# Beginner's Introduction to Computing at CARC

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Version 0.2

- •1) Basic Linux Literacy
- 2) Ability to run programs on CARC compute nodes

I hope to spend about an hour on each with a 15 minute break.

## Outline

- High Performance Computing Overview
- Logging in
- The BASH Shell and Scripts
- The Slurm Job Scheduler
- Storage at CARC
- Transferring data to and from CARC
- Accessing software and the module system
- Parallelization

## High Performance Computing

- What is high performance computing?
  - Really just means something that is a lot more powerful than your desktop or laptop.
  - Hardware:
    - That might mean more and faster processors to do the calculations more quickly (eg 400 CPUs instead of 4)
    - More RAM so you can work on bigger problems (3,000 GB instead of 8)
    - Bigger file systems so you can process larger datasets
    - More and bigger GPUs to accelerate your computations (12 GPUs at a time instead of 1)
  - People:
    - Someone else to manage the systems and keep them running and secure
    - Someone to answer your questions and help with problems



First login to the Linux **workstation** in front of you.

Use your CARC username and password.

Raise your hand if you have any trouble.

This is an "important step" so don't let me move on until you have logged in

First login to the **workstation** in front of you.

You will always login to CARC cluster remotely.

These clusters don't even have monitors.



We are going to use a program called secure shell.

Secure shell (ssh) is now built into every major operating system (Windows, OSX, and Linux).

You don't need third party programs like putty anymore.

Type to search









#### Should prompt you for a password...

Don't let me move on until you are able to login.

#### \$passwd

#### Welcome to Wheeler

Be sure to review the "Acceptable Use" guidelines posted on the CARC website.

For assistance using this system email help@carc.unm.edu.

Tutorial videos can be accessed through the CARC website: Go to http://carc.unm.edu, select the "New Users" menu and then click "Introduction to Computing at CARC".

Warning: By default home directories are world readable. Use the chmod command to restrict access.

Don't forget to acknowledge CARC in publications, dissertations, theses and presentations that use CARC computational resources:

"We would like to thank the UNM Center for Advanced Research Computing, supported in part by the National Science Foundation, for providing the research computing resources used in this work."

Please send citations to publications@carc.unm.edu.

Wheeler is our largest general purpose computational cluster.

There are two types of slurm partition on Wheeler:
1) Normal - this partition is for long running batch jobs.
2) Debug - for testing your code and interactive jobs. Short time limits so that nodes are usually available right away.

Type "qgrok" to get information about the partitions. Enter "quotas" to see your storage usage and limits. For a list of software installed on Wheeler enter "module spider"

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- \* Logging in under another person's account is strictly forbidden and will \*
- \* result in the account being locked.

Warning: Your home directory (~) usage is 195G\*/100G. Please reduce your usage. Use the ncdu command to see details.

Warning: Your personal scratch storage (/carc/scratch/users/mfricke) usage is 116.46GiB/100.00GiB)
mfricke@wheeler:~ \$

## Please enter the following command



cp -r /projects/shared/workshops/beginner/mystuff ~/

#### We will come help you if you have any trouble.

(Later I will go over what this command does)

# Linux and the BASH Shell

- The Kernel manages access to the hardware in a computer.
- An Operating System (OS) is the Kernel plus useful programs provided by the OS.
- The "shell" is the outermost layer of the OS.
- It is where the user interacts with the OS.







Microsoft Windows

## Graphical Shells (GUIs)





#### Linux and the BASH Shell



# The Borne-Again Shell (BASH)

Written in 1976 by Stephen Bourne for UNIX version 7.









