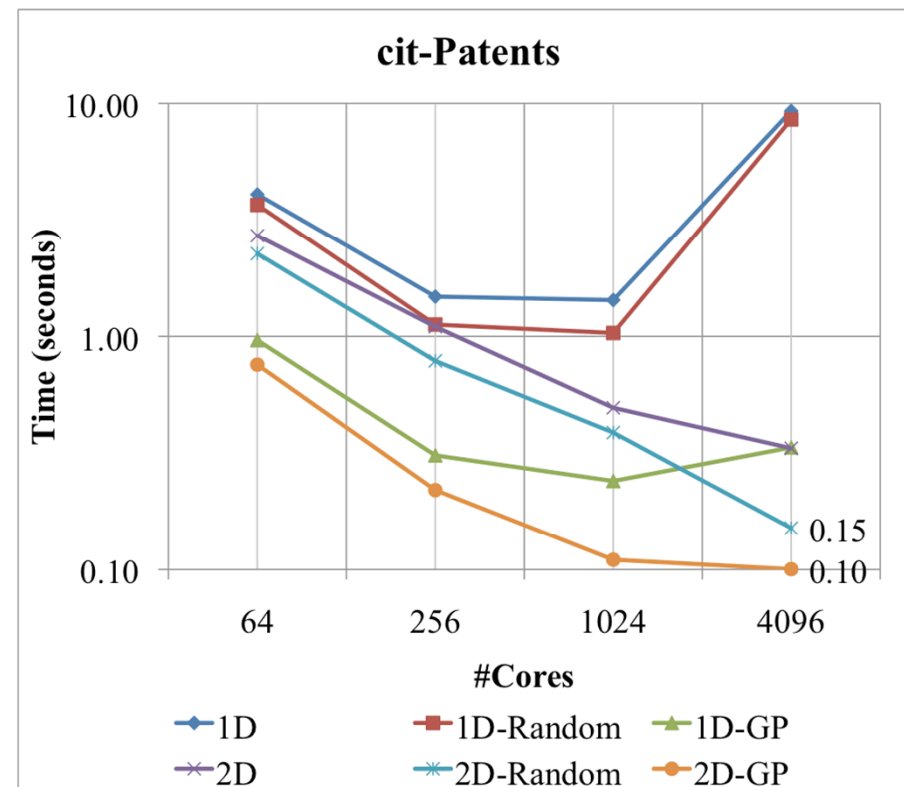
All matrices from UF collection (some originally from SNAP, etc.)

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1D-Block	12.8	1023	34.5M	2.14
1D-Random	1.3	1023	55.3M	1.52
1D-GP	1.2	1011	18.9M	0.53
2D-Block	11.4	62	43.4M	0.95
2D-Random	1.0	62	64.2M	0.43
2D-GP	1.4	62	22.4M	0.22

