



Interactive Tools for Combinatorial Supercomputing

John R. Gilbert

University of California, Santa Barbara

Viral Shah, Imran Patel (UCSB)

Alan Edelman (MIT and Interactive Supercomputing)

Ron Choy, David Cheng (MIT)

Parry Husbands (Lawrence Berkeley Lab)

Steve Reinhardt, Todd Letsche (SGI)

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Parallel Computing Today



Columbia,
NASA Ames Research Center

Departmental Beowulf cluster



But!

How do you program it?

C with MPI

```
#include <stdio.h>
#include <stdlib.h>
#include <math.h>
#include "mpi.h"
#define A(i,j) ( 1.0/((1.0*(i)+(j))*(1.0*(i)+(j)+1)/2 + (1.0*(i)+1)) )
void errorExit(void);
double normalize(double* x, int mat_size);
int main(int argc, char **argv)
{
    int num_procs;
    int rank;
    int mat_size = 64000;
    int num_components;
    double *x = NULL;
    double *y_local = NULL;
    double norm_old = 1;
    double norm = 0;
    int i,j;
    int count;
    if (MPI_SUCCESS != MPI_Init(&argc, &argv)) exit(1);
    if (MPI_SUCCESS != MPI_Comm_size(MPI_COMM_WORLD,&num_procs)) errorExit();
```

C with MPI (2)

```
if (0 == mat_size % num_procs) num_components = mat_size/num_procs;
else num_components = (mat_size/num_procs + 1);
mat_size = num_components * num_procs;
if (0 == rank) printf("Matrix Size = %d\n", mat_size);
if (0 == rank) printf("Num Components = %d\n", num_components);
if (0 == rank) printf("Num Processes = %d\n", num_procs);
x = (double*) malloc(mat_size * sizeof(double));
y_local = (double*) malloc(num_components * sizeof(double));
if ( (NULL == x) || (NULL == y_local) )
{
    free(x);
    free(y_local);
    errorExit();
}
if (0 == rank)
{
    for (i=0; i<mat_size; i++)
    {
        x[i] = rand();
    }
    norm = normalize(x,mat_size);
}
```

C with MPI (3)

```
if (MPI_SUCCESS !=
    MPI_Bcast(x, mat_size, MPI_DOUBLE, 0, MPI_COMM_WORLD)) errorExit();
count = 0;
while (fabs(norm-norm_old) > TOL) {
    count++;
    norm_old = norm;
    for (i=0; i<num_components; i++)
    {
        y_local[i] = 0;
    }
    for (i=0; i<num_components && (i+num_components*rank)<mat_size; i++)
    {
        for (j=mat_size-1; j>=0; j--)
        {
            y_local[i] += A(i+rank*num_components,j) * x[j];
        }
    }
    if (MPI_SUCCESS != MPI_Allgather(y_local, num_components, MPI_DOUBLE, x,
        num_components, MPI_DOUBLE, MPI_COMM_WORLD)) errorExit();
}
```

C with MPI (4)

```
    norm = normalize(x, mat_size);
}
if (0 == rank)
{
    printf("result = %16.15e\n", norm);
}
free(x);
free(y_local);

MPI_Finalize();
exit(0);
}
void errorExit(void)
{
    int rank;
    MPI_Comm_rank(MPI_COMM_WORLD, &rank);
    printf("%d died\n", rank);
    MPI_Finalize();
    exit(1);
}
```

C with MPI (5)

```
double normalize(double* x, int mat_size)
{
    int i;
    double norm = 0;
    for (i=mat_size-1; i>=0; i--)
    {
        norm += x[i] * x[i];
    }
    norm = sqrt(norm);
    for (i=0; i<mat_size; i++)
    {
        x[i] /= norm;
    }
    return norm;
}
```



```
A = rand(4000*p, 4000*p);  
x = randn(4000*p, 1);  
y = zeros(size(x));  
while norm(x-y) / norm(x) > 1e-11  
    y = x;  
    x = A*x;  
    x = x / norm(x);  
end;
```

Background

- Matlab*P 1.0 (1998): Edelman, Husbands, Isbell (MIT)
- Matlab*P 2.0 (2002-): MIT / UCSB / LBNL
- Star-P (2004-): Interactive Supercomputing / SGI



Data-Parallel Operations

< M A T L A B >

Copyright 1984-2001 The MathWorks, Inc.

Version 6.1.0.1989a Release 12.1

```
>> A = randn(500*p, 500*p)
A = ddense object: 500-by-500
```

```
>> E = eig(A);
```

```
>> E(1)
ans = -4.6711 +22.1882i
```

```
e = E(:, :);
```

```
>> ppwhos
```

Name	Size	Bytes	Class
A	500px500p	688	ddense object
E	500px1	652	ddense object
e	500x1	8000	double array (complex)

Task-Parallel Operations

```
>> quad('4./(1+x.^2)', 0, 1);  
ans = 3.14159270703219
```

```
>> a = (0:3*p) / 4  
a = ddense object: 1-by-4
```