#### mfricke@wheeler:~ \$

#### This is the current working directory. "~" is short for home directory





Example Filesystem Tree

• • •

# [vanilla@wheeler ~]\$ pwd /users/vanilla [vanilla@wheeler ~]\$

• • •

# [[vanilla@wheeler ~]\$ ls mystuff wheeler-scratch [vanilla@wheeler ~]\$

```
[[vanilla@wheeler ~]$ tree
    mystuff
     — myfile1
       - myfile2
    wheeler-scratch -> /wheeler/scratch/vanilla
2 directories, 2 files
```

[vanilla@wheeler ~]\$

```
[[vanilla@wheeler ~]$ tree
    mystuff
     — myfile1
       - myfile2
    wheeler-scratch -> /wheeler/scratch/vanilla
2 directories, 2 files
```

[vanilla@wheeler ~]\$

• • •

# [vanilla@wheeler ~]\$ tree This. means the current directory mystuff myfile1 myfile2 wheeler-scratch -> /wheeler/scratch/vanilla

2 directories, 2 files
[vanilla@wheeler ~]\$

#### 5 Minute Break

## "Absolute" paths vs "relative" paths

- A path is a list of directories and/or files. It is a path through the directory tree that tells one how to get somewhere in the filesystem.
- An absolute path tells one how to get to the destination from starting from the root of the filesystem. E.g "/users/vanilla/mystuff/"
- A relative path specifies how to get there \*starting from the current working directory\*. E.g vanilla/mystuff/



# [vanilla@wheeler ~]\$ ls mystuff/ myfile1 myfile2 [vanilla@wheeler ~]\$



# [vanilla@wheeler ~]\$ ls /users/vanilla/mystuff myfile1 myfile2 [vanilla@wheeler ~]\$

[vanilla@wheeler ~]\$ ls ./mystuff/ myfile1 myfile2 [vanilla@wheeler ~]\$ ls ~/mystuff/ myfile1 myfile2 [vanilla@wheeler ~]\$

[vanilla@wheeler ~]\$ ls -a mystuff .addressbook .oracle jre usage .addressbook.lu .pinerc .bashrc .pki .cache .rhosts .comsol .shosts .config .spack .flexlmrc .ssh

```
[vanilla@wheeler ~]$ ls -l
total 4
drwxr-xr-x 2 vanilla users 4096 Jun 14 22:05 mystuff
lrwxrwxrwx 1 vanilla users 24 Jun 14 21:20 wheeler-scratch -> /wheeler/scratch/vanilla
[vanilla@wheeler ~]$
```

```
[vanilla@wheeler ~]$ ls -l mystuff/
total 473704
-rw-r--r-- 1 vanilla users 483165473 Jun 14 23:20 myfile1
-rw-r--r-- 1 vanilla users 0 Jun 14 22:05 myfile2
[vanilla@wheeler ~]$
```

```
[vanilla@wheeler ~]$ ls -lh mystuff/
total 463M
-rw-r--r-- 1 vanilla users 461M Jun 14 23:20 myfile1
-rw-r--r-- 1 vanilla users 0 Jun 14 22:05 myfile2
[vanilla@wheeler ~]$
```
[vanilla@wheeler ~]\$ du -s
499704 .
[vanilla@wheeler ~]\$ du -sh
488M .

[vanilla@wheeler ~]\$

Figuring out where you going...

#### [vanilla@wheeler ~]\$ df -h

Filesystem Size Used Avail Use% Mounted on devtmpfs 24G 0 24G 0% /dev 24G 64K 24G 1%/dev/shm tmpfs 24G 968M 23G 5%/run tmpfs 24G 0 24G 0% /sys/fs/cgroup tmpfs /dev/mapper/centos-root 930G 567G 363G 61%/ /dev/sdc2 836G 72G 764G 9% /tmp /dev/md126p1 2.0G 333M 1.7G 17% /boot 172.17.2.254:/mnt/wheeler-scratch 37T 28T 8.7T 77% /wheeler/scratch 172.17.2.255:/mnt/wheeler-scratch2 37T 28T 9.0T 76% /wheeler/scratch2 beegfs\_nodev 110T\_51T\_60T\_46%/carc/scratch chama:/home/homes 65T 36T 30T 55% / users chama:/home/carc\_projects 65T 36T 30T 55% /projects

### Figuring out where you going...

[vanilla@wheeler ~]\$ quota -s

Disk quotas for user vanilla (uid 659):

