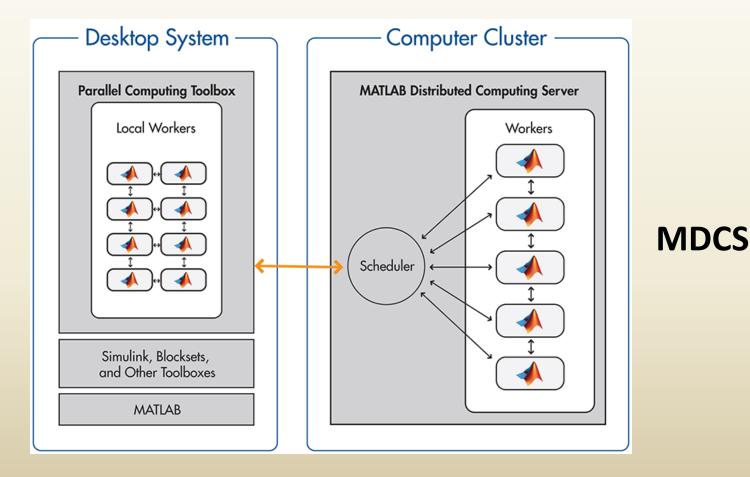
Parallel MATLAB: The Parallel Computing Toolbox, MDCS, and Red Cloud

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Seminar for the Bioinformatics Practitioners Club, Nov. 3, 2014

Overview of Parallel Computing Toolbox (PCT)

Parallel Resources: Local & Remote



PCT

PCT Opens Up Parallel Possibilities

- MATLAB does multithreading implicitly in core array ops.
- To exploit parallelism beyond this, a user needs to insert PCT commands. In order of increasing complexity:
 - Parallel for-loops: parfor
 - Single program, multiple data: spmd, pmode
 - Partitioned arrays for big-data parallelism: (co)distributed

PCT Opens Up Parallel Possibilities

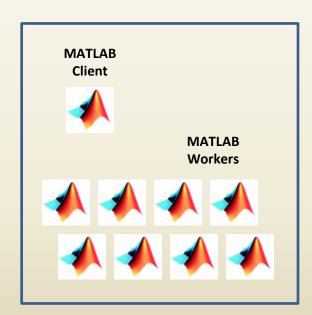
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 - Parallel for-loops: parfor
 - Single program, multiple data: spmd, pmode
 - Partitioned arrays for big-data parallelism: (co)distributed
 - Multiple batch-style runs of a serial function: createJob
 - Batch-style run of a parallel function: createParallelJob
 (= pmode), createMatlabPoolJob (if parfor/spmd sections)

Two Ways to Use PCT

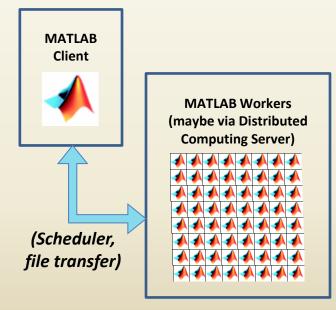
Interactively

- VS. -

batch-style



Set up matlabpool* – enter PCT commands at console



Select local pool **or** remote cluster – submit task script

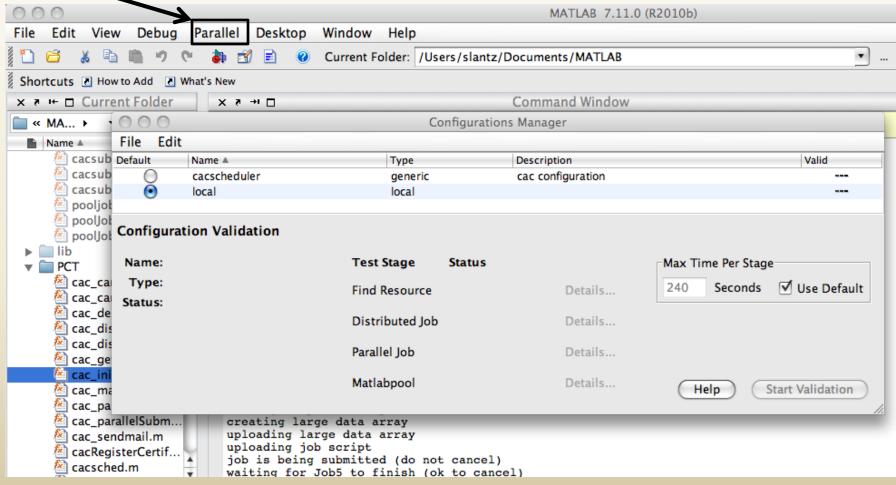
Major PCT Concepts

- matlabpool: pool of separate MATLAB processes = "labs"
 - Differs from multithreading! No shared address space
 - Ultimately allows same concepts to work on MDCS clusters
- parfor: parallel for-loop, iterations must be independent
 - Labs (workers) split up work; load balancing is built in
- spmd: single program, multiple data
 - All labs execute every command; labs can communicate
- (co)distributed: array is partitioned among workers
 - "Multiple data" to spmd, one array to MATLAB built-ins

What If You Outgrow Your Laptop?

