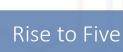


Research Computing
Orientation for Courses

Matt Gitzendanner

magitz@ufl.edu

**UF** Information Technology









# Course use of HiPerGator The University of Florida Supercomputer for Research

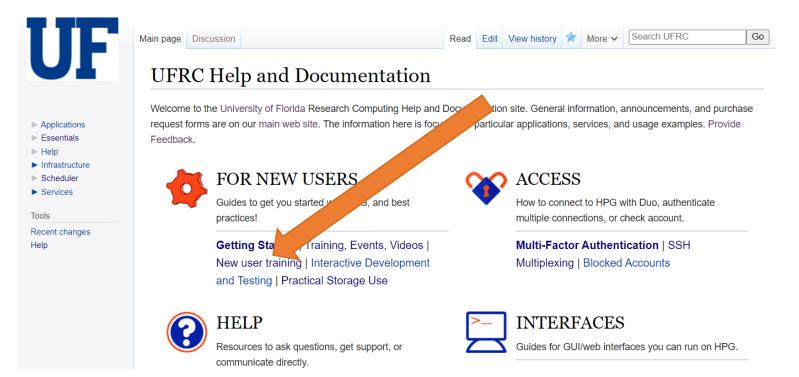
- Course is allocated 32 cores, 256GB RAM, 2TB Blue storage,
   GPUs as needed
  - Design projects with this in mind
  - Time your work with this in mind
  - Use resources efficiently
- Support requests should go through course TA
  - If TA cannot solve the issue, the TA should open support requests
- By using your account, you agree to the AUP
  - http://www.rc.ufl.edu/about/policies/
  - No restricted data





## **HiPerGator Account Training**

- Content and links at: <a href="https://hebr.nc.ufl.edu/doc/New user training">help.rc.ufl.edu/doc/New user training</a>
  - Page also has additional information for classes at the end



https://help.rc.ufl.edu/





# For users with an account

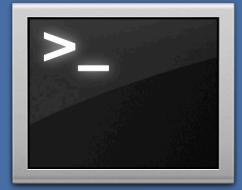
- If you already have a Research Computing account for research:
  - Make a folder for yourself at /blue/pre1234/<gatorlink>
  - When submitting jobs, add:
    - --account=pre1234 --qos=pre1234





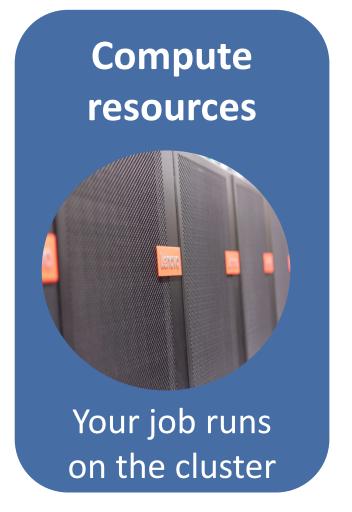
## Cluster overview

**User interaction** 



Login node (Head node)

**SLURM** Scheduler Tell SLURM what you want to do



## Tools for working with HiPerGator

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



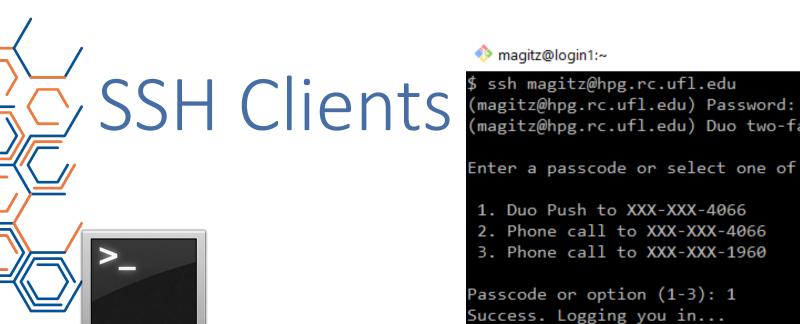
SFTP client to move files to/from your computer hpg.rc.ufl.edu