Filesystem space quota limit grace files quota limit grace chama:/home/homes

488M 100G 200G 315 4295m 4295m [vanilla@wheeler~]\$

#### Figuring out where you going...

[vanilla@wheeler ~]\$ stat mystuff/myfile1 File: 'mystuff/myfile1' Size: 483165473 Blocks: 947408 IO Block: 65536 regular file Device: 28h/40d Inode: 9232782834205560540 Links: 1 Access: (0644/-rw-r--r--) Uid: ( 659/ vanilla) Gid: ( 100/ users) Access: 2022-06-14 22:05:27.503289000 -0600 Modify: 2022-06-14 23:20:26.945918000 -0600 Change: 2022-06-14 23:20:48.754917000 -0600 Birth: -

### Figuring out what you've got...

[vanilla@wheeler ~]\$ find -name myfile2 ./mystuff/myfile2

[vanilla@wheeler ~]\$ find -name "myfile\*"
./mystuff/myfile1
./mystuff/myfile2
./mystuff/myfile3
./mystuff/myfile0

Wildcard

Figuring out what you've got...

[vanilla@wheeler ~]\$ cd mystuff/ [vanilla@wheeler ~/mystuff]\$

## Use the tab key to autocomplete

Going somewhere new...

Now it is your turn...



## • For this path: /projects/shared/workshops/beginner/vecadd

- What are the names of the files in that directory?
- When were they last modified?
- How large are the files?

You can find this information with the ls command.

Now it is your turn...



## • For this path: /projects/shared/workshops/beginner/vecadd

Now "cd" into that directory using <tab> autocomplete.

- Now you know how to find your way around filesystems using bash
- Let's see how to modify the filesystem.

- In bash to move a file we use the mv command.
- To copy a file it is cp.
- To copy files from CARC to a personal computer use scp or rsync.

[vanilla@wheeler beginner]\$ pwd /projects/shared/workshops/beginner [vanilla@wheeler beginner]\$ cd ~ [vanilla@wheeler ~]\$ pwd /users/vanilla [vanilla@wheeler ~]\$

First return to your home directory...

[vanilla@wheeler ~]\$ cd mystuff [vanilla@wheeler ~/mystuff]\$ mv myfile1 myfile0 [vanilla@wheeler ~/mystuff]\$ ls myfile0 myfile2 myfile3 [vanilla@wheeler ~/mystuff]

Modifying the filesystem... moving a file. [vanilla@wheeler ~/mystuff]\$ cp myfile0 myfile1
[vanilla@wheeler ~/mystuff]\$

**Source** Destination

[vanilla@wheeler ~/mystuff]\$ ls
myfile0 myfile1 myfile2 myfile3
[vanilla@wheeler ~/mystuff]\$

Modifying the filesystem... copying a file.

#### [vanilla@wheeler ~/mystuff]\$ mkdir mynewdir [vanilla@wheeler ~/mystuff]\$



Modifying the filesystem... create a new directory. [vanilla@wheeler ~]\$ cd ~
[vanilla@wheeler ~]\$ cp -r mystuff mystuff2
[vanilla@wheeler ~]\$

**Destination** 

Source

[vanilla@wheeler ~]\$ ls
mystuff mystuff2 wheeler-scratch

Copying a whole directory tree...

[vanilla@wheeler ~]\$ exit Lycaon:~ matthew\$ scp vanilla@wheeler.alliance.unm.edu:~/mystuff/myfile3 /tmp/





(vanilla@wheeler.alliance.unm.edu)Password:myfile3100%402.0KB/s00:00

Copying data to a personal computer from CARC...

Lycaon: matthew\$ scp -r vanilla@wheeler.alliance.unm.edu: /mystuff /tmp/





(vanilla@wheeler.alliance.unm.edu) Password:myfile1100% 1024KB 6.5MB/s 00:00myfile2100% 2048KB 382.5KB/s 00:05myfile3100% 40 3.2KB/s 00:00myfile0100% 1024KB 8.8MB/s 00:00

Copying data to a personal computer from CARC...