- This is where MDCS comes in: switch to batch-style.
- PCT's interfaces allow a third party (e.g., CAC) to write implementations of PCT functions that talk to an MDCS cluster, but look the same to you as when run locally.
- Select parallel resources by using a configuration/profile, or by issuing the findResource/parcluster command.
 - Choose "local" to stay local; choose "cacscheduler" to tie
 PCT methods to CAC-specific implementations
 - You don't ever call the underlying functions directly

Using a Configuration/Profile



findResource/parcluster

 If you download the CAC client-side code, cacsched.m shows you how to call the findResource function.

```
sched = findResource('scheduler','type','generic');
```

 Examine cac_initialize.m to see how the PCT interfaces are tied to to specific functions provided by CAC.

```
set(sched, 'SubmitFcn', {@cac_distributedSubmitFcn, clusterHost, Remo
set(sched, 'ParallelSubmitFcn', {@cac_parallelSubmitFcn, clusterHost
set(sched, 'GetJobStateFcn', @cac_getJobStateFcn);
set(sched, 'CancelJobFcn', @cac_cancelJobFcn);
set(sched, 'CancelTaskFcn', @cac_cancelTaskFcn);
set(sched, 'DestroyJobFcn', @cac_destroyJobFcn);
set(sched, 'ClusterOsType', 'pc');
set(sched, 'ClusterMatlabRoot', ClusterMLRoot);
```

The Great Name Change in R2012a

<u>Previous Name</u> <u>New Name</u>

findResource parcluster

scheduler object cluster object

Configuration Profile

createJob createJob (no change)

createParallelJob createCommunicatingJob (where 'Type' = 'SPMD')

createMatlabPoolJob createCommunicatingJob (where 'Type' = 'Pool')

createTask createTask (no change)

getAllOutputArguments fetchOutputs

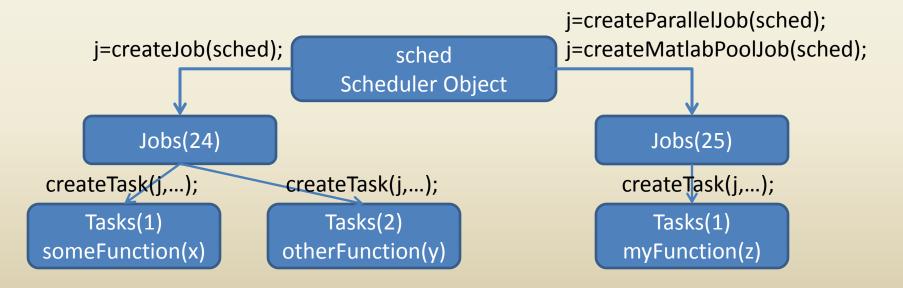
destroy delete

...etc., etc... See <u>www.mathworks.com/help/distcomp/release-notes.html</u>

Using Batch-Style PCT

Jobs and Tasks

 findResource creates a scheduler object, which allows you to create Jobs. In PCT, Jobs are containers for Tasks, which are where the actual work is defined.



Distributed Jobs

- PCT has 3 types of jobs: distributed, parallel, and pool.
- Distributed jobs have one or more tasks and no communication between tasks.
 - An MDCS scheduler runs each task as a one-core batch job
 - Useful for shifting a series of lengthy tasks to CAC, e.g.

Parallel and Pool Jobs

- Parallel and Pool jobs are multi-core or even multi-node.
 - Communication between cores/nodes must be possible.
 - The number of workers (labs) must be given.
 - These jobs have just one task!

```
job = createParallelJob(sched);
%10 core job
set(job, 'MaximumNumberOfWorkers', 10);
set(job, 'MinimumNumberOfWorkers', 10);
%All 10 cores will execute [out1] = parallelFunction(0.25,'lswt');
task=createTask(job, @parallelFunction, 1, {0.25,'lswt'});
submit(job);
```

Parallel Jobs

- All workers (labs) run the task function.
- The task function is responsible for implementing the actual parallelism using "labindex" logic.
- PCT supports MPI-style commands inside parallel jobs.

```
Size and rank are available from the start of the job.

labindex = MPI_Comm_rank+1
numlabs = MPI_Comm_size+1
Initialization is done for you
(no MPI_Init).
```

```
fprintf('I am %d', labindex);

if labindex == 1
    A = labBroadcast(1, magic(numlabs));
else
    A = labBroadcast(1);
end
```