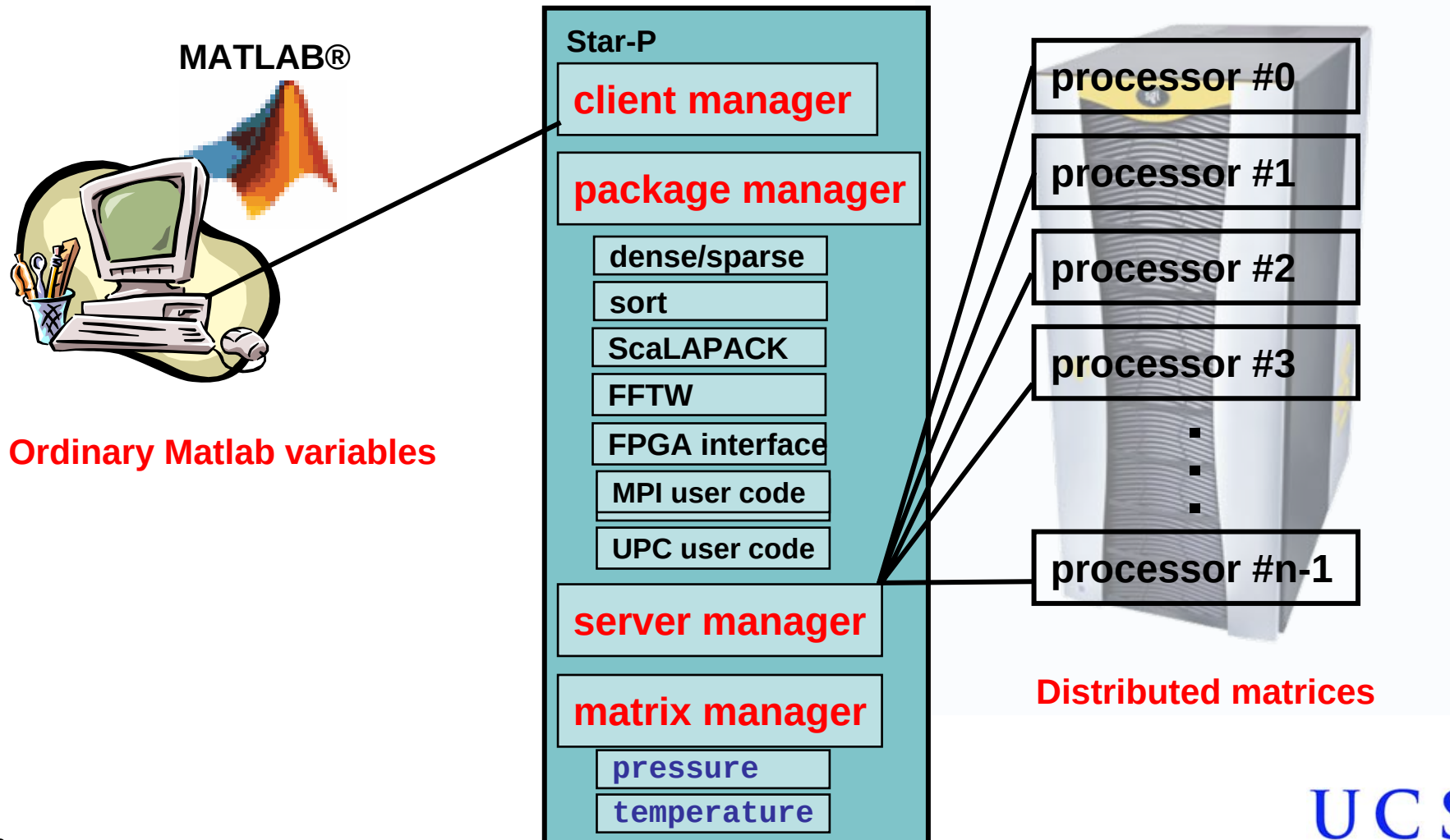
```
>> a(:, :)  
ans =  
  
0  
0.2500000000000000  
0.5000000000000000  
0.7500000000000000
```

```
>> b = a + .25;
```

```
>> c = ppeval('quad', '4./(1+x.^2)', a, b);  
c = ddense object: 1-by-4
```

```
>> sum(c)  
ans = 3.14159265358979
```