Lycaon: matthew\$ scp -r /tmp/mystuff vanilla@wheeler.alliance.unm.edu: //





(vanilla@wheeler.alliance.unm.edu) Password:									
myfile1	100%	1024	KB	591.5KB/s	00:01				
myfile0	100%	1024	KB	2.0MB/s	00:00				
myfile2	100%	2048	KB	2.1MB/s	00:00				
myfile3	100%	40	2	.1KB/s 00	:00				

To copy from a personal computer to CARC...

### 15 Minute Break

#### ssh vanilla@wheeler.alliance.unm.edu

Log back into wheeler...

[vanilla@wheeler ~]\$ file mystuff/myfile0 mystuff/myfile0: data

[vanilla@wheeler ~]\$ file mystuff/myfile3
mystuff/myfile3: ASCII text

Figuring out file types ...

[vanilla@wheeler ~]\$ cat mystuff/myfile3 Welcome to the CARC Beginner's Workshop



#### [vanilla@wheeler ~]\$ nano mystuff/myfile3



[vanilla@wheeler ~]\$ date Wed Jun 15 03:08:15 MDT 2022

[vanilla@wheeler ~]\$ echo Hello from \$HOSTNAME Hello from wheeler

[vanilla@wheeler ~]\$ hostname wheeler

Programs we will use as examples...

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#### • • •

🛅 Matthew — ssh wheeler — 71×19

#### File Edit Options Buffers Tools Sh-Script Help

<mark>#</mark>!/bin/bash

## echo Hello from \$HOSTNAME date

Shell Scripts

#### [vanilla@wheeler ~]\$ bash myscript.sh

Programs we will use as examples...

#### Software Access



#### [vanilla@wheeler ~]\$ module spider matlab

matlab: Versions: matlab/R2017a matlab/R2018b matlab/R2019a matlab/R2020a matlab/R2021a

#### Getting access to software...

[vanilla@wheeler ~]\$ module load matlab/R2021a Lmod has detected the following error: Matlab may only be run on compute nodes. wheeler is not a compute node. Exiting... While processing the following module(s): Module fullname Module Filename

matlab/R2021a /opt/local/modules/matlab/R2021a.lua

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Getting access to software...

[vanilla@wheeler ~]\$ module load matlab/R2021a Lmod has detected the following error: Matlab may only be run on compute nodes. wheeler is not a compute node. Exiting... While processing the following module(s): Module fullname Module Filename

matlab/R2021a /opt/local/modules/matlab/R2021a.lua

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## What is a compute node?

### Getting access to software...

# HPC Cluster











# Never run computations on the head node

# Always use compute nodes

vanilla@wheeler:~ \$ qgrok queues free busy offline jobs nodes CPUs Idle\_CPUS GPUs CPUs/node Memory/node time\_limit CPU\_limit GPU\_limit RAM\_limit

50 170 1360 58 8 48G 2-00:00:00 400 0 2415 144 0 normal 3 23 48G 96580M debug 2 4:00:00 16 0 0 0 2 16 16 0 8 0 144 23 50 172 1376 74 totals: 5 0

#### Compute nodes and partitions...


### Head Node (wheeler)



### **Compute Nodes**





wheeler172

[vanilla@wheeler ~]\$ lscpu Architecture: x86 64 CPU op-mode(s): 32-bit, 64-bit Byte Order: Little Endian CPU(s): 8 On-line CPU(s) list: 0-7 Thread(s) per core: 1 Core(s) per socket: 4 Socket(s): 2 NUMA node(s): 2 Vendor ID: GenuineIntel CPU family: 6 Model: 26 Model name: Intel(R) Xeon(R) CPU X5550 @ 2.67GHz

Wheeler has 8 cores per node....

mfricke@hopper:~ \$ lscpu Architecture: x86 64 CPU op-mode(s): 32-bit, 64-bit Byte Order: Little Endian CPU(s): 64 On-line CPU(s) list: 0-63 Thread(s) per core: 2 Core(s) per socket: 16 Socket(s): 2 NUMA node(s): 2 Vendor ID: GenuineIntel CPU family: 6 Model: 85 Intel(R) Xeon(R) Gold 6226R CPU @ 2.90GHz Model name:

Hopper has 32 real cores (64 virtual cores) per node....