Strong scaling



Orkut social network
3.1M rows; 237M nonzeros
Max nonzeros/row = 33K

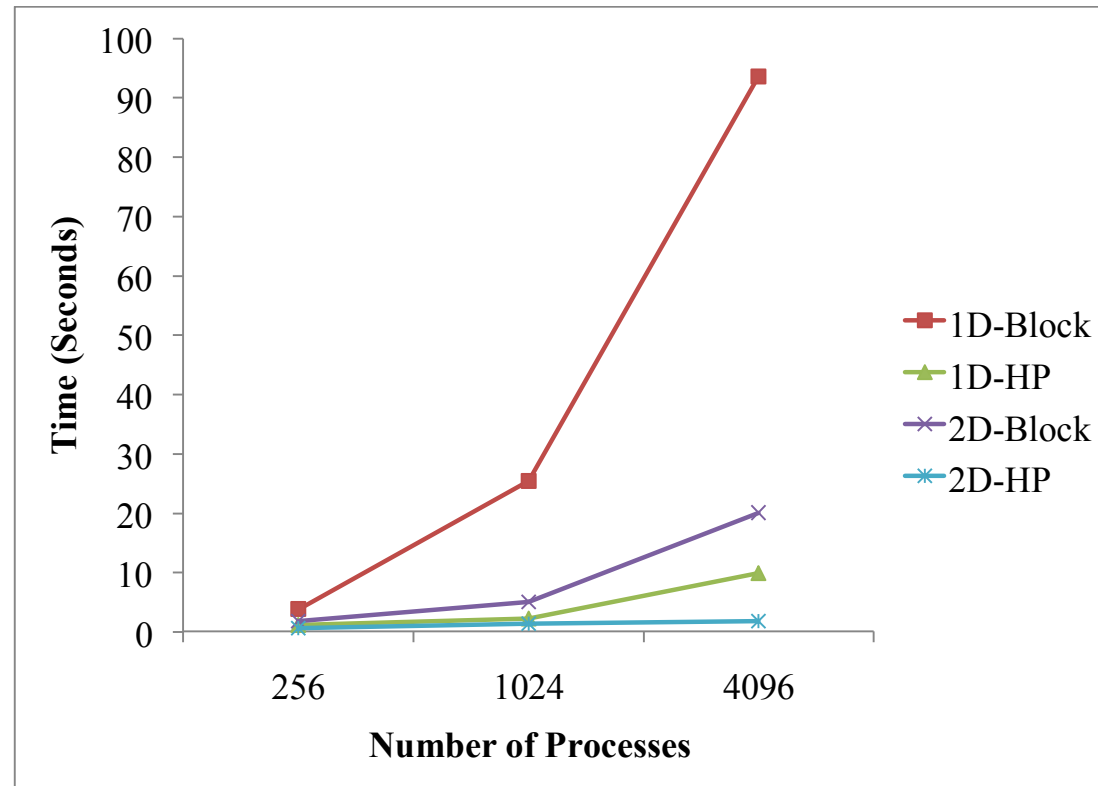


Patent citations network
3.8M rows; 37M nonzeros
Max nonzeros/row = 1K

1D stops scaling around 1024 processes due to high communication cost.

“Weak Scaling”

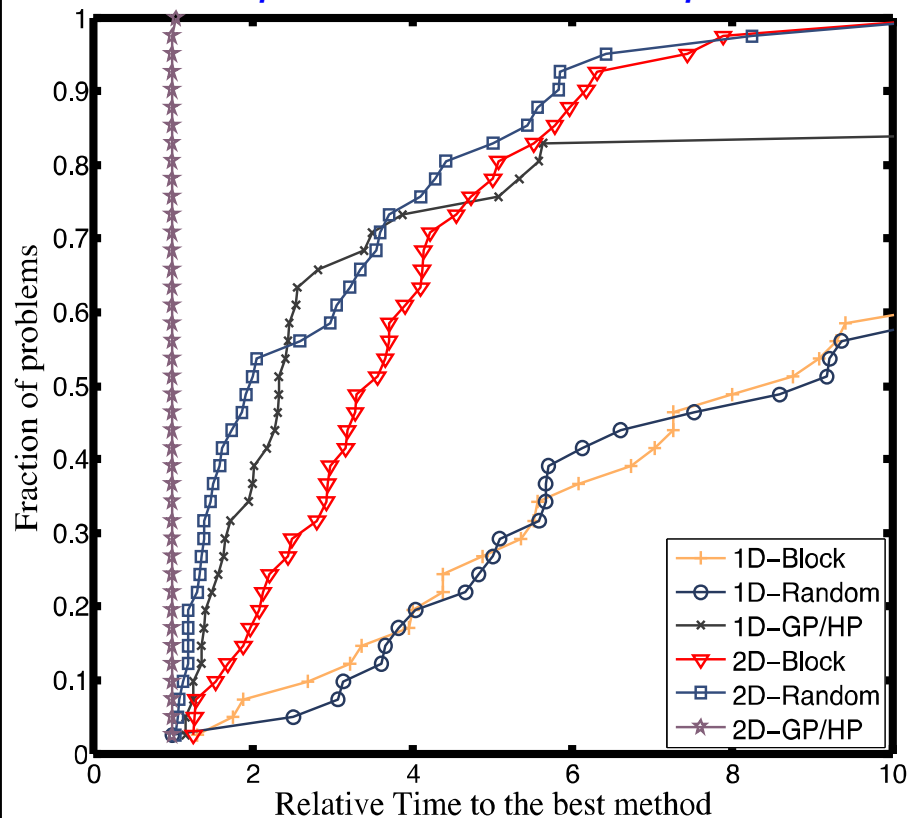
- R-MAT matrices (Chakrabarti et al., 2004) with Graph-500 parameters ($a=0.57$; $b=c=0.19$; $d=0.05$)
 - rmat_22 on 256 procs
 - 4.2M vertices
 - 38M edges
 - rmat_24 on 1024 procs
 - 16.8M vertices
 - 151M edges
 - rmat_26 on 4096 procs
 - 67.1M vertices
 - 604M edges
- Times for 100 SpMV
- 2D-HP maintains best weak scaling.