## Text editor

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





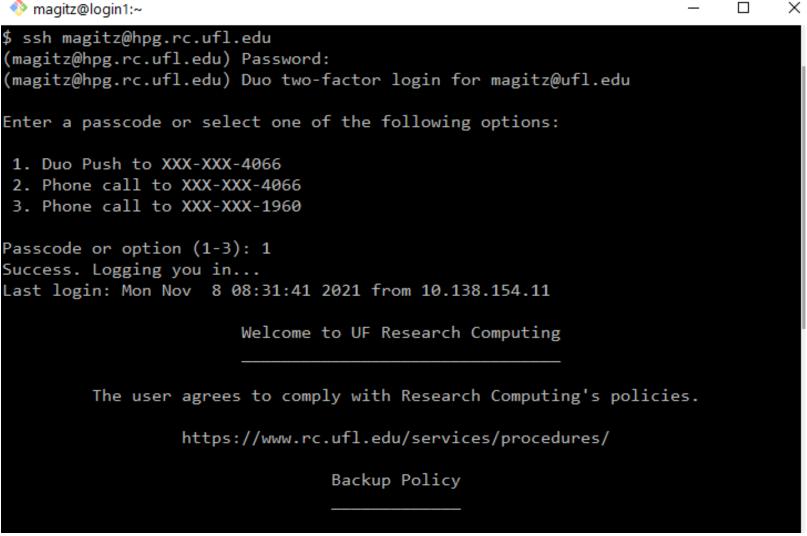
Mac/Linux: Terminal







Windows: Git Bash, MobaXterm, PuTTY, Bitvise



ssh user@hpg.rc.ufl.edu







npg.tlp - magitz@hpg.rc.ufl.edu:22 - Bitvise SSH Client										-	- 1		×
Profile: hpg.tlp													
( 🖺 )	Login	Options	Terminal	RDP	SFTP	Serv	ices	C2S	S2C	SSH	Notes	About	
ě	Server							Authentication					
Save profile	Host hpg.rc.ufl.edu							Username magitz					
0	Port 22 Enable obfuscation							Initial method keybo			ard-interactive ∨		
Save profile as	Obfuscation keyword							Submethods bsdauth,pan			,pam,tot	p,pw,p\	
	Kerberos							Elevation De		Default	Default ∨		
Bitvise SSH	SPN												
Server Control	GSS/Kerberos key exchange												
Panel	Request delegation												
	✓ gssapi-keyex authentication												
New terminal console	Prox	ry settings		Host	key mana	<u>qer</u>		Client ke	ey mana	ager_		<u>He</u>	elp

