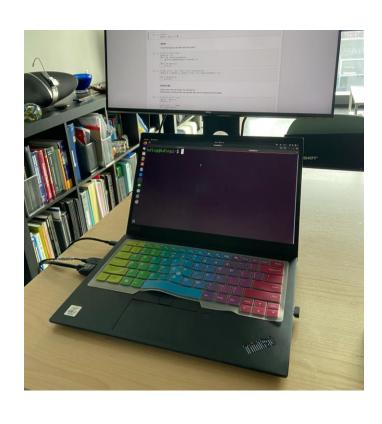
Slurm Workshop

Introduction to High Performance Computing (HPC)

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Slurm skill self-assessment

PC and HPC





Need large amount of data

Solve my problems faster

Why HPC?

To use GPU

PC cannot run my program

Connecting to HPC — Tutorial 0





ESCRIPTION

ssh (SSH client) is a program for logging into a remote machine and for executing commands on a remote machine. It is intended to provide secure encrypted communications between two untrusted hosts over an insecure network. X11 connections, arbitrary TCP ports and UNIX-domain sockets can also be forwarded over the secure channel.

ssh connects and logs into the specified destination, which may be specified as either [user@]hostname or a URI of the form ssh://[user@]hostname[:port]. The user must prove his/her identity to the remote machine using one of several methods (see below).

HPC can only be used through the internet!

HPC storage and data transfer?

- File system -- since we are sharing the cluster, we each need our separate copies of storage!
- Data mapping & viewing:
 - http hypertext transfer protocol
 - SAMBA -- re-implementation of the SMB networking protocol
- Data transfer
 - scp -- Secure copy protocol
 - Globus GridFTP, file transfer protocol

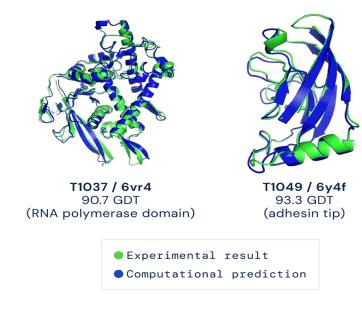
Using internet protocols to manage data transfer and data

https://rcc.uchicago.edu/docs/data-transfer/index.html#http-web-access

Software on HPC



231GB - Call of Duty: Modern Warfare



2.2TB - AlphaFold

Huge software needs to be managed differently.



Environment modules (http://modules.sourceforge.net/)

Benefit of HPC

Benefit 1: software that require large memory and large storage will be able to run on HPC.

Benefit 2: paralleled software will run much faster on HPC than on PC

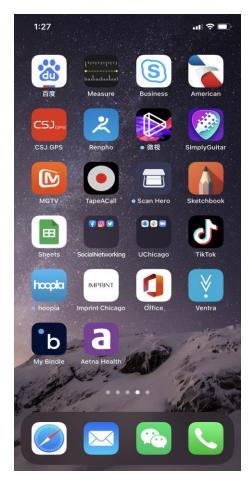
Supercomputer



More than one computer

Serial software will not magically run faster on supercomputer. The programmer need to parallel them.

HPC computation



Just click and run



- What if 1000 users are clicking on the same app?
- What if 1000 AlphaFold computations are requested at the same time?
- Whose job should start first?

HPC job scheduler – a simplified toy problem

Setup:

- One shared resource (e.g., a processor).
- Many "jobs" to do (e.g., processes).

Question: In what order should we sequence the jobs?

Assume: Each job has a:

- weight w_i ("priority")
- length l_J

HPC job scheduler – a simplified toy problem

Definition: The completion time C_j of job j = Sum of job lengths up to and including j.

```
Example: 3 jobs, l_1 = 1, l_2 = 2, l_3 = 3.
```

Schedule:

Question: What is C_1 , C_2 , C_3 ? 1, 3, 6

HPC job scheduler – a simplified toy problem

Question: What if $w_i > w_j$ but $l_i > l_j$?