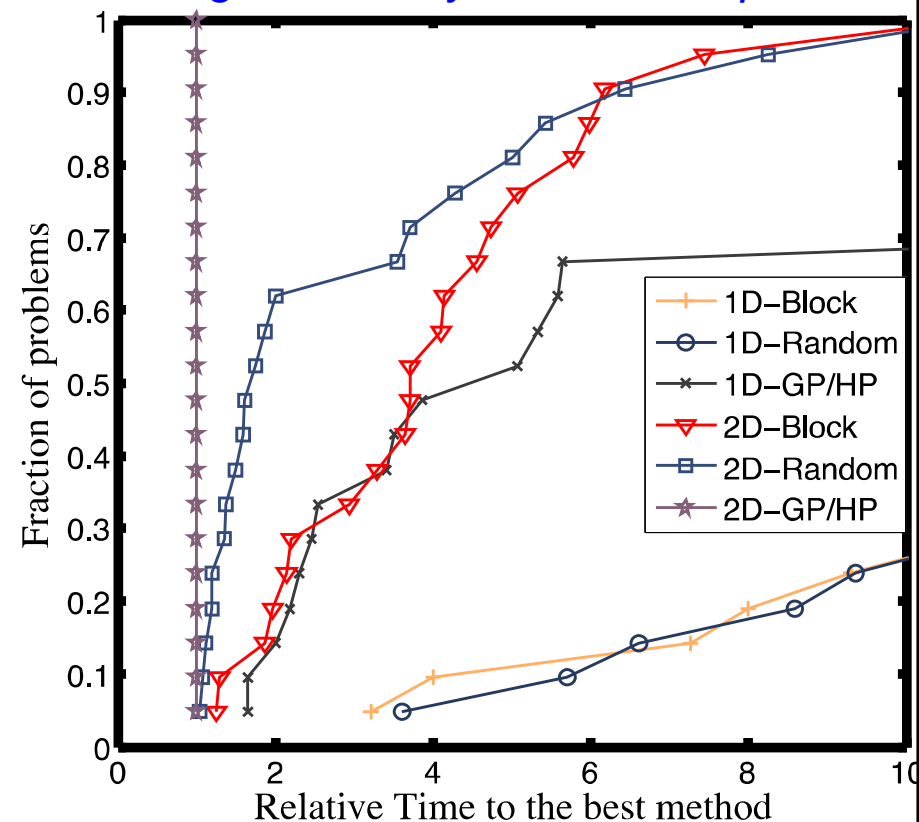
Performance Profile

- 10 matrices: 1.1M - 67.5M rows; 36M-1.6B nonzeros
- 2D-GP/HP best in all but one experiment
- Benefit of 2D even greater for large numbers of processes