## Architecture

Wheeler Cluster Grid > Wheeler Cluster > --Choose a Node

~



### 5 Minute Break



Technology, IT etc.

# SLURM

means

## Simple Linux Utility for Resource Management



by acronymsandslang.com





[vanilla@wheeler ~]\$ squeue

	JOBID PA	RTITION	NAME	USER S	Т	TIME	NODES	S NODELIST(I	REASON)	
	159914	normal	co-mcpdf	nsharma	2 CG	2-00:0	0:26	1 wheeler2	.57	
	159915	normal	co-mcpdf	nsharma	2 CG	2-00:0	0:26	1 wheeler2	.57	
	159916	normal	co-mcpdf	nsharma	2 CG	2-00:0	0:26	1 wheeler2	.57	
	159917	normal	co-mcpdf	nsharma	2 CG	2-00:0	0:26	1 wheeler2	.57	
	159918	normal	co-mcpdf	nsharma	2 CG	2-00:0	0:26	1 wheeler2	.57	
	159919	normal	co-mcpdf	nsharma	2 CG	2-00:0	0:26	1 wheeler2	.57	
	159912	normal	co-mcpdf	nsharma	2 CG	2-00:0	0:28	1 wheeler2	.57	
	159913	normal	co-mcpdf	nsharma	2 CG	2-00:0	0:28	1 wheeler2	.57	
16680	0_[21-100	)%10] r	normal Ja	annat jar	nnat	PD	0:00	1 (JobArray	TaskLimit)	
	167067	normal	WINDENE	R rubelda	as PD	0:0	00 36	6 (QOSMaxC	puPerUserLim	nit)
	167068	normal	WINDENE	R rubelda	as PD	0:0	00 24			



- [vanilla@wheeler ~]\$ srun --partition debug --nodes 2 hostname
- srun: Account not specified in script or ~/.default\_slurm\_account, using latest project wheeler302.alliance.unm.edu
- You have not been allocated GPUs. To request GPUs, use the -G option in your submission script.
- wheeler301.alliance.unm.edu
- [vanilla@wheeler ~]\$ srun --partition debug --nodes 2 hostname

#### The srun command...

#### We finally used the HPC Cluster!



- [vanilla@wheeler ~]\$ srun --partition debug --nodes 2 hostname
- srun: Account not specified in script or ~/.default\_slurm\_account, using latest project wheeler302.alliance.unm.edu
- You have not been allocated GPUs. To request GPUs, use the -G option in your submission script.
- wheeler301.alliance.unm.edu
- [vanilla@wheeler ~]\$ srun --partition debug --nodes 2 hostname

#### The srun command...

[vanilla@wheeler ~]\$ srun --partition debug --ntasks 8 hostname srun: Account not specified in script or ~/.default\_slurm\_account, using latest project wheeler302.alliance.unm.edu wheeler302.alliance.unm.edu wheeler302.alliance.unm.edu wheeler302.alliance.unm.edu wheeler302.alliance.unm.edu wheeler302.alliance.unm.edu wheeler302.alliance.unm.edu wheeler302.alliance.unm.edu You have not been allocated GPUs. To request GPUs, use the -G option in your submission script. wheeler302.alliance.unm.edu wheeler302.alliance.unm.edu

#### The srun command...

[vanilla@wheeler ~]\$ cp -r /projects/shared/workshops/beginner/vecadd ~
[vanilla@wheeler ~]\$

Review, what does this command do?

[vanilla@wheeler ~]\$ cd vecadd/

[vanilla@wheeler ~/vecadd]\$ module load openmpi/4.1.2-q2zi

What do these commands do?

[vanilla@wheeler ~/vecadd]\$ srun --partition debug --ntasks 4 vecaddmpi

Now run the program with "srun"...