More on Parallel Jobs

- All basic message-passing methods are available: Send, Receive, Broadcast, Barrier, gop (allreduce or allgather)
- Source and tag are the same as in MPI. MATLAB figures
 out datatypes for you.
 - labSend(data,dest,[tag]);
 - labReceive(source,tag);
 - labReceive(); % take any

```
if labindex == 1
    labSend(rand(5,5),2,5);
elseif labindex == 2
    randMatrix = labReceive('any', 5);
    labSend(randMatrix,3);
else
    randMatrix = labReceive(2);
end
```

 (Co)distributed arrays are sliced across workers so huge matrices can be operated on. Collect slices with gather.

Pool Jobs

- One worker acts as the proxy for your MATLAB client.
 This "master" runs the task function.
- The rest of the workers act as the labs in a matlabpool.
 These labs run parfor/spmd sections of the task function.

```
a(1) = gpuDeviceCount();
spmd
    b = labindex + gpuDeviceCount();
    c = numlabs;
end
for i=1:c{1}
    a(i+1) = b{i};
end % for 8 GPUs, expected answer is a = [1:8]
```

State of Jobs

 After a job is submitted, "job.state" is just one of several different ways to learn the state of the job.

```
submit(job);
%Periodically check the status (qstat -j jid)
job.state
%Block until the job is finished
waitForState(job); %==waitForState(job,'finished');
```

- waitForState is a PCT interface to block on job state,
 which can be problematic if jobs take a long time or fail.
- If more control is desired, check job.state periodically to see if the job finished or failed.

Retrieving Results

- Once your job completes, you need to get the results in two steps: (1) download files, (2) load into workspace.
- Download is only needed for MDCS jobs. It is triggered automatically by checking on job.state for a completed job, or by a blocking call to waitForState(job).
- Loading the results requires a separate function call
 - a = getAllOutputArguments(job) returns cell array a{Task,Output}
 - $a\{1,2\} = Task 1$, second output

Works the Same Everywhere!

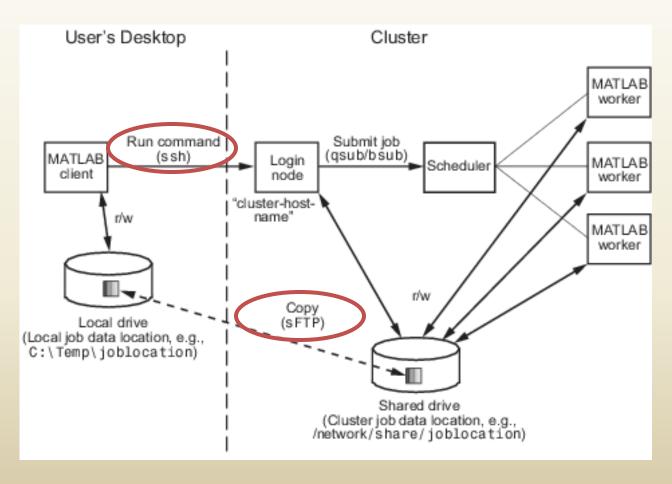
```
Command Window
  >> sched = findResource('scheduler','configuration','cacscheduler');
  >> j = createJob(sched);
  >> t = createTask(j,@rand,1,{3,3});
                                          We can control which resource is
 >> set(t,'CaptureCommandWindowOutput',1);
  >> submit(j);
                                          used to execute the job simply by
 Retrieved R2011b as current matlab version
                                          swapping out the scheduler object!
 Need assistance? Contact help@cac.cornell
                                          Command Window
                                            >> sched = findResource('scheduler','configuration','local');
                                            >> j = createJob(sched);
                                            >> t = createTask(j,@rand,1,{3,3});
  [2012-05-25 18:37:56,411] Using certificat
                                            >> set(t,'CaptureCommandWindowOutput',1);
                                            >> submit(j);
  >> wait(j);
                                            >> wait(j);
 Downloading completed job: Job227.
                                            >> a = getAllOutputArguments(j);
 >> a = getAllOutputArguments(j);
                                            >> a\{1,1\}
  >> a\{1,1\}
                                            ans =
  ans =
                                                0.9173
                                                         0.4809
                                                                   0.4626
     0.9173
               0.4809
                        0.4626
                                                0.6839
                                                         0.4612
                                                                   0.8009
     0.6839
               0.4612
                        0.8009
                                                0.8661
                                                         0.1562
                                                                   0.2155
      0.8661
               0.1562
                        0.2155
```

How to Do It Without PCT or MDCS

- Create a MATLAB .m file that takes one or more input parameters (such as the name of an input file).
- Apply the MATLAB C/C++ compiler (mcc), which converts the script to to C, then to a standalone executable.
- Run N copies of the executable on an N-core machine or a cluster, each with a different input parameter
 - mpirun can launch non-MPI processes, too
- Matlab runtimes (free!) must be available on all nodes
- For process control, write a master script in Python, say

Overview of MATLAB Distributed Computing Server (MDCS) and File Transfer

Connect to MDCS with ssh and sftp



When Is File Transfer Needed?