Star-P Architecture

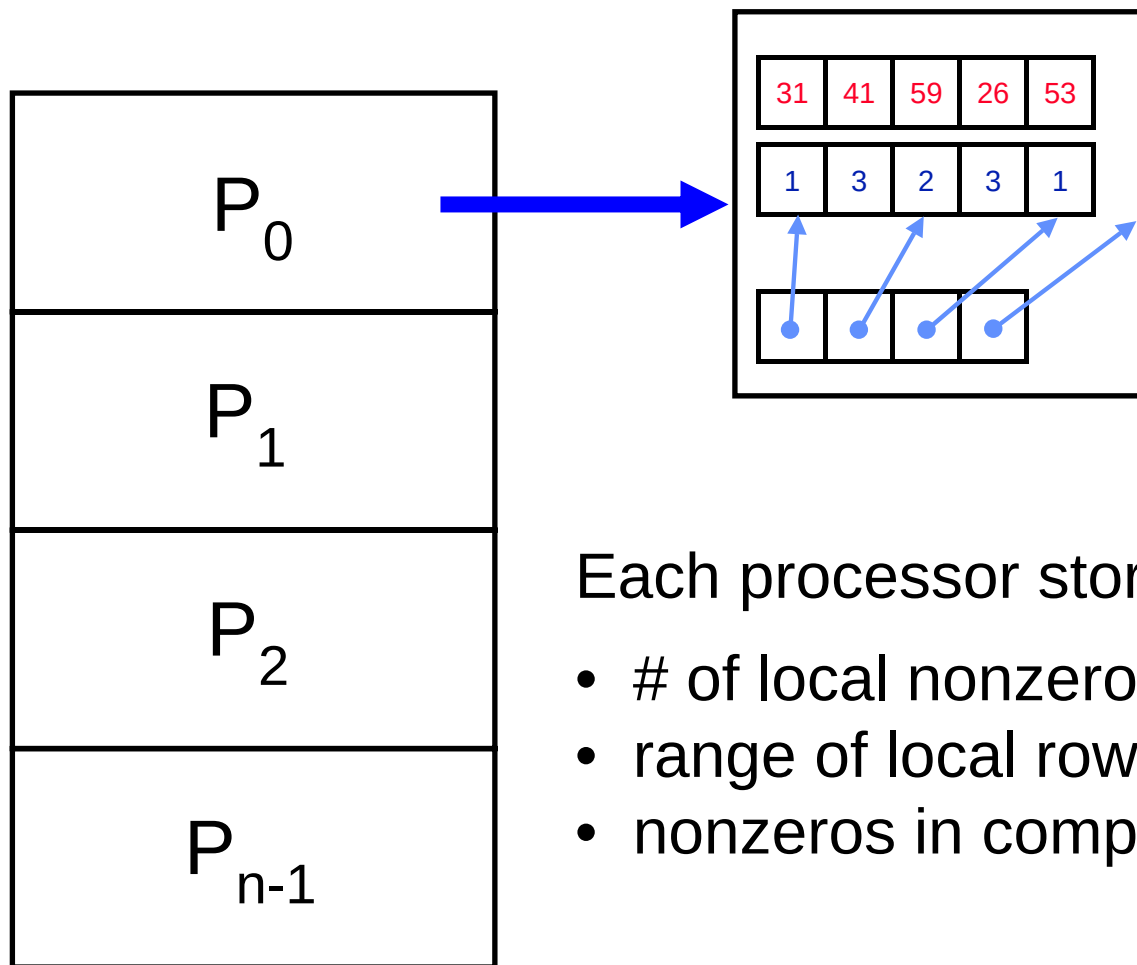


Matlab sparse matrix design principles

- All operations should give the same results for sparse and full matrices (almost all)
- Sparse matrices are never created automatically, but once created they propagate
- Performance is important -- but usability, simplicity, completeness, and robustness are more important
- Storage for a sparse matrix should be $O(\text{nonzeros})$
- Time for a sparse operation should be $O(\text{flops})$ (as nearly as possible)

**Star-P dsparse matrices: same principles,
but some different tradeoffs**

Distributed sparse data structure



Each processor stores:

- # of local nonzeros
- range of local rows
- nonzeros in compressed form

The sparse() constructor

- **$A = \text{sparse} (I, J, V, nr, nc)$**
- Input: ddense vectors **I, J, V** , dimensions **nr, nc**
- Output: **$A(I(k), J(k)) = V(k)$**
- Sort triples **$\langle i, j, v \rangle$** by **$\langle i, j \rangle$**
- Locally convert to compressed row indices
- Sum values with duplicate indices

Sparse array and matrix operations

- `dsparse` layout, same semantics as `ddense`
- Data distribution by rows only, for now
- Matrix arithmetic: `+`, `*`, `max`, `sum`, etc.
- Matrix indexing and concatenation
$$A(1:3, [4\ 5\ 2]) = [B(:, J)\ C];$$
- Linear solvers: `x = A \ b`; using SuperLU (MPI)
- Eigensolvers: `[V, D] = eigs(A)`; using PARPACK (MPI)

Sorting in Star-P

- `[V, perm] = sort (V)`
- Common primitive for several sparse array algorithms, including the `sparse` constructor
- Star-P uses a parallel sample sort

Sparse matrix times dense vector

- $y = A * x$
- First matvec with A caches a communication schedule
- Later matvecs with A use the cached schedule
- Communication and computation overlap

Combinatorial Scientific Computing

- Sparse matrix methods
- Knowledge discovery
- Web search and information retrieval
- Graph matching
- Machine learning
- Geometric modeling
- Computational biology
- Bioinformatics
- . . .

How will combinatorial methods be used by nonexperts?

Analogy: Matrix division in Matlab

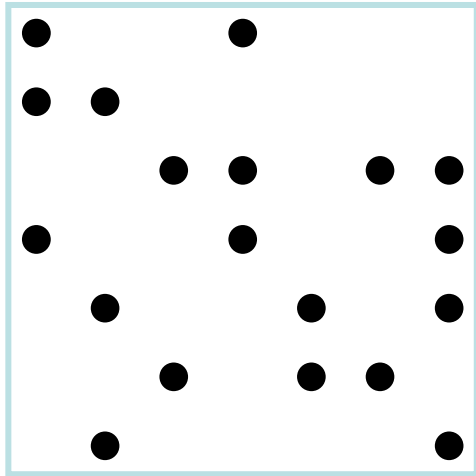
$$\mathbf{x} = \mathbf{A} \setminus \mathbf{b};$$

- Works for either full or sparse A
- Is A square?
 - no** => use QR to solve least squares problem
- Is A triangular or permuted triangular?
 - yes** => sparse triangular solve
- Is A symmetric with positive diagonal elements?
 - yes** => attempt Cholesky after symmetric minimum degree
- Otherwise
 - =>** use LU on $\mathbf{A}(:, \text{colamd}(\mathbf{A}))$

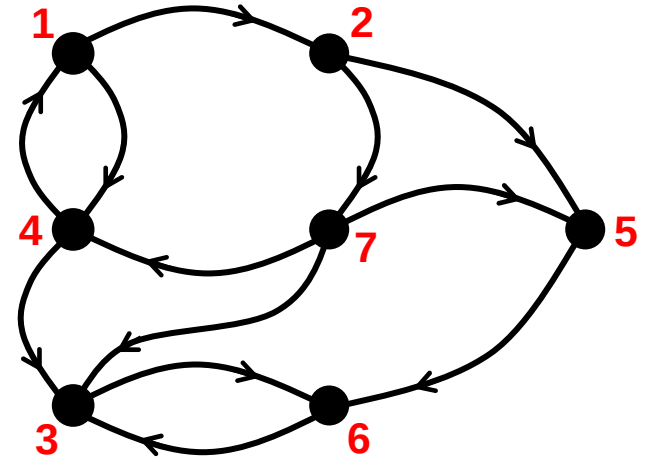
Combinatorics in Star-P

- Represent a graph as a sparse adjacency matrix
- A sparse matrix language is a good start on primitives for computing with graphs
 - Random-access indexing: **$A(i, j)$**
 - Neighbor sequencing: **$\text{find } (A(i, :))$**
 - Sparse table construction: **$\text{sparse } (I, J, V)$**
 - Breadth-first search step : **$A * v$**