Idea: Assign "scores" to jobs that are:

- inscreasing in weight
- decreasing in length

HPC batch job scheduling is NP-complete

- https://www.mendeley.com/catalogue/9ed825e9-8bae-303d-b7bc-84e0f4051411/
- Clay Mathematics Institute
- https://www.claymath.org/millennium-problems

HPC job scheduler -- SLURM

- Slurm stands for "simple linux utility for resource management"
- Slurm fair tree algorithm
 - https://slurm.schedmd.com/SLUG19/Priority_and_Fair_Trees.pdf
- Slurm documentation
 - https://slurm.schedmd.com/documentation.html

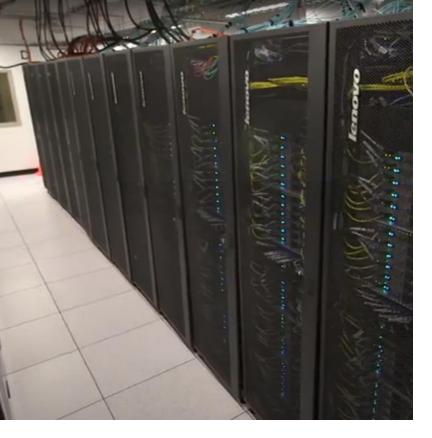
HPC job scheduler -- Slurm

- Slurm is just a software performing scheduling!
- Over 60% of TOP500 user Slurm
 - https://en.wikipedia.org/wiki/Slurm_Workload_Manager
- Job scheduler (Slurm) will be here to stay and running jobs on clusters cannot be fully automated
 - Halting problems is undecidable
 - https://en.wikipedia.org/wiki/Halting_problem
 - NP-hard, even harder than NP-complete

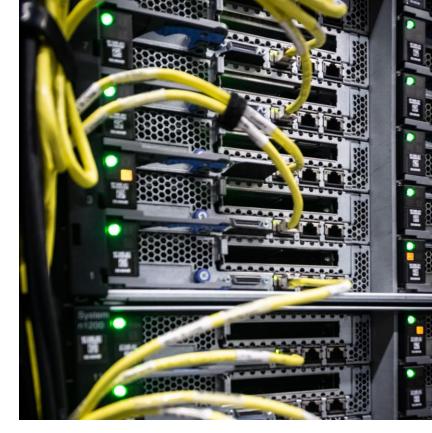
First poll: what is Slurm?



5-minute break

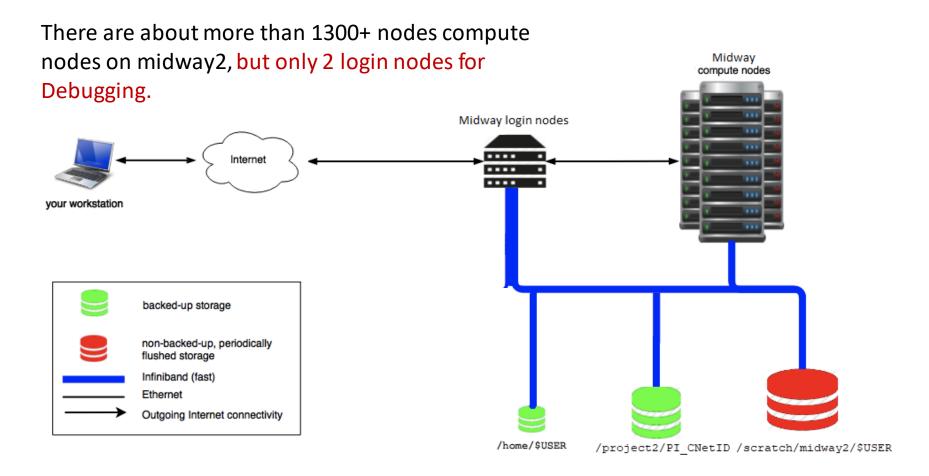






- . Midway is a constellation a of many compute systems and storage with various architectures coupled together in one system.
- . SLURM is the software used to manage the workload on Midway
- https://top500.org/

Schematic of the Midway Cluster



RCC Slurm

- Standard HPC resources
- Completely free, only for UChicago affiliates
- Require faculty approval
- Managed according to SU (service units), 1 SU = 1 core x 1 hour (https://rcc.uchicago.edu/accounts-allocations/calculationsservice-units)

Q&A

Summary

- The RCC system is pretty common among Top500
- Management according to SU
- Free for all UChicago affiliates
- State-of-the art hardware

Slurm syntax

```
#!/bin/bash
# Here is a comment
#SBATCH --time=1:00:00
                               => Time your job is allowed to run
                               => Number of nodes to run on
#SBATCH –nodes=1
                               => Number of cores on each node to use
#SBATCH –ntasks-per-node=1
#SBATCH --mem-per-cpu=2000
                               => Memory per cpu => 2000Mb or 2Gb
#SBATCH –job-name=MyJob
                               => Name of the job.
#SBATCH -output= MyJob-%j.ou => Job output file behaves as stdout for the code.
#SBATCH -error=MyJob-%j.err
                               => Error file. behaves as stderr for the code.
module load <module name>
                               => Load any modules you need for your application
                               => run the code you want
#Run your code
```

