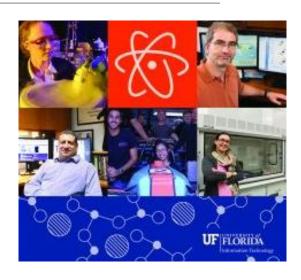
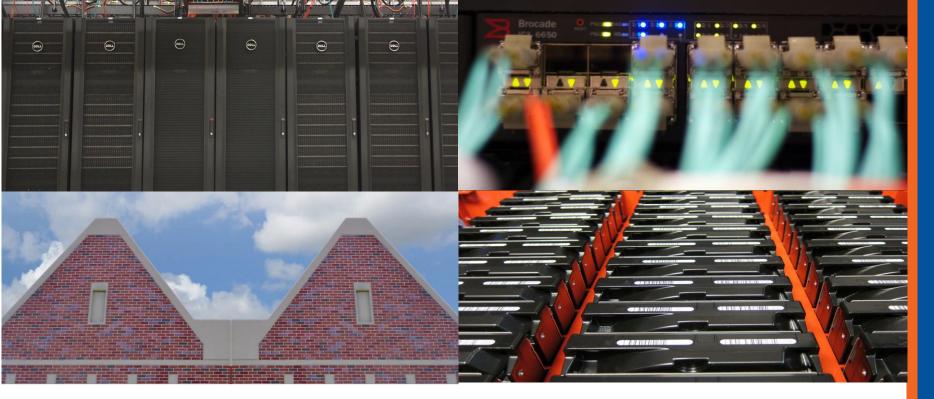
Research Computing Orientation for UF Courses

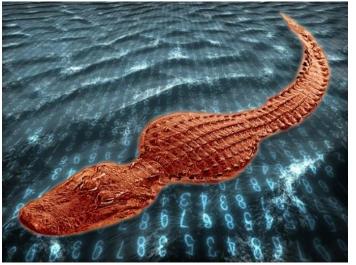
Matt Gitzendanner magitz@ufl.edu







Providing computational resources for a top 10 university





OneIT for the #GatorGood



- 51,000 Cores
- 6PB Storage
- High-speed networks



Course is allocated 32 cores & 112GB RAM

- Design projects with this in mind
- Time your work with this in mind
- Use resources efficiently

Support requests should go through course TA

- If TA cannot solve the issue, the TA should open support requests
- By using your account, you agree to the AUP
 - http://www.rc.ufl.edu/about/policies/
 - No restricted data

For users with an account

- If you already have a Research Computing account for research:
- module load class/pre1234
 - Ensures that jobs run under course allocation
 - Storage is under course allocation

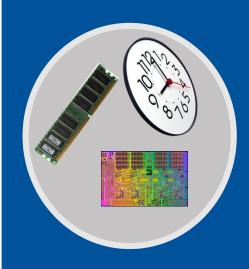
Cluster basics

User interaction



Login node (Head node)

Scheduler



Tell the scheduler what you want to do

Compute resources



Your job runs on the cluster

Tools

ssh client to connect to hpg.rc.ufl.edu





e.g.: Terminal, MobaXterm

SFTP client to move files to/from your computer





e.g.: Cyberduck, FileZilla

Text editor

Especially on Windows, be sure to convert DOS line breaks to Unix, and don't use Word





e.g.: VS Code

SSH Clients

• •

nagitz — magitz@login2:~ — ssh magitz@hpg.rc.ufl.edu — 80×30

flmnh-C02N85F3G:~ magitz\$ ssh magitz@hpg.rc.ufl.edu magitz@hpg.rc.ufl.edu's password:

Last login: Fri May 25 12:11:54 2018 from login4.ufhpc

Welcome to UF Research Computing

The user agrees to comply with Research Computing's policies.

https://www.rc.ufl.edu/about/policies

Backup Policy

Data on Research Computing storage systems is NOT backed up without investment in backup services. Please visit the link below for more information.

https://www.rc.ufl.edu/about/policies/storage/protection/

UFIT Policy Notice

The user understands and acknowledges that the computer and the network are the property of the University of Florida (UF). The user agrees to comply with the UF Acceptable Use Policy and Guidelines. UF monitors computer and network activities without user authorization. UF may provide information about computer or network usage to UF officials, including law enforcement when warranted. Therefore, the user should have limited expectations of privacy.