Sparse adjacency matrix and graph

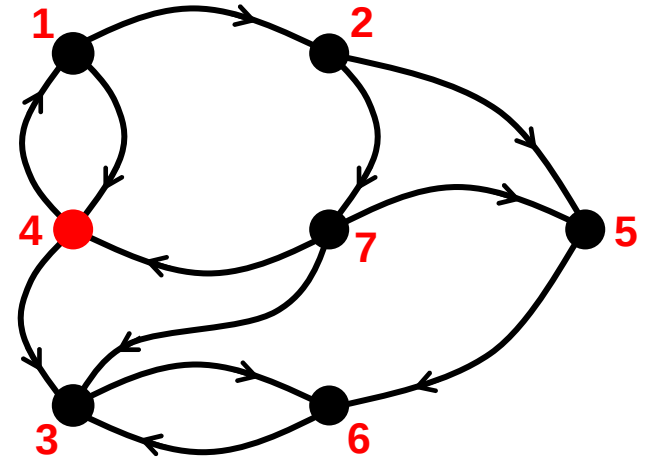
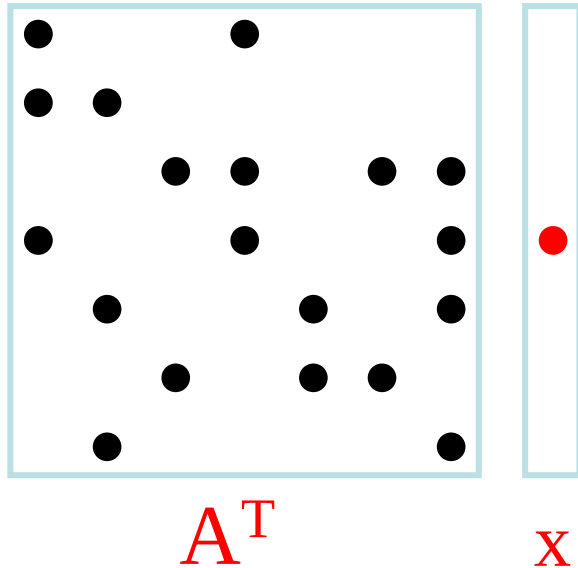


A^T



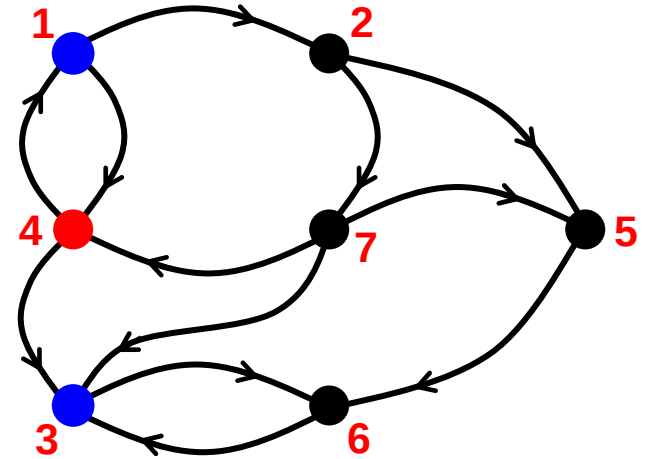
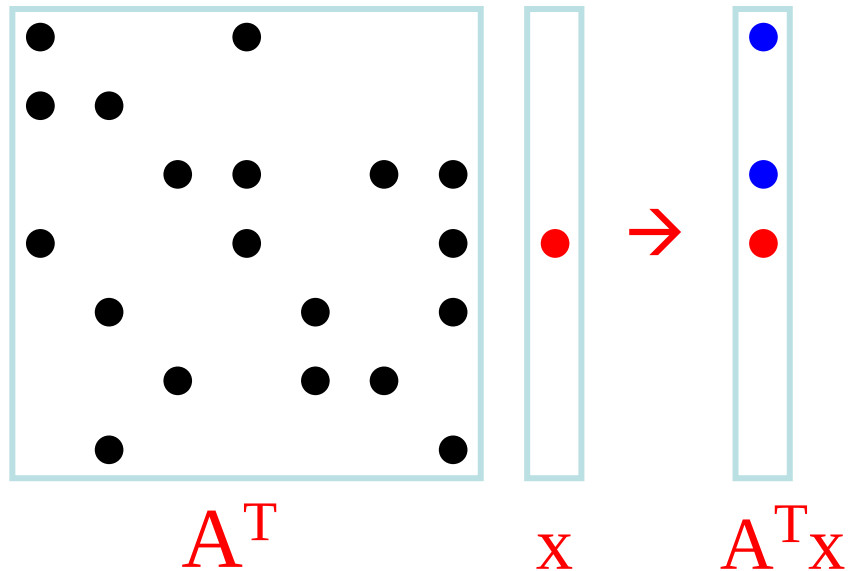
- Adjacency matrix: sparse array w/ nonzeros for graph edges
- Storage-efficient implementation from sparse data structures