Second poll: Slurm vs parallel computing

Tutorial 1 interactive job

.../Slurm_10142021/Turorial0_Interactive

Tutorial 2: how to submit OpenMP jobs?

```
#!/bin/bash
#Here is a comment
#SBATCH --job-name=MyJob
#SBATCH --output=MyJob-%j.out
#SBATCH --error=MyJob-%j.err
#SBATCH --time=1:00:00
#SBATCH --partition=broadwl
#SBATCH --nodes=1
#SBATCH --ntasks-per-node=10
#SBATCH --mem=16000 #Per Node
##SBATCH --mem-per-cpu=2000 #Per CPU
module load gcc/9.2.0
make -f Makefile
export OMP NUM THREADS=8
#Run your code
./norm prog
```

Slurm has some variables you can use. %j is the job number. When the job runs %j will be expanded to the job number.

Specify number of cores > 1.

OMP_NUM_THREADS is an environment variable.

.../Slurm_10142021/Tutorial2_openmp

Tutorial 3: how to submit GPU jobs?

```
#!/bin/bash
#SBATCH --time=1:00:00
#SBATCH --time=00:10:00
#SBATCH --partition=gpu2
#SBATCH --gres=gpu:1
#SBATCH --nodes=1
#SBATCH --ntasks-per-node=4
#SBATCH --mem=2000
#SBATCH --job-name=MyJob
#SBATCH --output=MyJob %j.out
#SBATCH --error=MyJob_%j.err
```

module load cuda/10.1

make -f Makefile

./deviceQ

Specify number of gpus, like gpu:1

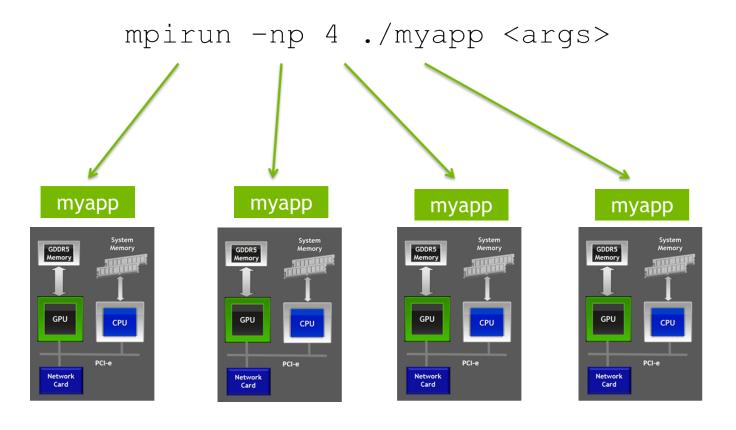
Specify partition gpu2

```
.../Slurm_10142021/Tutorial3_GPU
```

Third poll: GPU

Tutorial 4: How to submit MPI + GPU jobs?

CUDA Aware MPI



Tutorial 4: How to submit MPI + GPU jobs?

Compilation

```
#!/bin/bash
module load openmpi/3.1.2
module load cuda/10.1
# Compiling the device code
nvcc -c dev.cu
#Compiling the host code
mpicc c hostname.c
# Linking the host and device code
mpicc -o HostMap dev.o hostname.o -lcudart
#Submitting the job as batch script
sbatch mpijob.sh
```

Job Submission

```
#!/bin/bash.brown.edu/oscar/gpu-computing/mpi-cuda

#SBATCH -t 00:30:00

#SBATCH --nodes=2

#SBATCH --ntasks-per-node=4 slide Show Review

#SBATCH --partition=gpu2

#SBATCH --gres=gpu:2dd-ins | Shapes | Icons | 30 Models |

#SBATCH --job-name=MyJob

#SBATCH --output=MyJob-%j.out

#SBATCH--asse_deerror=MyJob-%j.err

#SBATCH --qos=stafftest

mpirun ./HostMap
```