[vanilla@wheeler ~]\$ qgrok queues free busy offline jobs nodes CPUs

normal 0299 1973002400debug 400432totals: 4299 1973042432

srun is good but HPC centers are busy!



#### Compute Node 01

#### Compute Node 02

#### Compute Node 03

Compute Node 04

Compute Node 05









# Scheduler

qgrok

sbatch

## squeue – u USERNAME

## Running Programs on Compute Nodes

- qgrok
- Intro to the Slurm Scheduler
- The srun command
- sbatch
- sinfo
- squeue
- squeue -u username --start
- <u>https://www.cism.ucl.ac.be/Services/Formations/slurm/2016/slurm.pdf</u>
- https://www.nrel.gov/hpc/assets/pdfs/slurm-advanced-topics.pdf

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#### File Edit Options Buffers Tools Sh-Script Help

#!/bin/bash

**#SBATCH** --job-name=demo

#SBATCH --ntasks=4

#SBATCH --time=00:10:00

```
#SBATCH --mem-per-cpu=4G
```

```
#SBATCH --mail-user=yourusername@unm.edu
```

```
#SBATCH --mail-type=All
```

# Enter the commands you want to run below here: sleep 60 echo Hello from node \$HOSTNAME

## [vanilla@wheeler ~/vecadd]\$ cat vecaddmpi.sh

- #!/bin/bash
- #SBATCH --job-name=vecaddmpi
- #SBATCH --ntasks=4
- #SBATCH --time=00:10:00
- #SBATCH --mem-per-cpu=4G
- #SBATCH --mail-user=mfricke@unm.edu
- #SBATCH --mail-type=All
- #SBATCH --output=vecaddmpi.out

#### module load openmpi/4.1.2-q2zi srun ./vecaddmpi Slurm Script

vanilla@wheeler:~/vecadd \$ sbatch vecaddmpi.sh
sbatch: Using account 2016199 from ~/.default\_slurm\_account
Submitted batch job 167571

vanilla@wheeler:~/vecadd \$ squeue --me JOBID PARTITION NAME USER ST TIME NODES NODELIST(REASON) 167571 normal vecaddmp vanilla R 0:07 1 wheeler145

vanilla@wheeler:~/vecadd \$ tail -f vecaddmpi.out

- You have not been allocated GPUs. To request GPUs, use the -G option in your submission script.
- Assigning compute node to rank 1.
- ComputeNode: Starting with rank 1.
- ComputeNode (1): Waiting for vectors from dataserver with rank 3...
- Assigning compute node to rank 2.
- ComputeNode: Starting with rank 2.
- ComputeNode (2): Waiting for vectors from dataserver with rank 3...
- Will try to allocate a vector of size 1 GB.

vanilla@wheeler:~/vecadd \$ tail -f vecaddmpi.out

- You have not been allocated GPUs. To request GPUs, use the -G option in your submission script.
- Assigning compute node to rank 1.
- ComputeNode: Starting with rank 1.
- ComputeNode (1): Waiting for vectors from dataserver with rank 3...
- Assigning compute node to rank 2.
- ComputeNode: Starting with rank 2.
- ComputeNode (2): Waiting for vectors from dataserver with rank 3...
- Will try to allocate a vector of size 1 GB.

### Useful Slurm Commands

squeue --me --long squeue --me --start scancel jobid sacct shows information about jobs you submitted shows when slurm expects your job to start cancels a job shows your job history

vanilla@wheeler:~/vecadd \$ seff 167573 Job ID: 167573 **Cluster: wheeler** User/Group: mfricke/users State: COMPLETED (exit code 0) Nodes: 1 Cores per node: 4 CPU Utilized: 00:01:03 CPU Efficiency: 78.75% of 00:01:20 core-walltime Job Wall-clock time: 00:00:20 Memory Utilized: 39.55 MB (estimated maximum) Memory Efficiency: 0.24% of 16.00 GB (4.00 GB/core)

### CARC Resources

- Tutorial Videos
- Written Tutorials





# Getting Help

# help@carc.unm.edu

## Office hours

### Bonus – sview and graphical programs