[magitz@login2 ~]\$ ■



Mac/Linux: Terminal



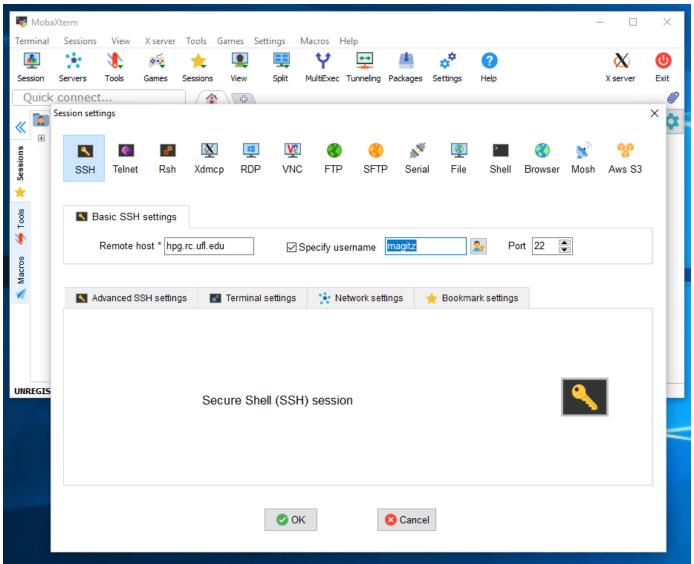


Windows: MobaXterm or PuTTY

ssh gatorlink@hpg.rc.ufl.edu

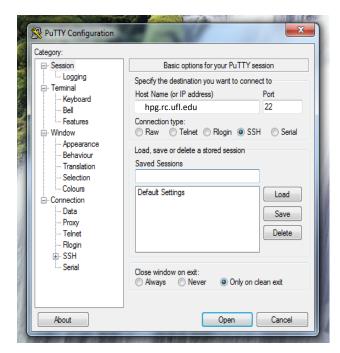


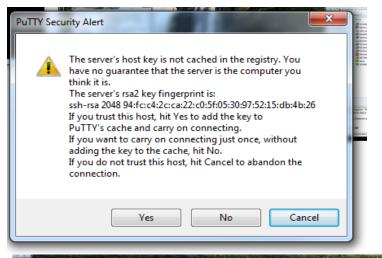
MobaXterm



PuTTY



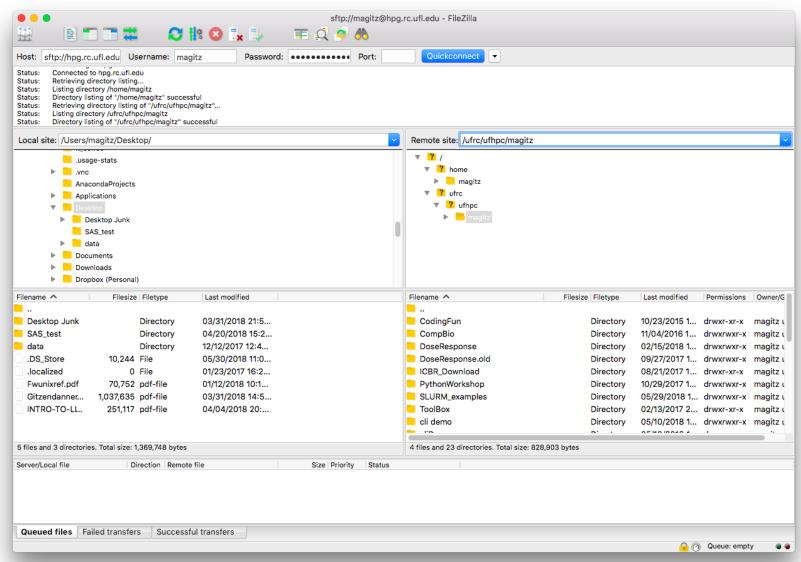






FileZilla





Text Editors

- Not Microsoft Word or other word processor
- Contextual coloring
- Built-in SFTP Client
- Regular expression find/replace

Unix line breaks



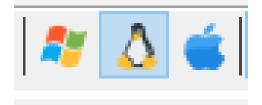




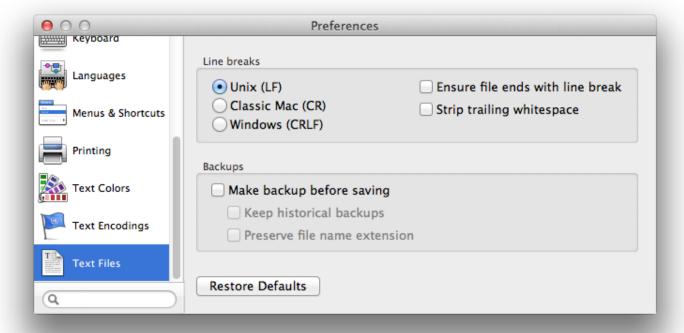


Unix line breaks









Storage

- /home/\$USER
 - 20GB limit
 - scripts, code, small data
 - Do NOT use for job input/output
- /ufrc/pre1234/\$USER
 - 2TB limit per group
 - ALL input/output from jobs should go here