Tutorial 5: How to submit array based jobs?

```
#!/bin/bash
# Job Name
#SBATCH --job-name=arrayjob
# Walltime requested
#SBATCH --time=0:10:00
#Add partition
#SBATCH --partition=broadwl-lc
# Provide index values (TASK IDs)
#SBATCH --array=1-16
# Use '%A' for array-job ID, '%J' for job ID and '%a' for task ID
#SBATCH --error=maths%A-%a.err
#SBATCH --output=maths%A-%a.out
# single core
#SBATCH --ntasks-per-node=1
#SBATCH --mem-per-cpu=2000
# Use the $SLURM ARRAY TASK ID variable to provide different inputs for each job
input=$((SLURM_ARRAY_TASK_ID*1000+2))
echo "Running job array number: "$SLURM_ARRAY_TASK_ID "input " $input
```

.../Slurm_10142021/Tutorial5_GPU



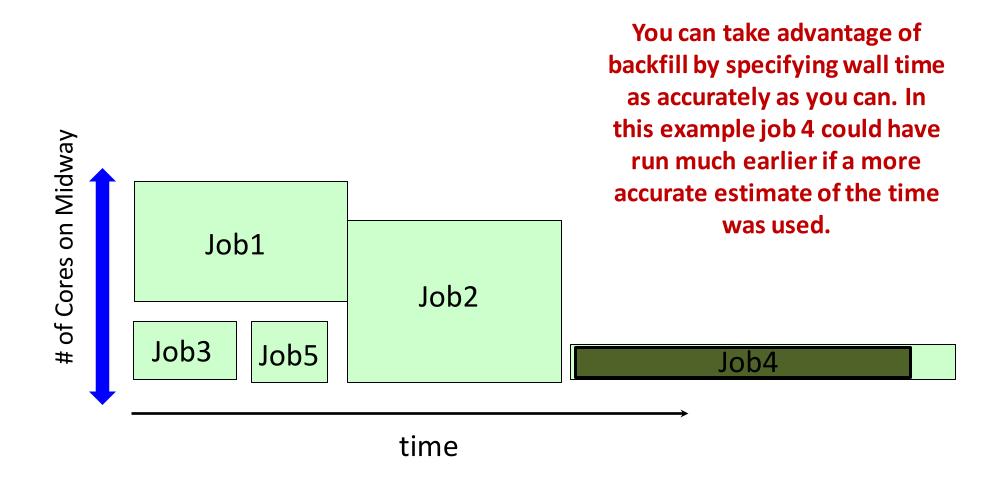
5-minute break

What if I need an entire node, large memory or specific features?

- Exclusive node --- #SBATCH —exclusive
- Useful commands
 - Jobinfo –j jobid
 - rcchelp qos
- Specific hardware -- #SBATCH -constraint=v100
 - o *nodestatus* command

Job Priority

Why has someone else's job started before mine?



Make an informed decision Slurm setups

dingkaihua@midway2-login2 cheat_sheets]\$ rcchelp qos									
Name	MaxNodes	MaxCPUs	MaxCPUsPerUser	MaxNodesPerUser	MaxJobsPerUser	MaxSubmitJobs	MaxWall	Partition	
amd bigmem bigmem2 broadwl broadwl-large broadwl-lc build cpp-staging			 112 2800 	 100 	256 100 5 100 100	 100 100 500 100	1-12:00:00 1-12:00:00 1-12:00:00 1-12:00:00 12:00:00 1-12:00:00 12:00:00	build	
cron debug gpu gpu2	5 2 	 4 	5 	5	10 1 16 10	10 1 100 100	12:00:00 00:15:00 1-12:00:00 1-12:00:00	cron broadwl westmere sandyb gpu gpu	

Qos table shows the association and limitation of each partition.

There is an optimal number of nodes / cores to request to achieve the best speedup.

- Amdahl's law
- https://en.wikipedia.org/wiki/Amdahl%27s_law
- More nodes does not lead to faster computation eventually

Job submission and monitoring

Slurm Commands

Command	Description			
sbatch script.sbatch	Submits script.sbatch job script			
squeue -u \$USER or myq	Reports the status of your jobs			
sacct -u \$USER	Displays accounting data for your job(s)			
scancel jobid	Cancels a running job or removes it from the queue			
scontrol show job jobid or jobinfo	Displays details of a running job			

Recommended online resources

- User guide on running jobs on Midway
 - https://rcc.uchicago.edu/docs/running-jobs/index.html

- Details Slurm documentation
 - https://slurm.schedmd.com/sbatch.html

- SLURM Cheat Sheet
 - https://slurm.schedmd.com/pdfs/summary.pdf

Q&A

Slurm skill self-assessment

Thank you!

Please feel free to reach out for more questions!

help@rcc.uchicago,

dingkaihua@uchicago.edu