HPC at CCR, and Intro to OnDemand

Jeanette Sperhac

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Welcome to Center for Computational Research (CCR)



We are an academic research computing center at

University at Buffalo,

State University of New York (SUNY),

Buffalo, New York, USA.

High-Performance Computing (HPC) at CCR



- CCR houses about 1600 computing nodes
- each node has up to 40 processing cores (CPUs)
- some have Graphics
 Processing Units (GPUs)

In total, CCR has about 30,000 CPUs (cores).

What makes HPC, HPC?

Overall high performance computing features:

- Fast compute
- Data storage
- Substantial memory
- Fast networking
- Specialized software

Fast compute

CCR has more than 1 PFlop/second peak performance compute capacity

petaflop/s = one quadrillion floating-point operations per second

- processor density (up to 40 cores/node)
- Iots of memory (up to 800 GB/node)
- specialized hardware (think GPUs)
- specialized architectures (tuned to scientific problems)

Lots of data storage



3+ PB high-performance parallel filesystem

(recall: $1 PB = 2^{50} bytes = 1024$ terabytes = one million gigabytes)

High performance networks



- fast connections to data storage
- fast interconnects between compute nodes

About SLURM



SLURM (Simple Linux Utility for Resource Management) is batch scheduling software. It's the gatekeeper on the CCR computing resource.

You tell it about your job's requirements:

how many cores?

- how much memory?
- for what duration?

(Yes, the name is a Futurama reference!)

About SLURM II



We must also tell SLURM where to run our job:

- under whose account?
- on which cluster?
- on which partition?

And SLURM schedules your job.

(Yes, the name is a Futurama reference!)

CCR has three computing clusters:

- general-compute
- industry
- faculty (that's us!)

Faculty cluster

On the faculty cluster, we will use the valhalla partition and these parameters:

cluster=faculty account=cyberwksp21 partition=valhalla

qos=valhalla

See for yourself! On the command line, type: slimits

Different ways to run

We can tell SLURM to run:

sbatch:

schedule a "batch" job when possible

salloc/srun:

schedule the resources with salloc, run interactively with srun

See salloc and srun in action:

salloc demo

Monitoring: So, what's cooking on the cluster?

What may I access on the cluster?

slimits

Show me the faculty cluster status:

sqstat --faculty

Show me valhalla's allocated nodes:

snodes all faculty/valhalla | grep alloc

output:

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About OnDemand at CCR

During the workshop, we will use a web browser and OnDemand to access CCR computing resources. In some cases the negotiation with SLURM happens behind the scenes.

https://ondemand.ccr.buffalo.edu

We will use OnDemand three ways:

Jupyter Notebooks

notebooks run on a valhalla compute node

Faculty Cluster App

command line access to a valhalla compute node

Faculty Shell

command line access to CCR front-end node, vortex

OnDemand at CCR

This is where we begin: https://ondemand.ccr.buffalo.edu



OnDemand Jupyter Notebooks

Schedule a SLURM job that runs a Jupyter session on a valhalla compute node:

Interactive Apps -> Jupyter Notebook Quantum Chemistry

 Start the Interactive App:



OnDemand Jupyter Notebooks

Schedule a SLURM job that runs a Jupyter session on a valhalla compute node:

Interactive Apps -> Jupyter Notebook Quantum Chemistry

2. Configure the session(SpecifySLURMparameters):



OnDemand Jupyter Notebooks

Schedule a SLURM job that runs a Jupyter session on a valhalla compute node:

Interactive Apps -> Jupyter Notebook Quantum Chemistry

3. Run Jupyter:

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Imput_tmp.in 8 days ago	0 B
□ □ job-8528567.tar.gz 3 years ago	426 B
junk-activate-libra-env.sh 3 months ago	430 B
monitorSLURM.py a year ago	21.4 kB
Rplots.pdf 3 years ago	3.61 kB
this-file.txt a day ago	93 B
C trythis.txt a day ago	38 B
yum-list-installed.txt 3 years ago	86.1 kB

OnDemand Faculty Cluster App

Schedule a SLURM job that runs a Linux desktop on a valhalla compute node:

Interactive Apps -> Faculty Cluster Desktop - Advanced Options

Share the cores!

 Start the Interactive App:

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OnDemand Faculty Cluster App

Schedule a SLURM job that runs a Linux desktop on a valhalla compute node:

Interactive Apps -> Faculty Cluster Desktop - Advanced Options

Share the cores!

Specify
 SLURM
 parameters:



OnDemand Faculty Cluster App

Schedule a SLURM job that runs a Linux desktop on a valhalla compute node:

Interactive Apps -> Faculty Cluster Desktop - Advanced Options

Share the cores!

3. Run Cluster Desktop:



OnDemand Faculty Shell

Run a command line shell on CCR's front-end node, vortex:

Clusters -> Faculty Cluster Shell Access

Note: It's not a running job, just a shell. Use the shell to run a SLURM script, or a modest test (few minutes' duration, low memory requirements).

1. Start the shell:

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2. Run the shell:

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View your active jobs

View jobs you are running right now:

Jobs -> Active Jobs

1. View in OnDemand:



2. Or run squeue -u \$USER in a terminal:

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Troubleshoot and debug

View your OnDemand sessions (Click on the box icon, My Interactive Sessions):



Troubleshoot and debug: zoom in on one session

For each session we see useful information:

- Note the hostname where the job is running, cpn-p27-15
- You can click Session ID to access session log files



Files app



Use the OnDemand Files app (e.g. Files -> Home Directory) for:

- Browsing directories
- Upload files
- Download files
- Simple file edits*

* Potentially

Editing your files

You have multiple options for file editor:

- OnDemand Files app* (simplest)
- nano (easy)
- vi (just right)
- emacs (grrr)
 - * Potentially

Contact us

Have a question, comment, or issue?

- Join us on the workshop Slack channel: quantumdynamicshub.slack.com
- Check the CCR documentation
- Enter a CCR help ticket:
 email: ccr-help@buffalo.edu
 webpage: https://ubccr.freshdesk.com/