Breadth-first search: sparse mat * vec



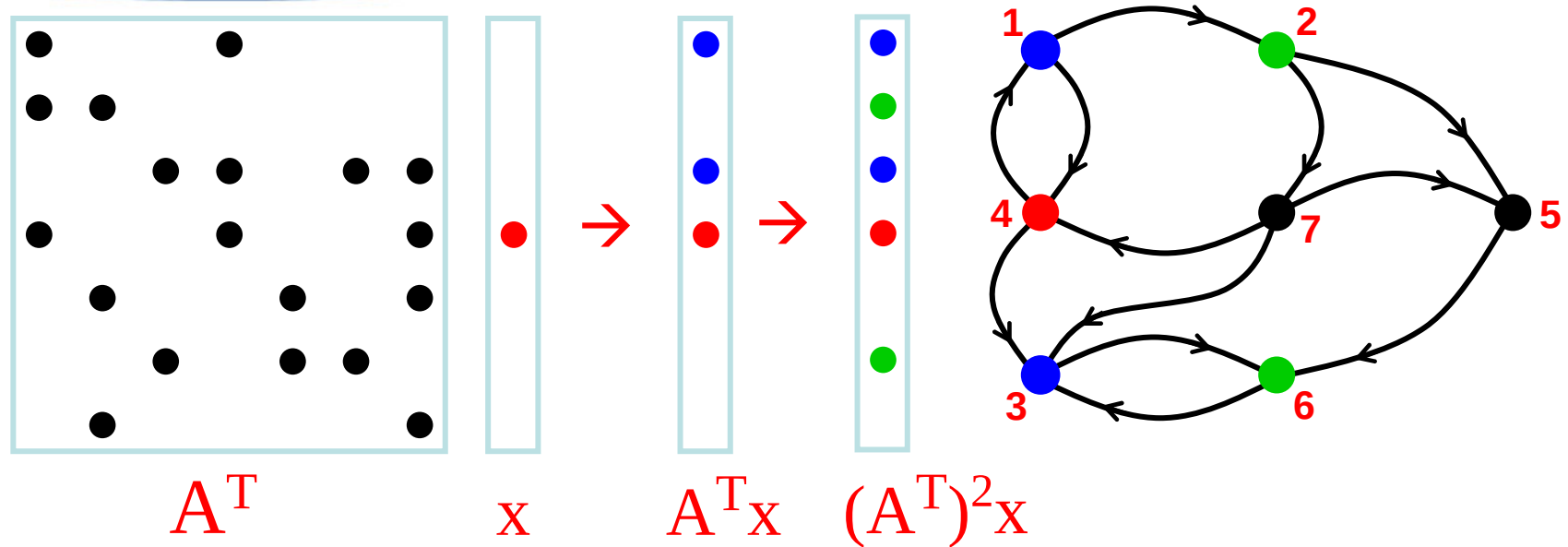
- Multiply by adjacency matrix $\rightarrow$ step to neighbor vertices
- Efficient implementation from sparse data structures

Breadth-first search: sparse mat * vec



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Breadth-first search: sparse mat * vec



- Multiply by adjacency matrix $\rightarrow$ step to neighbor vertices
- Efficient implementation from sparse data structures

Connected components of a graph

- Sequential Matlab uses depth-first search (**dmperm**), which doesn't parallelize well
- Pointer-jumping algorithms (Shiloach/Vishkin & descendants)
 - repeat
 - Link every (super)vertex to a neighbor
 - Shrink each tree to a supervertex by pointer jumping
 - until no further change
- Other coming graph kernels:
 - Shortest-path search (after Husbands, LBNL)
 - Bipartite matching (after Riedy, UCB)
 - Strongly connected components (after Pinar, LBNL)