Initial method: keyboard-interactive





## ood.rc.ufl.edu





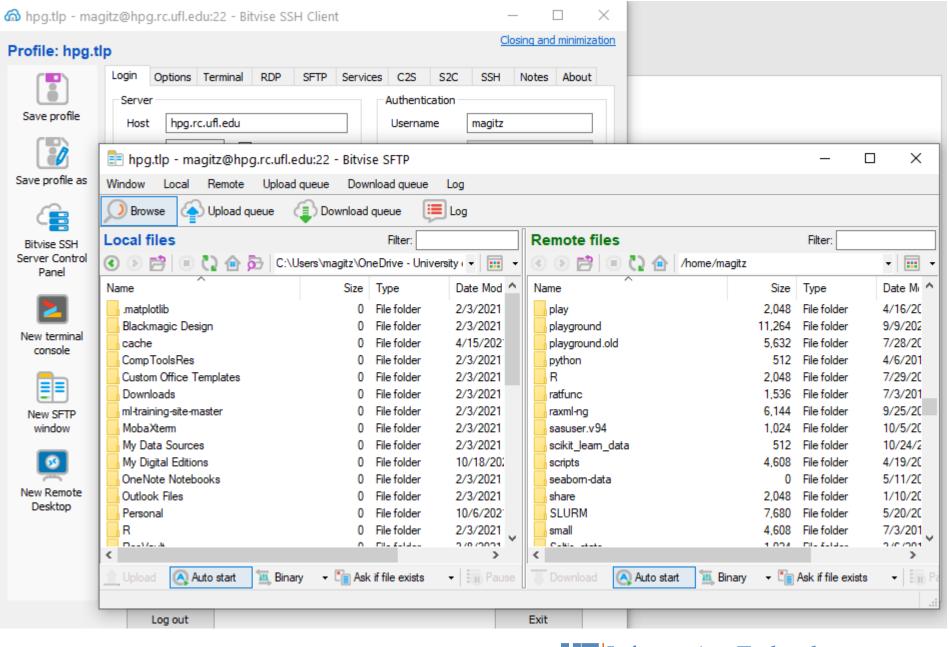
A

Need to be on UF network (VPN if off campus)



## SFTP Client

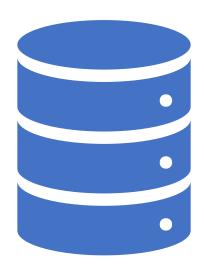






# Storage on HiPerGator

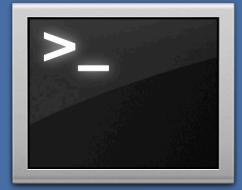
- •Home storage: /home/<user>
  - 40GB limit
  - Scripts, code, compiled applications
  - Do NOT use for job input/output
  - Week of snapshots at ~/.snapshot/
- •Blue storage: /blue/pre1234/<user>
  - 2TB limit per class
  - ALL input/output from jobs should go here
- All storage systems are for research and coursework data only
- Nothing is backed up
- All course accounts are deleted at the end of the semester