If you have a custom function and/or require a datafile:

```
j = createJob(sched);
createTask(j,@rand,1,{3,3});
createTask(j,@myfunction,1,{3,3});
submit(j);
waitForState(j);
a = getAllOutputArguments(j);
```

- The rand function is no problem at all, it's built in, but myfunction.m does not exist on the remote computer.
- Transfer this file and get it added to the path.

MATLAB Can Copy the Files...

- Setting the FileDependencies property tells MATLAB to copy the files for you.
- Specify the directories and files the task will need. All files and directory structure will be copied.
- Not very efficient, though: file transfer occurs separately for each worker running a task for that particular job.

```
>> set(j,'FileDependencies',{'/home/username/src/myfunction.m',... '/home/username/data/dfile.mat');
```

...Or Copy the Files Yourself

- *FileDependencies* is best for smaller projects with only a couple of files.
- Alternative for larger files:
 - 1. Copy the file(s) using sftp, or GridFTP
 - 2. Add the path to the worker sessions
- PathDependencies is used to make the task function available at run time.
 - >> set(j,'PathDependencies',{' \\matlabstorage01\matlab\username'});

Remote File Storage at CAC

- Subscriptions have 50GB of storage space
 - Intended for MATLAB scripts, job data, etc.
 - Accessible to all MATLAB jobs run by the same user
 - Can be expanded by adding extra storage to a subscription
- General access is provided through GridFTP

```
>> help gridFTP
>> ftp = gridFTP();
>> ftp.list(");
```

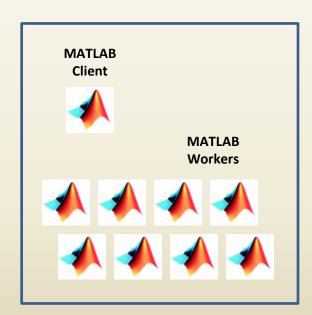
Red Cloud

Two Ways to Use PCT

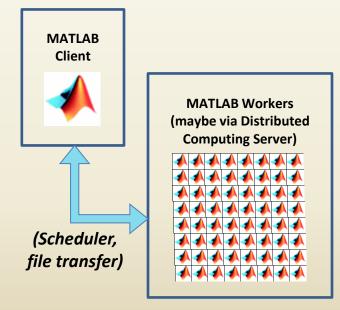
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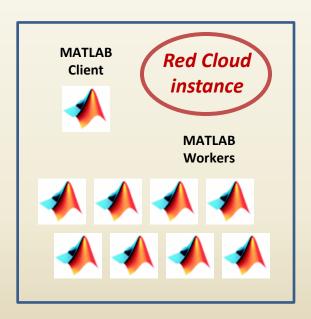
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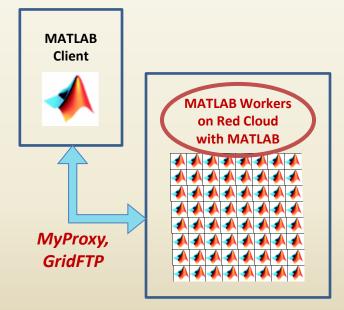
Two Ways to Use PCT at CAC

Red Cloud



Log in to an instance based on an image with MATLAB

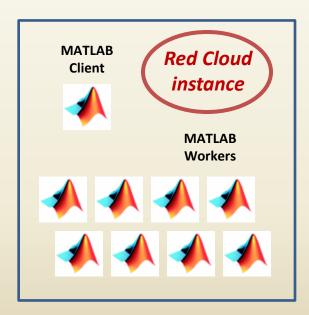
Red Cloud with MATLAB



Select CAC as your remote cluster – submit task script

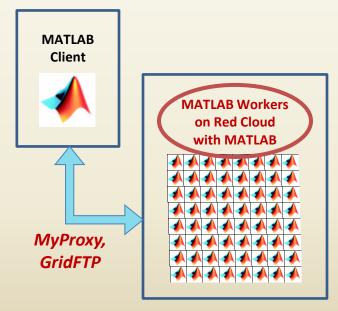
Two Ways to Use PCT at CAC

Infrastructure as a Service



Log in to an instance based on an image with MATLAB

Software as a Service



Select CAC as your remote cluster – submit task script

Red Cloud with MATLAB

- This "Software-as-a-Service" (SaaS) enables a broad research community to run MATLAB on CAC's highperformance resources in a secure, useable manner.
- Both hardware and software components make up the system. They integrate with the end user's MATLAB client at different levels.
- All functions are provided by various "services", meaning you never actually log on to any CAC systems. The client software simply makes requests to CAC systems.