- All storage systems are for research and coursework data only
- Nothing is backed up
- All course accounts are deleted at the end of the semester



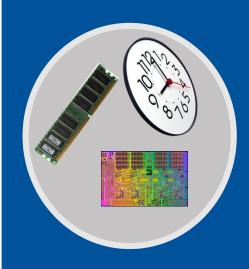
Cluster basics

User interaction



Login node (Head node)

Scheduler



Tell the scheduler what you want to do

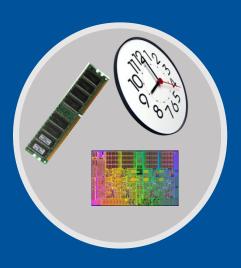
Compute resources



Your job runs on the cluster

Cluster basics

Scheduler



Tell the scheduler what you want to do

Development servers



GUI servers



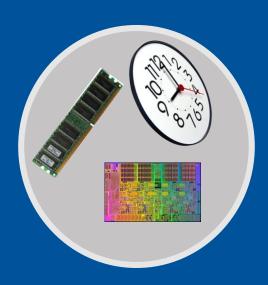
Compute servers



Scheduling a job

- Need to tell scheduler what you want to do
 - How many CPUs you want and how you want them distributed
 - How much RAM your job will use
 - How long your job will run
 - The commands that will be run

Scheduler



Tell the scheduler what you want to do

Basic SLURM job script

```
#!/bin/sh
#SBATCH --ntasks=1
                                      # Run on a single CPU
#SBATCH --mem=1qb
                                      # Memory limit
#SBATCH --time=00:05:00
                                      # Time: hrs:min:sec
#SBATCH --job-name=serial job test  # Job name
#SBATCH --mail-type=ALL
                                     # Mail events
#SBATCH --mail-user=email address # Where to send mail
#SBATCH --output=serial test %j.out
                                     # Output and error log
pwd; hostname; date
module load python
echo "Running plot script on a single CPU core"
python /ufrc/data/training/SLURM/plot template.py
date
UF Information Technology
```

OneIT for the #GatorGood

```
#!/bin/sh
#SBATCH --ntasks=1
```

- Nodes: --nodes or -N
 - Request a certain number of physical servers
- Tasks: --ntasks or -n
 - Total number of tasks job will use
- CPUs per task: --cpus-per-task or -c
 - Number of CPUs per task

For single processor jobs

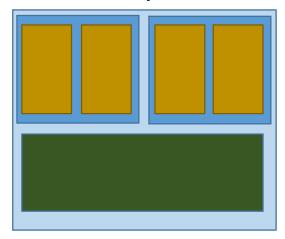
```
#SBATCH --nodes=1

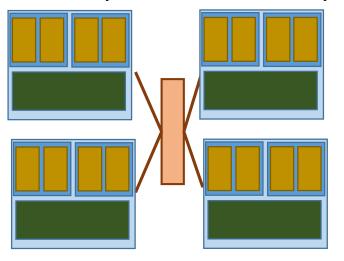
#SBATCH --ntasks=1

#SBATCH --cpus-per-task=1

Optional
```

- Parallel applications
 - OpenMP, Threaded, Pthreads
 - All cores on one sever, shared memory
 - MPI
 - Can use multiple servers
 - See: https://wiki.rc.ufl.edu/doc/Sample_SLURM_Scripts



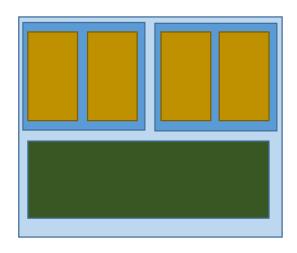




For threaded applications (single node):

```
#SBATCH --nodes=1
#SBATCH --ntasks=1

#SBATCH --cpus-per-task=8
```



Node information

- HiPerGator 2.0 Compute Servers
 - 32 cores
 - Two 16-core sockets
- HiPerGator 1.0 Compute Servers
 - 64 cores
 - Four 16-core sockets

SLURM Memory Requests

- --mem=1gb (total memory)
- --mem-per-cpu=1gb (memory per core)
 - Can use mb or gb
 - No decimal values: use 1500mb, not 1.5gb

HiPerGator 2.0 Compute Servers:

- ~ 120 GB available RAM
- HiPerGator 1.0 Compute Servers:
- ~ 250 GB available RAM

SLURM Time Request

- Time: --time or -t
 - 120 (minutes)
 - 2:00:00 (hh:mm:ss)
 - 7-0 (days-hours)
 - 7-00:00 (days-hh:mm)
 - 7-00:00:00 (days-hh:mm:ss)

Emails

Job ID: 94392

Cluster: hipergator

User/Group: magitz/ufhpc

State: COMPLETED (exit code 0)