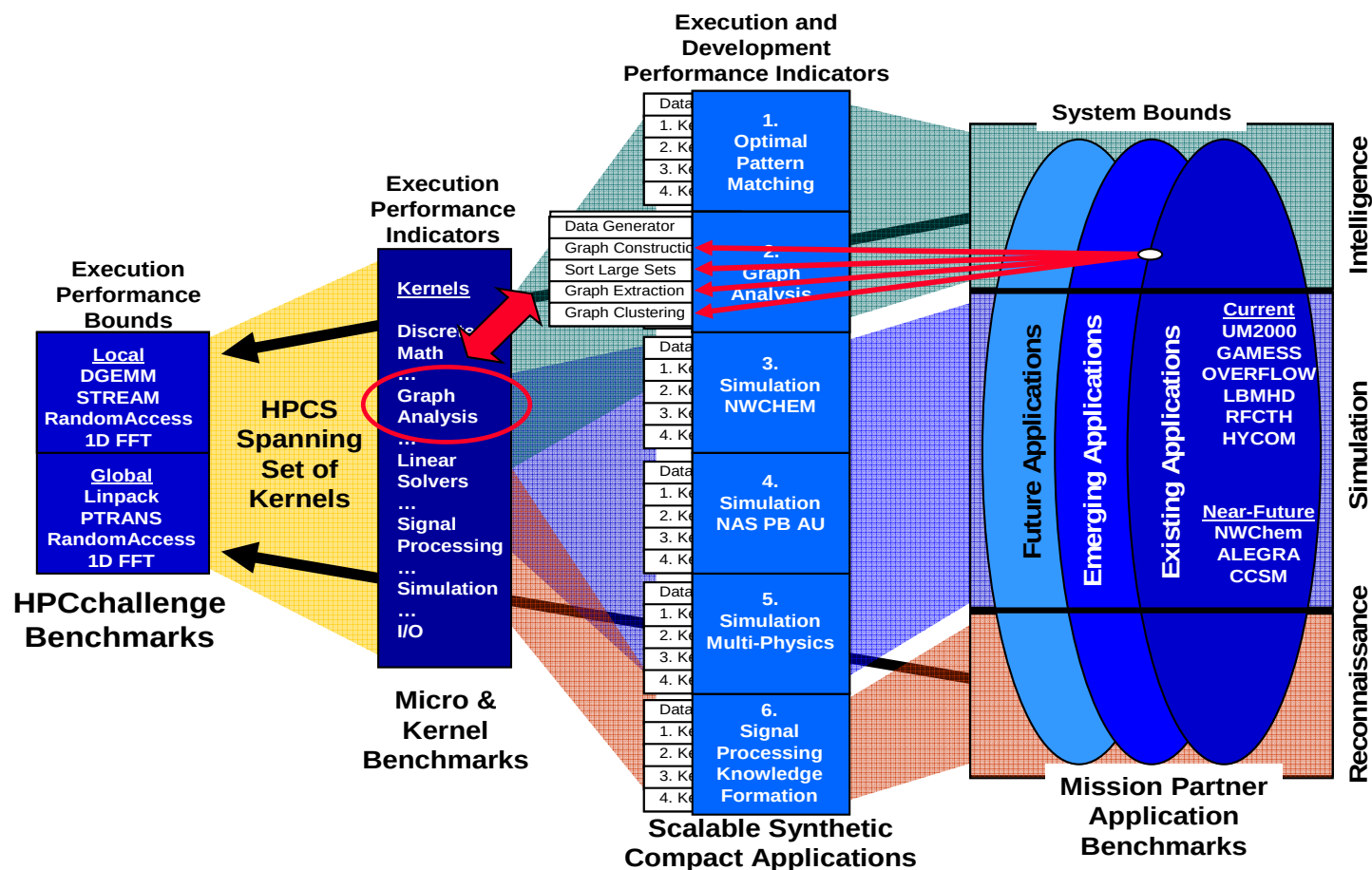
HPCS Scalable Synthetic Compact Apps



SSCA#1: Bioinformatics / Pattern Matching

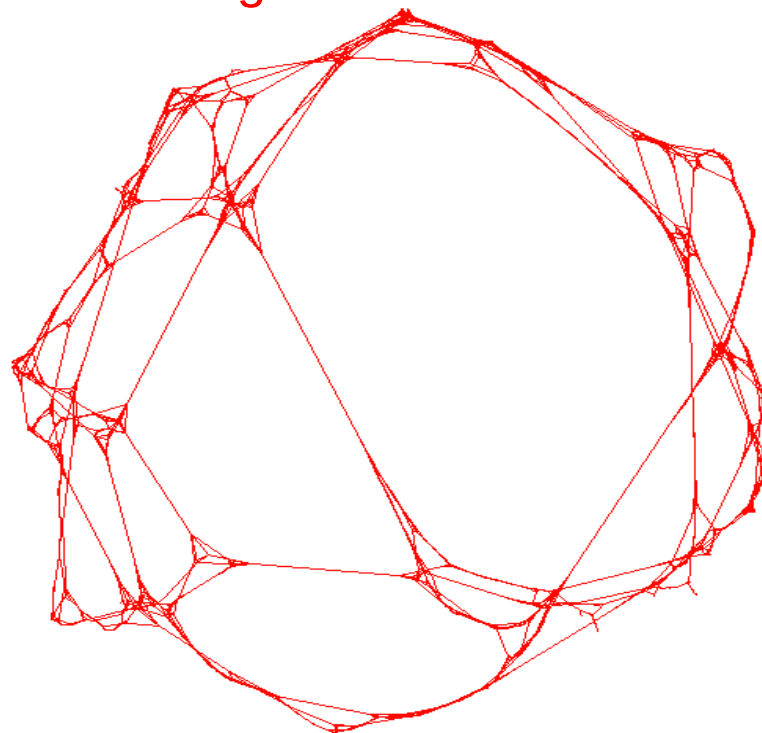
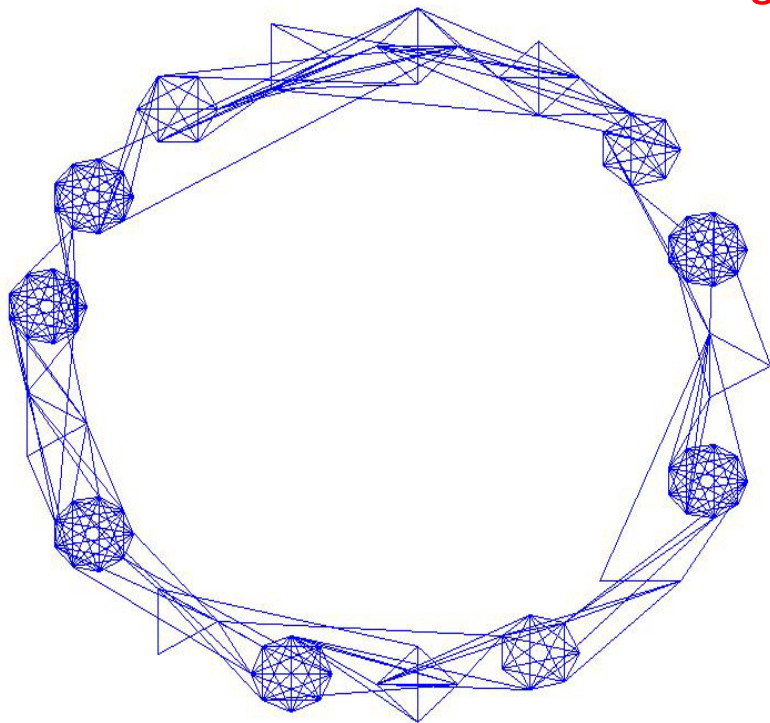
SSCA#2: Graph Analysis