Current System Specifications

- Microsoft Windows HPC Server 2008 cluster
 - Supports MATLAB clients on Windows, Mac, and Linux
 - Releases R2010b, R2011a, R2011b, R2012a, and R2013a
- 64 Intel cores in 8 Dell C6100 blade servers
 - Per server: 2 4-core Xeon E5620s @ 2.4 GHz
 - In Dell C410x: 8 NVIDIA Tesla M2070s, 1 Tflop/s, 6 GB each
 - 8 GPU-linked cores have 10GB RAM each; others have 2GB
- 8TB DataDirect Networks storage: RAID-6, error correction
 - Accessible by all servers and externally at 10 Gb/s

Services and Security

File transfer service

 Move files through a GridFTP (specialized FTP) server to a network file system that is mounted on all compute nodes

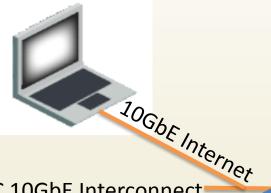
Job submission service

 Submit and query jobs on the cluster (via TLS/SSL); jobs are executed by MATLAB workers on the compute nodes

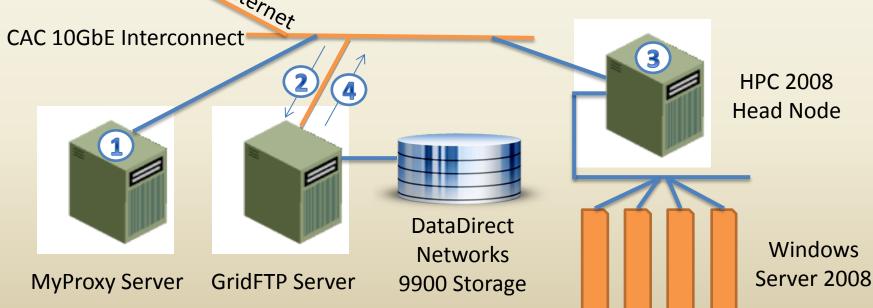
Security and credentials

 Send username/password over a TLS encrypted channel to MyProxy; get a short-lived X.509 certificate granting access

Hardware View

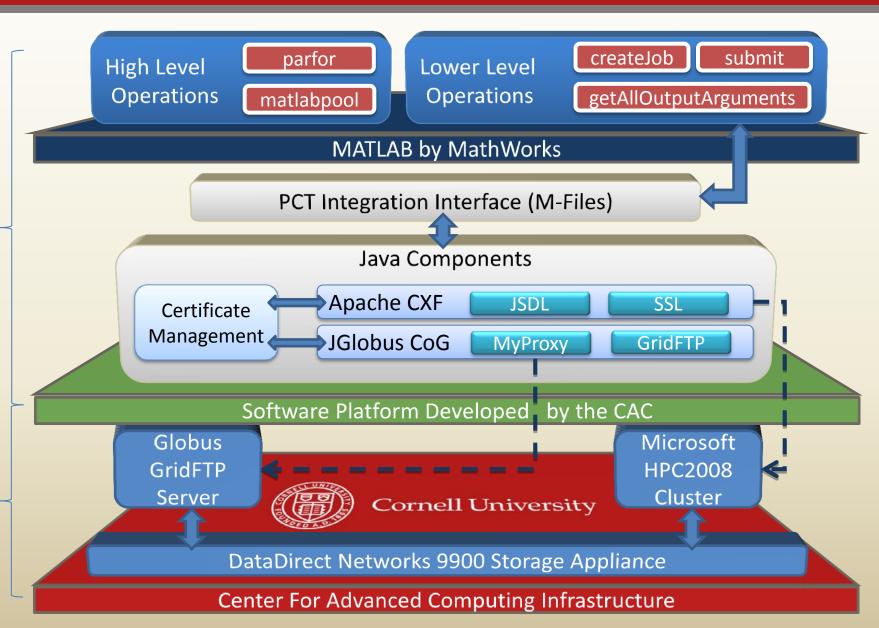


- 1. Retrieve certificate
- 2. Upload files to storage via GridFTP
- 3. Submit job to run MATLAB workers on cluster
- 4. Download files via GridFTP



Software View

- File movement and job submission interactions are largely hidden by software integrated with MATLAB.
- CAC's client code for MATLAB is a mix of Java and M-files that enable access to the HPC cluster directly from your MATLAB client through the PCT "generic scheduler" interface.
- Client code communicates as needed with server-side software at CAC to run distributed and parallel jobs on the HPC cluster's 64 CPU cores and 8 GPUs.