Nodes: 1

Cores per node: 4

CPU Utilization: 00:00:44

CPU Efficiency: 52.38% of 00:01:24 core-walltime

Memory Utilization 1.52 MB

Memory Efficiency: 0.04% of 4.00 GB

Emails

```
Job ID: 5019
Cluster: hpg1
User/Group: magitz/ufhpc
State: CANCELLED (exit code 0)
Cores: 1
CPU Utilization: 00:00:00
CPU Efficiency: 0.00% of 00:00:00 core-walltime
Memory Utilization 1.26 MB
Memory Efficiency: 126.17% of 1.00 MB
```

Job error file:

```
slurmstepd: Job 5019 exceeded memory limit (1292 > 1024), being
killed
slurmstepd: Exceeded job memory limit
slurmstepd: *** JOB 5019 ON dev1 CANCELLED AT 2016-05-16T15:33:27
***
```



Quality of Service (--qos)

- Each group has two QOS options
 - Investment QOS:
 - The NCUs the group has purchased
 - --qos=pre1234 (or leave off as this is default)
 - Burst QOS: 9X the allocation
 - The burst capacity, available when idle resources are available on the cluster
 - --qos=pre1234-b
- Users can choose higher priority, or larger pool of resources

SLURM

Note that multi-letter directives are double-dash:

```
sbatch: error: distribution type
'ail-type=ALL' is not recognized
```

- --ntasks
- --mem-per-cpu
- Do not use spaces with =
 - --mail-user=magitz@ufl.edu 🗸
 - --mail-user magitz@ufl.edu 🗸
 - not: --mail-user= magitz@ufl.edu

Examples

See example job scripts at: help.rc.ufl.edu

Batch System

- SLURM Commands
- Using Variables in SLURM
- SLURM Job Arrays

- Account and QOS
 Limits Under SLURM
- GUI Partition
- Big Memory Partition

- Annotated Job Script
 Walk-through
- Sample SLURM Scripts
- UFRC environment module

Or: /ufrc/data/training/SLURM/

Submit your job

```
[magitz@login3 SLURM_examples]$ sbatch single_job.sh
Submitted batch job 30592170

[magitz@login3 SLURM_examples]$ Squeue -u magitz

JOBID PARTITION NAME USER ST TIME NODES NODELIST(REASON 30592170 hpg2-comp serial_j magitz R 0:30 1 c24b-s15

[magitz@login3 SLURM_examples]$
```



So what is this "module" thing?

- Imod—Implementation of Environment Modules developed at TACC
- Allows easy management of user's environment



TEXAS ADVANCED COMPUTING CENTER



Powering Discoveries That Change The World

lome About Resources User Services Research & Development Partnerships Education & Outreach News

Lmod: Environmental Modules System



Uneil for the #GatorGood

The standard way

```
PATH=$PATH:/some/long/path/to/application
export $PATH
LD_LIBRARY_PATH=$LD_LIBRARY_PATH:/long/path/to/place/I/probably/cant/find
export $LD_LIBRARY_PATH
```

- Need to track down paths to applications, libraries, etc.
- Multiple compilers, and MPI implementations
- Manage dependencies
- Multiple versions of apps





Module discovery

- module spider
 - List everything
- module spider cl
 - List applications that have cl in name
- module spider amber/12
 - List details about this version of AMBER
- module key molecular
 - Keyword search for applications

Multiple versions

[magitz@submit1 ~]\$ module spider gaussian
Rebuilding cache file, please wait done
gaussian:
Description:
A software for electronic structure modeling
Versions:
gaussian/e01
gaussian/g03
gaussian/g09
To find detailed information about gaussian please enter the full name
For example:

\$ module spider gaussian/g09



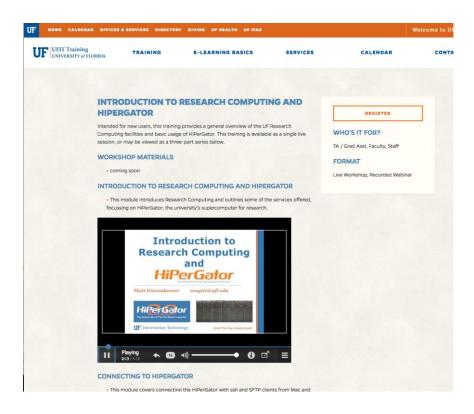
Module loading

- module load raxml
- module load intel raxml
- module load intel openmpi raxml
- module load intel/12 openmpi/1.6 raxml/3.2

module unload raxml

Training resources

https://training.it.ufl.edu





UF Research Computing

- Help and Support
 - http://help.rc.ufl.edu
 - Documents on hardware and software resources
 - Various user guides
 - Many sample submission scripts
 - http://rc.ufl.edu/
 - Frequently Asked Questions
 - Account set up and maintenance