SSCA#3: Sensor Processing / Imaging



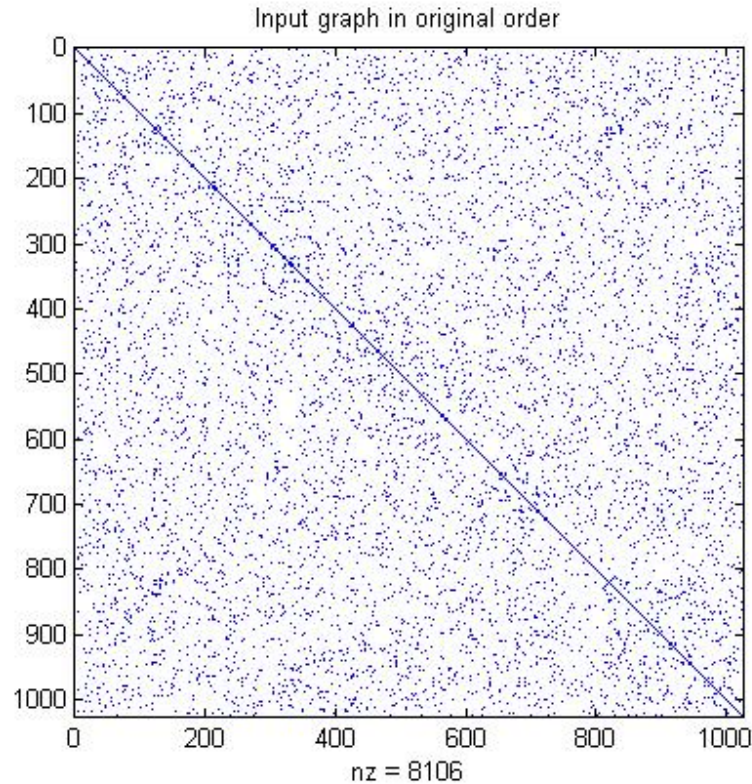
SSCA#2: “Graph Analysis”

Fine-grained, irregular data access
Searching and clustering



- Many tight clusters, loosely interconnected
- Input data is edge triples $\langle i, j, \text{label}(i,j) \rangle$
- Vertices and edges permuted randomly

Concise SSCA#2 in Star-P



Kernel 1: Construct graph data structures

- Graphs are dsparse matrices, created by `sparse( )`

Kernels 2 and 3

Kernel 2: Search by edge labels

- About 12 lines of executable Matlab or Star-P

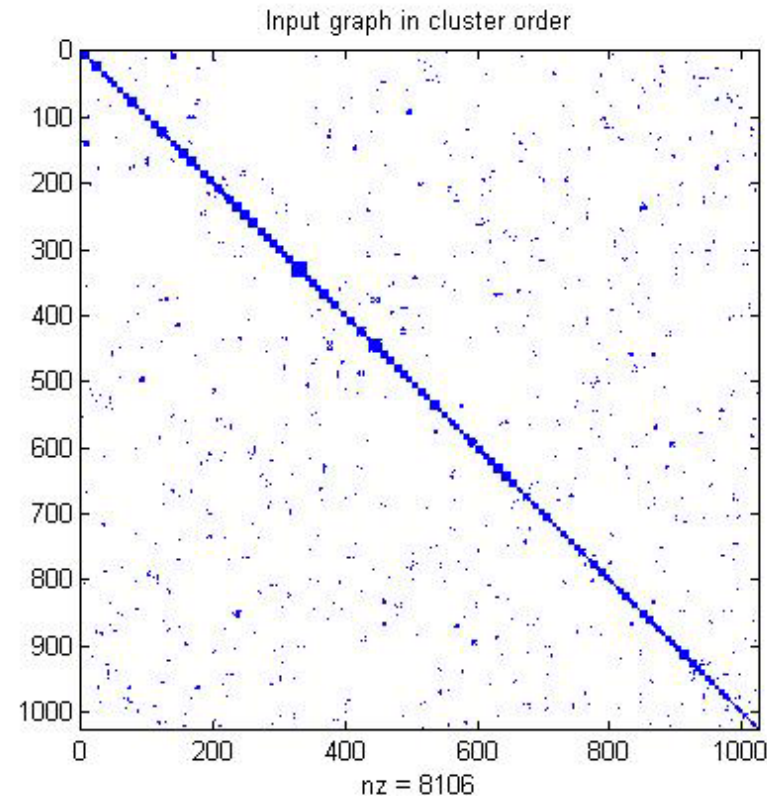
Kernel 3: Extract subgraphs

- Returns subgraphs consisting of vertices and edges within fixed distance of given starting vertices
- Sparse matrix-vector product for breadth-first search
- About 25 lines of executable Matlab or Star-P

Kernel 4: Vertex clustering

- Grow local clusters from many seeds in parallel
- Breadth-first search by sparse matrix * matrix

```
% Grow each seed to vertices  
%   reached by at least two  
%   paths of length 1 or 2  
  
C = sparse(seeds, 1:ns, 1, n, ns);  
C = A * C;  
C = C + A * C;  
C = C > 1;
```



Expressive Power: SSCA#2 Kernel 3

Star-P (25 lines)

```
A = spones(G.edgeWeights{1});
nv = max(size(A));
npar = length(G.edgeWeights);
nstarts = length(starts);
for i = 1:nstarts
    v = starts(i);
    % x will be a vector whose nonzeros
    % are the vertices reached so far
    x = zeros(nv,1);
    x(v) = 1;
    for k = 1:pathlen
        x = A*x;
        x = (x ~= 0);
    end;
    vtxmap = find(x);
    S.edgeWeights{1} = G.edgeWeights{1}(vtxmap,vtxmap);
    for j = 2:npar
        sg = G.edgeWeights{j}(vtxmap,vtxmap);
        if nnz(sg) == 0
            break;
        end;
        S.edgeWeights{j} = sg;
    end;
    S.vtxmap = vtxmap;
    subgraphs{i} = S;
end
```