A Note on the Platform

- The compute nodes that run your MATLAB jobs are running Windows HPC 2008 (64 bit).
 - Your client need not be running on a Win64 platform.
 - Files requiring compilation might need to be recompiled on the HPC cluster; a utility is provided for mex files, e.g.
 - MATLAB is resilient to paths with the wrong direction of slashes, but the difference can cause problems.
 - C:\Users\naw47\myfiles\this.dat
 Windows path
 - /home/naw47/myfiles/this.dat
 Mac, Linux path

Support

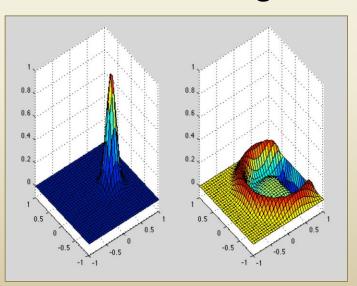
- A subscription comes with basic support to help you get started (contact help@cac.cornell.edu).
 - The basic rate allows CAC to recover hardware and software maintenance costs.
- You have the option to add more extensive consulting support to your subscription.
 - Troubleshooting
 - Guidance on optimizing your application
 - General help with parallel MATLAB

Case Study: GPGPU and MATLAB PCT

A Word About GPUs

- Red Cloud with MATLAB features 8 nodes with dedicated NVDIA Tesla M2070 GPUs capable of 1 Tflop/s each!
- MATLAB PCT has built-in GPU functions that provide an easy way to program the GPUs without learning CUDA

Stop by after the lecture to see a demo of how to run a wave simulator on Red Cloud's NVIDIA GPUs

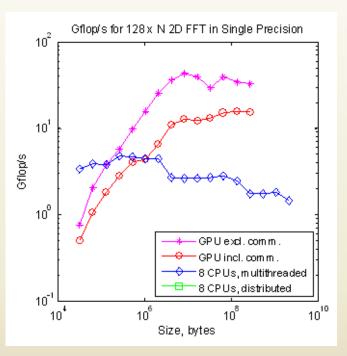


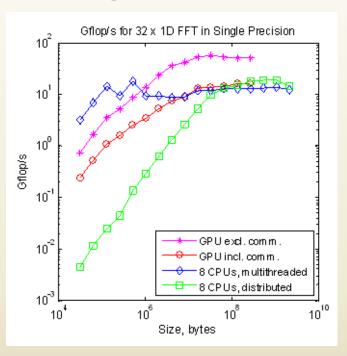
GPGPU in MATLAB: Fast and Easy

- Initial benchmarking with large 1D and 2D FFTs shows excellent acceleration on 1 GPU vs. 8 CPU cores
 - Including communication: up to 10x speedup
 - Excluding communication: up to 20x speedup
- MATLAB code changes are trivial
 - Move data to GPU by declaring a gpuArray
 - Methods are overloaded to use internal CUDA code on gpuArrays
 gpuArrays
 g = gpuArray(r);

```
f = fft2(g);
```

GPU Excels at Large FFTs





- 2D FFT > 8 MB can be 9x faster on GPU (including data transfers), but array of 1D FFTs is equally fast on 8 cores
- Limited to 256 MB due to bug in cuFFT 3.1; fixed in 3.2

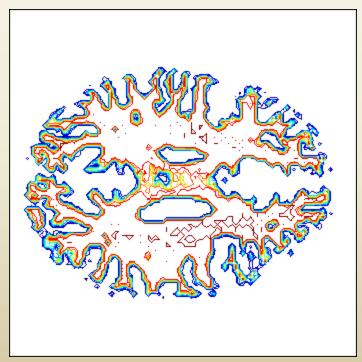
Analysis of MRI Brain Scans

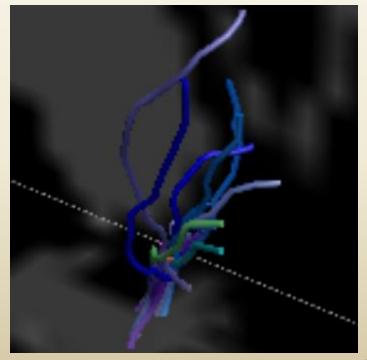
- Work by Ashish Raj and Miloš Ivković, Weill-Cornell Medical College
- Research question: Given two different regions of the human brain, how interconnected are they?
- Potential impact of this technology:
 - Study of normal brain function
 - Understanding medical conditions that damage brain connections, such as multiple sclerosis, Alzheimer's, TBI
 - Surgical planning

