

# Lecture 5 - Introduction to ISAAC

DSE 512

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# From Last Time

- No announcements
- Questions about Docker? EC2?
- Questions about anything else?
- *What scientists must know about hardware to write fast code*  
<https://biojulia.net/post/hardware/>

# Using ISAAC

# What is ISAAC

- Infrastructure for Scientific Applications and Advanced Computing
- Campus cluster
- Heterogeneous
- System overview <https://oit.utk.edu/hpsc/isaac-open-enclave-new-kpb/system-overview-cluster-at-kpb/>

# Where to get help

- Read the documentation! <https://oit.utk.edu/hpsc/isaac-open-enclave-new-kpb/>
- Ask question in class
- ISAAC Office Hours TRF 11:00am-11:59am <https://oit.utk.edu/hpsc/>
- Email and/or schedule office hours with instructor

## Similarities

- Hardware is entirely managed
- "Someone else's computer"

## Differences

- Software environment is *largely managed*
- No administrative privileges
- No Docker

# Logging in with ssh

- ~~ISAAC acf login.acf.tennessee.edu~~
- ISAAC-NG login.isaac.tennessee.edu
- Credentialing:
  - Username is netid
  - Enter netid password when asked
  - Use 2-factor method specified

# Logging in with ssh

```
$ ssh username@login.isaac.tennessee.edu
```

```
$ ssh mschmid3@login.isaac.tennessee.edu
```



# ssh config

Add a similar line to `~/.ssh/config`

```
Host isaac
HostName login.isaac.tennessee.edu
User mschmid3
ServerAliveInterval 30
Port 22
```

```
$ ssh isaac
```

# Node Tiering

- Login nodes
  - Where you land after logging in
  - *Not where jobs should run*
  - Shared resource
  - file editing, code compilation, job submission
- Compute nodes
  - You have to go out of your way to get here
  - *Where jobs should run*
  - Semi-isolated resource

# Accessing the Compute Nodes



# Basic Slurm

Task	Command	Example
Submit job	<code>sbatch</code>	<code>sbatch myjob.slurm</code>
Run interactive job	<code>srun</code>	<code>srun ... --pty bash -i</code>
Show jobs	<code>squeue</code>	<code>squeue</code>
Show my jobs	<code>squeue</code>	<code>squeue --me</code>
Cancel job	<code>scancel</code>	<code>scancel 123456</code>

# Project Account

- Course account
  - ACF-UTK0188
  - `/lustre/isaac/proj/UTK0188`
- Opportunistic account
  - ACF-UTK0011

# Software

- No root
  - no `sudo apt install ...`
  - no Docker
- Instead
  - modules
  - building from source
  - pip/conda/...
  - singularity

# Modules

Task	Command
Show available modules	<code>module avail</code>
Load module	<code>module load \${MODULE}</code>
Unload module	<code>module unload \${MODULE}</code>
Show loaded modules	<code>module list</code>
Module description	<code>module whatis \${MODULE}</code>
Search modules for string	<code>module apropos \${MY_STRING}</code>
Get help	<code>module help</code>

# Python on ISAAC

- Load with `module load Python`
- `pip`
- `conda (module load anaconda3)`



- /sw/isaac/applications/r\_studio/4.0.4/singularity\_r\_studio/R.4.0.4.simg

```
RIMGPATH="/sw/isaac/applications/r_studio/4.0.4/singularity_r_studio/R.4.0.4.simg"
alias R="singularity exec ${RIMGPATH} R --no-save --quiet"
alias Rscript="singularity exec ${RIMGPATH} Rscript"

$ Rscript -e "1+1"
[1] 2
```

# Job Files

```
#!/bin/bash
#SBATCH --account UTK0188
#SBATCH --partition=campus
#SBATCH --job-name=R_hello_world
#SBATCH --nodes=1
#SBATCH --time=00:00:30

RIMGPATH="/sw/isaac/applications/r_studio/4.0.4/singularity_r_studio/R.4.0.4.simg"
singularity exec ${RIMGPATH} Rscript -e "print('hello world')"
```

```
sbatch myjob.slurm
```

# Next Time

- Multi-node jobs
- Singularity
- Homework?

# Live Demo