All experiments: 64-4096 procs



Large runs only: 1024-4096 procs



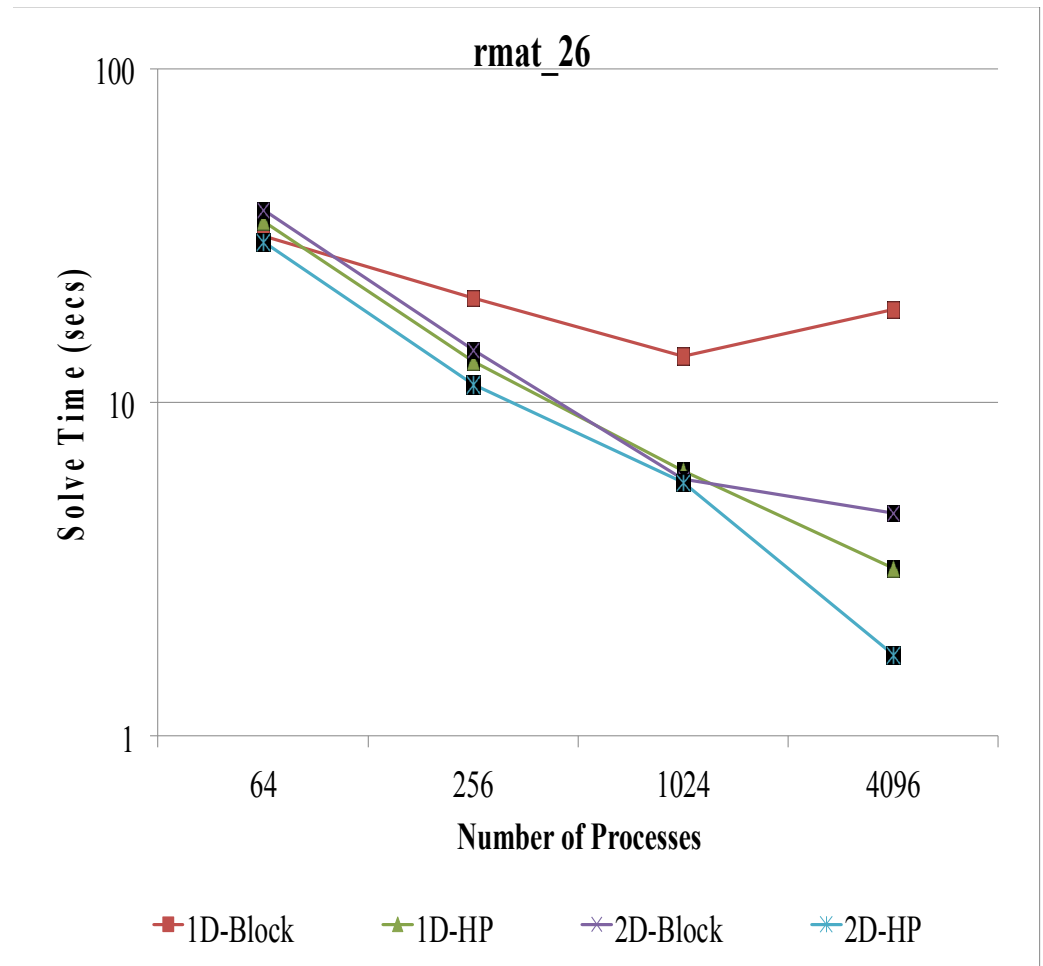
Eigensolver Experiments

■ Anasazi Toolkit in Trilinos

- Baker, Hetmaniuk, Lehoucq, Thornquist; ACM TOMS 2009
- Block-based eigensolvers: Solve $AX = X\Lambda$ or $AX = BX\Lambda$

■ Experiment:

- Find 10 largest eigenvalues of Laplacian using Block Krylov-Schur (BKS) solver
- rmat_26 matrix: 67.1M rows; 604M nonzeros