Connecting Two Types of MRI Data

• 3D MRI scans to map the brain's white matter

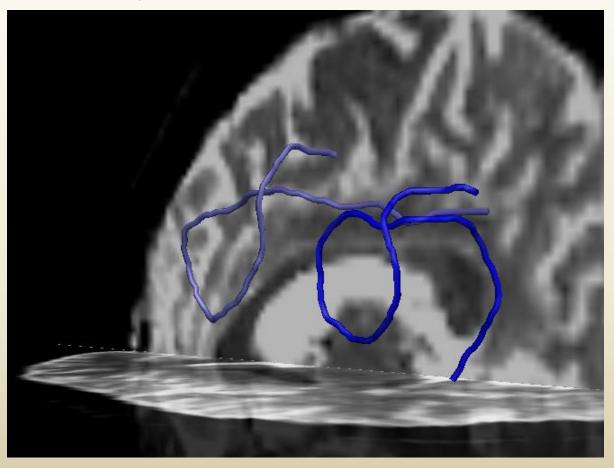
 Fiber tracts to show lines of preferential diffusion





Need for Computational Power

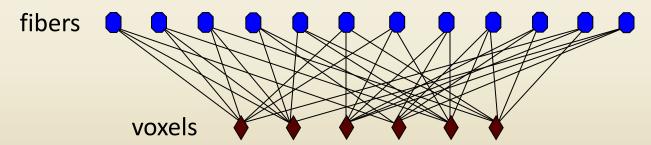
- Problem: long, spurious fibers arise in firstpass analysis
- Solution: use MATLAB to reweight fibers according to importance in connections



Examples of improbable fibers eliminated by analysis

Connections in a Bipartite Graph

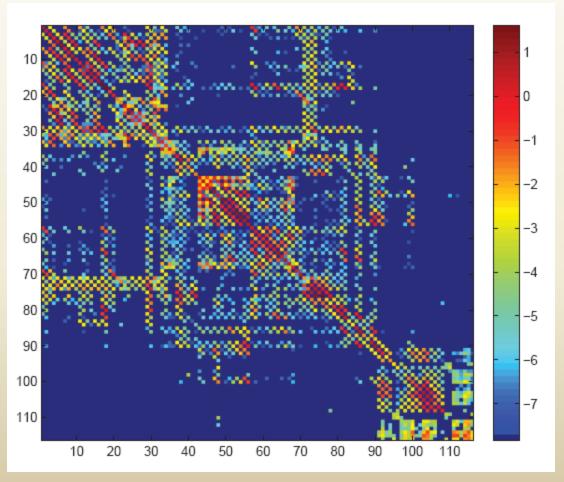
- Ivković and Raj (2010) developed a message-passing optimization procedure to solve the weighting problem
- Operates on a bipartite graph: nodes = fibers + voxels,
 edge weights = connection strength

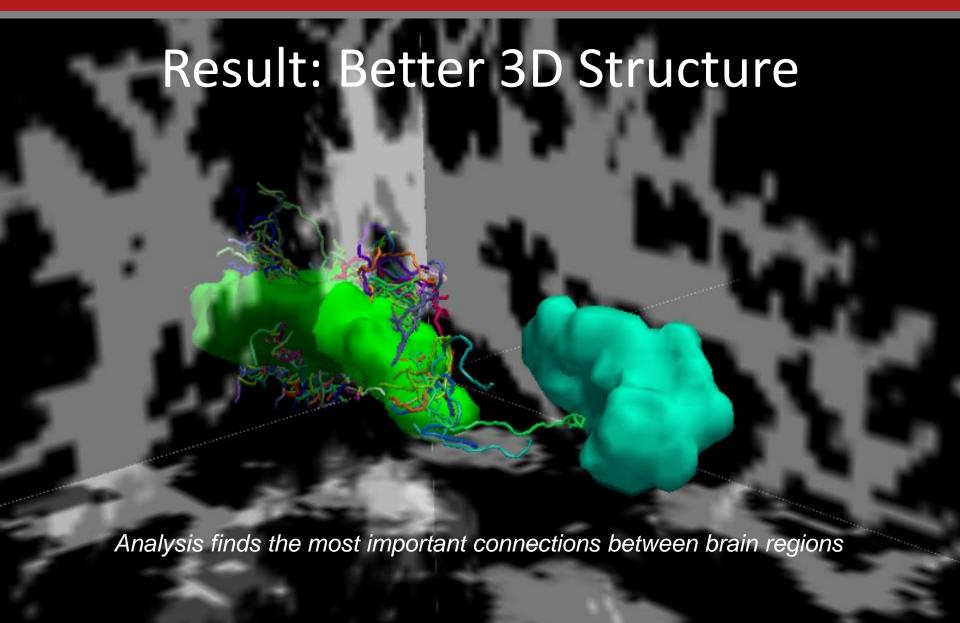


 MATLAB computations at each voxel are independent of all other voxels, likewise for fibers; inherently parallel