MATLABmpi (91 lines)

```
declareGlobals;

% Wait for a response for each request we sent out.
for unused = 1:nunReqst
    [src tag] = probeSubgraphs(G, [P tag K3 dataReq]);
    [starts newEdges] = MPI_Recv(src, tag, P.comm);
    subg.edgeWeights{1}(:, starts) = newEdges;
    [newEdges unused] = find(newEdges);
    allNewEdges = [allNewEdges, newEdges];
end
end % of if -P paral

% Eliminate any new ends already in the all starts list.
newStarts = setdiff(allNewEdges, allStarts);
allStarts = [allStarts, newStarts];

% ENABLE_PLOT_K3DB
plotEdges(subg.edgeWeights{1}, startVertex, endVertex, k);
end % of ENABLE_PLOT_K3DB

if isempty(newStarts) % if empty we can quit early.
    break;
end
end
```

<u>Lines of code</u>	Star-P cSSCA2	MATLABmpi spec	C/Pthreads/ SIMPLE
Kernel 1	29	68	256
Kernel 2	12	44	121
Kernel 3	25	91	297
Kernel 4	44	295	241

```
% For each processor which has any of the vertices we need.
startDests = floor((newStarts - 1) / P.myV);
uniqDests = unique(startDests);
for dest = uniqDests
    starts = newStarts(startDests == dest);

    if dest == P.myRank
        newEdges = G.edgeWeights{1}(:, starts - P.myBase);
        subg.edgeWeights{1}(:, starts) = newEdges;
        [allNewEdges unused] = find(newEdges);
    end
    MPI_Send(dest, P.tag K3 dataReq, P.comm, starts);
    numReqst = numReqst + 1;
end
```

Scaling up

Early results on SGI Altix (up to 64 processors):

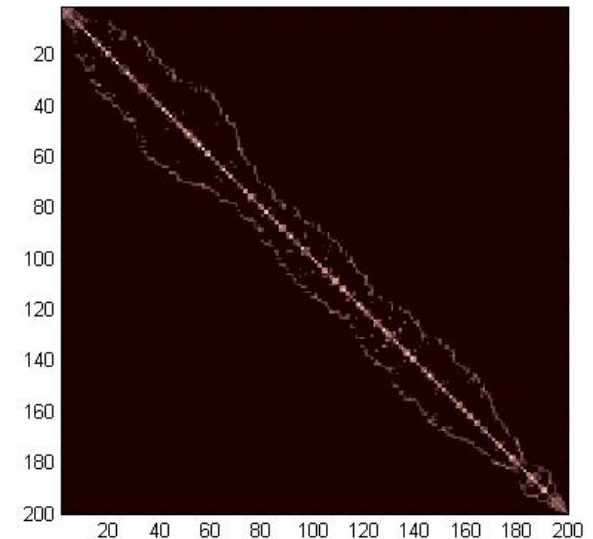
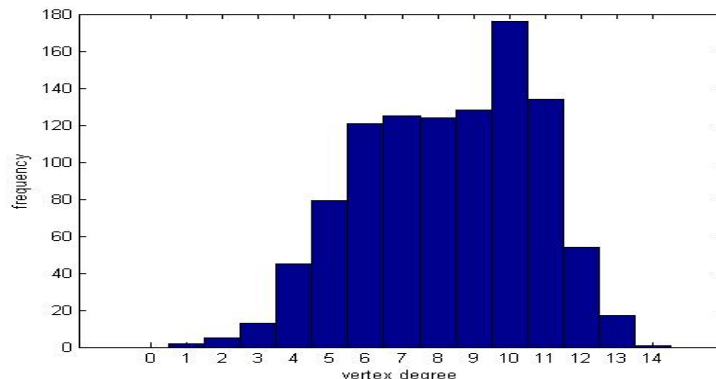
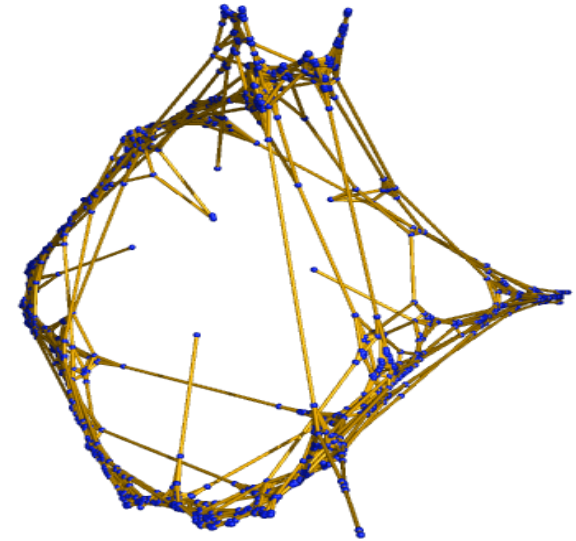
- Have run cSSCA2 on graphs with $2^{27} = 134$ million vertices
- Have manipulated graphs with 400 million vertices and 4 billion edges
- Timings scale well – for large graphs,
 - 2x problem size $\rightarrow$ 2x time
 - 2x problem size & 2x processors $\rightarrow$ same time

Using this benchmark to tune lots of infrastructure

Work in progress: Toolbox for Graph Analysis and Pattern Discovery

Layer 1: Graph Theoretic Tools

- Graph operations
- Global structure of graphs
- Graph partitioning and clustering
- Graph generators
- Visualization and graphics
- Scan and combining operations
- Utilities



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