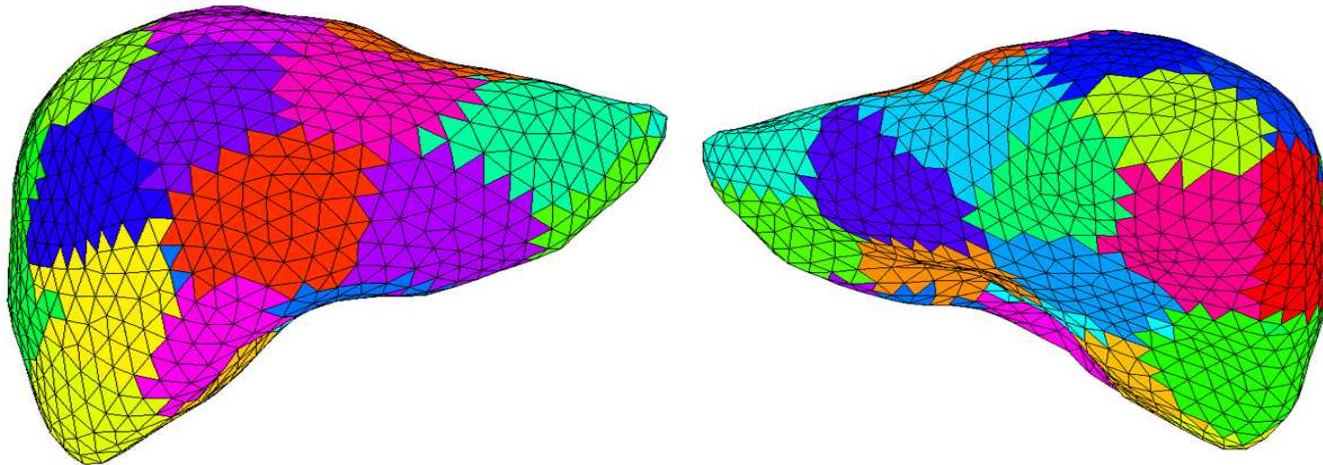
Conclusions

- 2D distributions allow scalable parallel computations for small-world (scale-free) graphs.
 - For 1D, must use “vertex delegates” or “disaggregation”
- 1D (hyper)graph partitioning is effective on scale-free graphs for moderate number of processes.
 - Good load balance, low communication volume
- Combining 2D distribution with (hyper)graph partitioning gives best results.
 - Low number of messages, low communication volume, low imbalance.
 - Allows reuse of existing partitioning software.
- Ongoing/future work:
 - Compare to other 2D partitioning methods.
 - Use faster partitioning method in 1st step (e.g., PULP)
 - Optimize 2nd step in algorithm (Cartesian layout)

Extra Slides

Data Partitioning

- (Hyper-)graph partitioning generally reduces communication for SpMV
- Software tools (e.g., Metis, Scotch, Zoltan) were designed for meshes and PDE discretizations
 - Not optimized for scale-free graphs
 - Focus has been on cut edges and communication volume
 - We also wish to reduce #messages



Test Matrices & Platform

- Compare times for 100 matrix-vector products with 1D and 2D distributions
- Platform: cab cluster at LLNL (1200 Intel Xeon E5 16-core nodes operating at 2.6 GHz, 32 GB memory/node, Infiniband)
- Matrices from the University of Florida matrix collection. (Symmetrized, if needed)

Name	Description	Number of Rows	Number of Nonzeros
Hollywood-2009	Hollywood movie actor network (Boldi, Rosa, Santini, Vigna)	1.1M	113M
Wikipedia-20070206	Links between wikipedia pages (Gleich)	3.5M	85M
Ljournal-2008	LiveJournal social network (Boldi, Rosa, Santini, Vigna)	5.6M	99M
Wb-edu	Links between *.edu webpages (Gleich)	8.9M	88M
Cit-Patents	Citation network among US patents (Hall, Jaffe, Trajtenberg)	3.8M	33M