Data Product: Connectivity Matrix

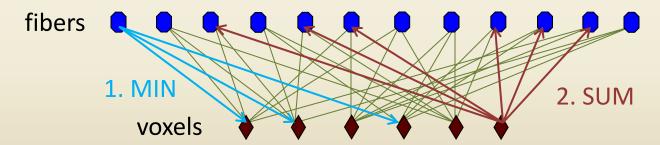
- Graph with 360K nodes, 1.8M edges, optimized in 1K iterations
- The reduced digraph at right is based on 116 regions of interest





Message-Passing Algorithm

- Iterative procedure also known as "min-sum"
- Fiber-centric step: for each fiber, find the minimum of all its edge weights; reset the edges to that value (or to the second smallest value, if already at min)



 Voxel-centric step: for each voxel, sum up its current edge weights; distribute WM value back proportionately

Round One: Parallelization

- Min/sum can be done independently for each fiber/voxel
- Loops can be converted into parfor-loops
 - On 8 cores: 375 sec/iteration shrinks to 136 sec/iteration
 - After pre-packing the WM data structure to eliminate voxels not traversed by at least one fiber: 42 sec/iteration
 - After eliminating redundant searches via improvements to indexing, and removing parfor: 32 sec/iteration!
- A better algorithm with good memory locality beats parallelization!

MATLAB Loves Matrices!

- Original code was written using structs
 - Advantage: little wasted space; handles variable-length lists of edges connected to a voxel (1–274) or fiber (2–50)
 - Disadvantage: poor data locality, because structs hold lots of extraneous info about voxels/fibers
 - Disadvantage: unsupported on GPU in MATLAB
- Better to store data in matrices!
 - Column-wise operations are often multithreaded
 - Matrix operations are often vectorized on CPUs or GPUs

Round Two: Vectorization

- So, just throw everything into one giant matrix?
 - Problem #1: row-major ordering = bad stride
 - Problem #2: mixing of dissimilar data = poor data locality
 - Due to these problems, the initial matrix-based version of the serial min-sum algorithm ran slower, 53 sec/iteration
- Initial optimization steps were easy...
 - Make columns receive all edge weights (messages)
 - Pull out only necessary info and store in separate,
 condensed matrices

Round Three: CPU Optimization

- Tighten up memory utilization by grouping fibers and voxels according to numbers of coordinating edges
 - Different matrices for fibers that connect to 2, 3, 4... edges
 - Yields full columns in the matrix for all 2-edge fibers, etc.
- Resulting code is much more complex
 - New inner for-loops over fiber count, voxel count
 - Challenge to construct the necessary indexing
- Excellent performance in the end: 0.25 sec/iteration
- Good outcome, but days and days of work

Round Four: GPU Optimization

- Since R2011b, min and sum will work on gpuArrays!
- Go back to big, simple matrices with top-heavy columns
 - Reason 1: GPU doesn't deal well with nested for-loops
 - Reason 2: Want vectorized, SIMD ops on millions of items
- Resulting code is actually less complex
 - Keep data in a few huge arrays
- Best result (after a few tricks): 0.15 sec/iteration
 - 350x speedup over un-optimized, matrix-based version
 - 2500x speedup over initial struct-based version

Are GPUs Really That Simple?

- No. Your application must meet four important criteria.
 - 1. Nearly all required operations must be implemented natively for type GPUArray.
 - 2. The computation must be arranged so the data seldom have to leave the GPU.
 - 3. The overall working dataset must be large enough to exploit 100s of thread processors
 - 4. The overall working dataset must be small enough that it does not exceed GPU memory.

PCT and MDCS: The Bottom Line

- PCT can greatly speed up the analysis of large datasets
- GPU functionality is a good addition to the arsenal
- Yes, a learning curve must be climbed...
 - General knowledge of how to restructure code for parallel and vector computing
 - Specific knowledge of PCT functions
- But speed matters!...
 - MRI image analysis, e.g., is transformed from a research curiosity into a diagnostic tool for real-time, clinical use