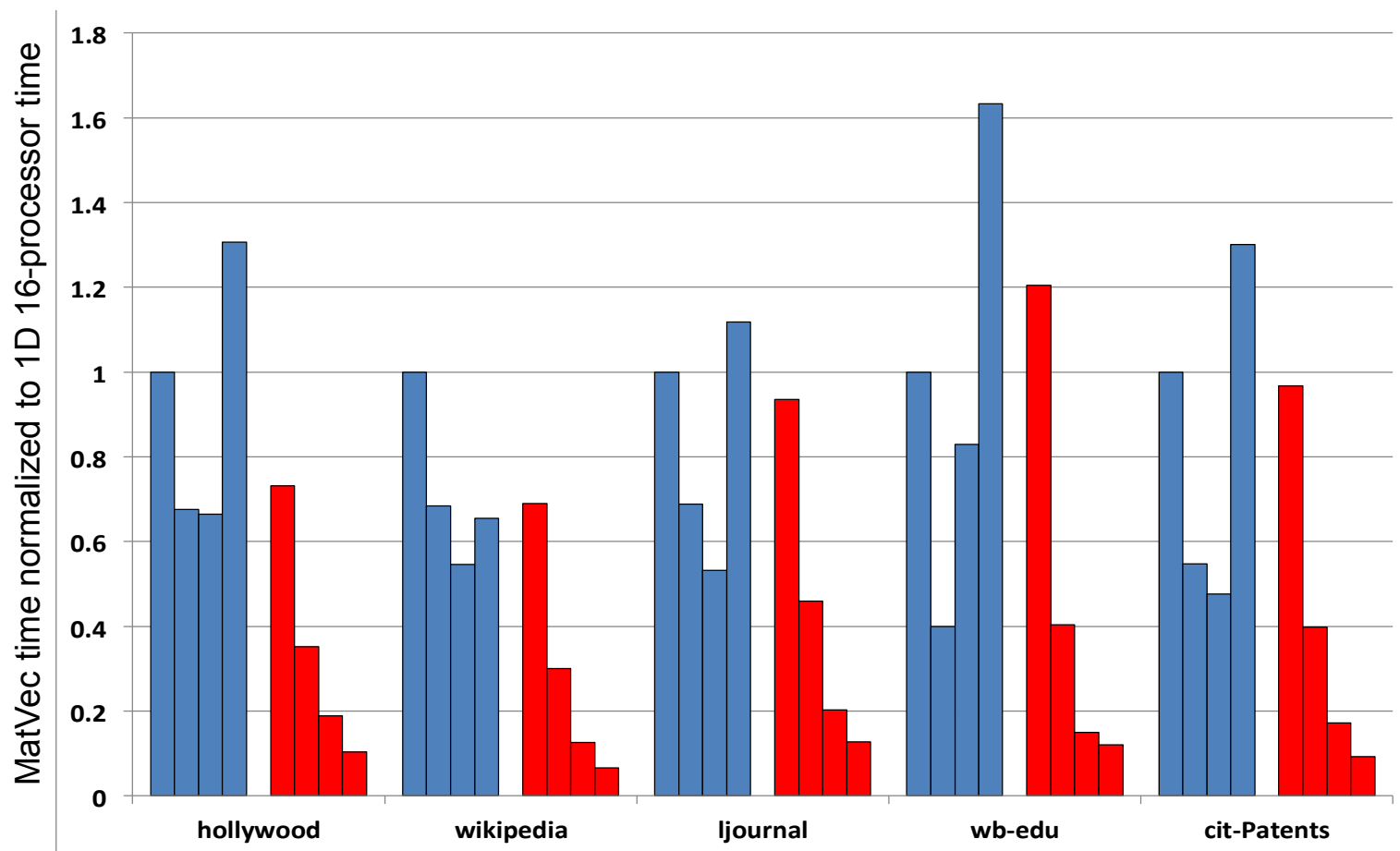
1D vs 2D Strong Scaling experiments

For each matrix:

Blue = Trilinos 1D Matrix Distribution on 16, 64, 256, 1024 processors (left to right)

Red = Trilinos 2D Matrix Distribution on 16, 64, 256, 1024 processors (left to right)

Times are normalized to the 1D 16-processor runtime for each matrix.



Trilinos: Petra Object Model

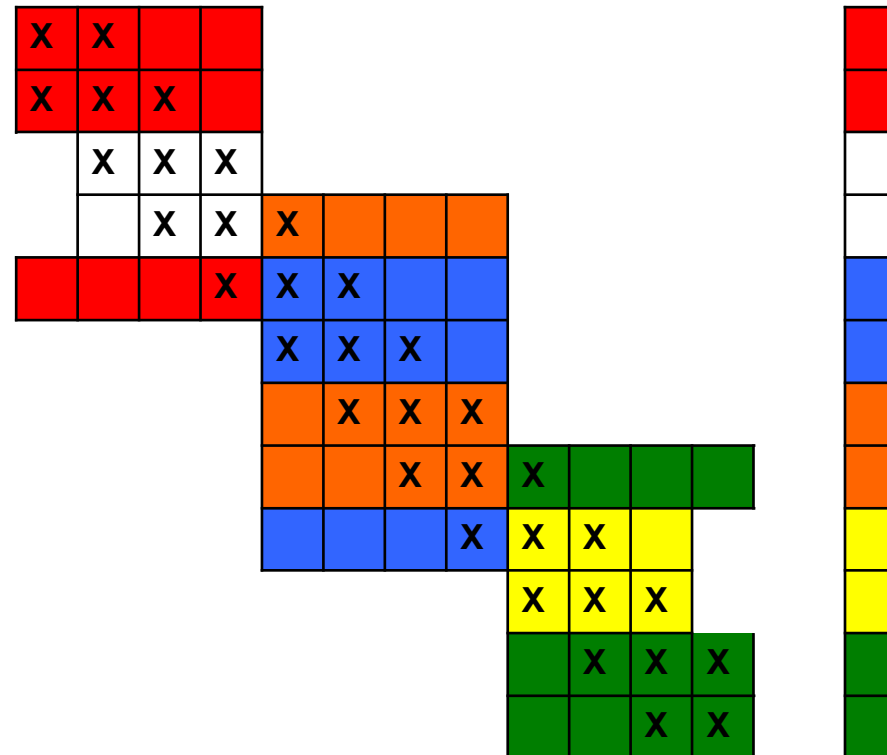
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Rank 3 (Blue)

Row Map = {4, 5, 8}

Column Map = {4, 5, 6, 7}

Range/Domain Map = {4, 5}



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