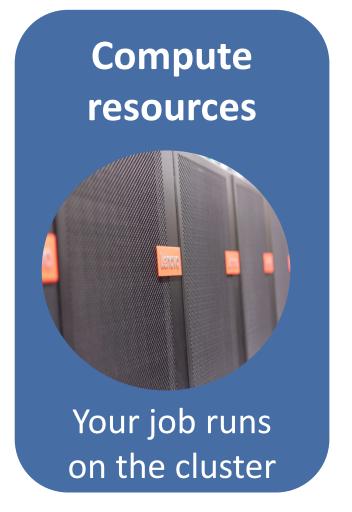
## Cluster overview

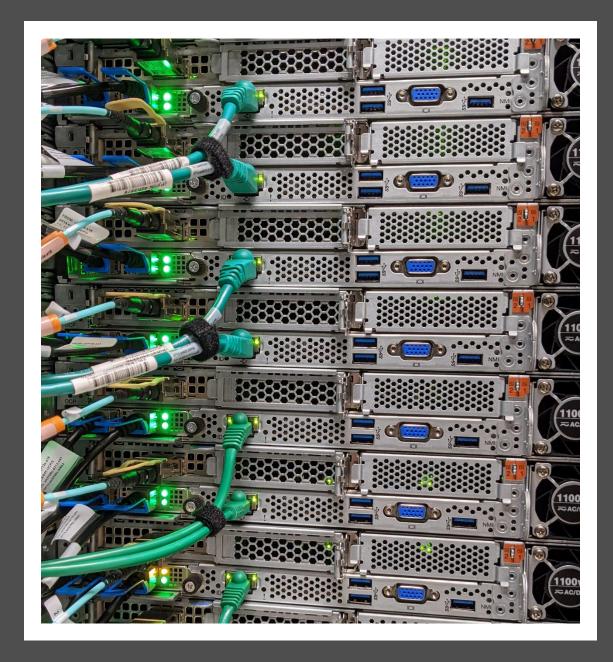
**User interaction** 



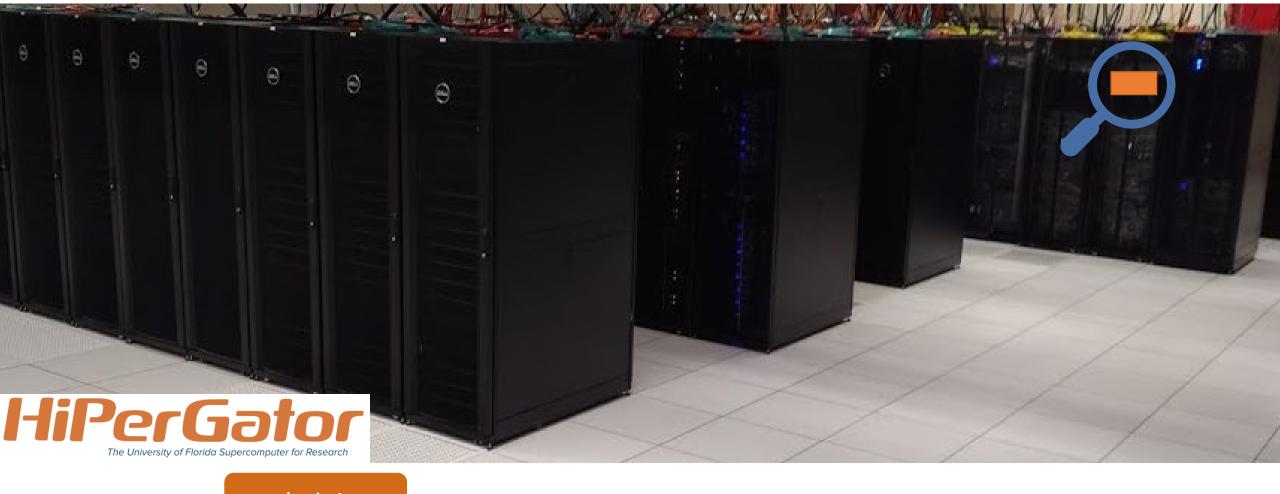
Login node (Head node)

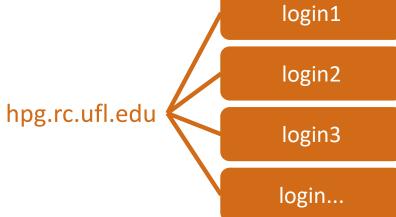
**SLURM** Scheduler Tell SLURM what you want to do













**UF** Information Technology



## Appropriate use of login nodes

- Login nodes are for:
  - File and job management
  - Short-duration interactive testing and development
- Limit your use to **no more than**:





## Resources

Development servers

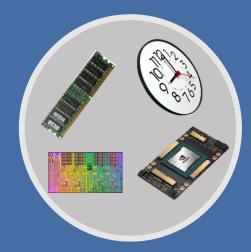


**GPU Servers** 





SLURM Scheduler



Tell SLURM what you want to do

Jupyter severs



**GUI** servers



Batch compute resources





# Jupyter Hub and on Demand

# jhub.rc.ufl.edu ood.rc.ufl.edu



To setup link to the class blue directory, open a Terminal (File> New > Terminal) and run (e.g. for class ast4930):

ln -s /blue/ast4930 blue\_ast4930



VPN required if

off campus



## Jupyter and conda environments

## Be careful with pip install

- Can lead to conflicting versions of packages
- pip installs packages in
   ~/.local/lib/python3.x/site-packages

## Use conda/mamba

- Create isolated environments
- To use in Jupyter, create custom kernel folder. See help page.
- To use in script:

```
module purge; module load conda
conda activate my_env
python my_script.py
```

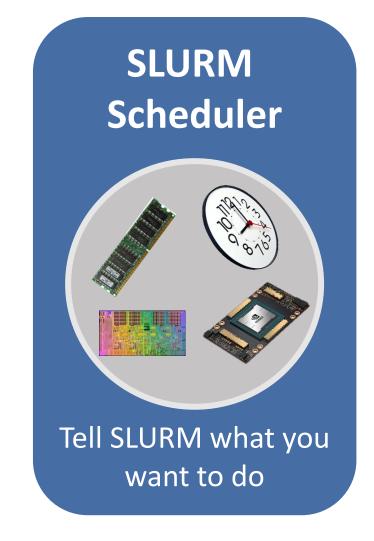
Script should start with:
#!/usr/bin/env python





# Scheduling a job

- What resources does your job need?
  - How many CPUs you want and how you want them grouped?
  - How much RAM your job will use?
  - How long your job will run?
  - How many GPUs?
  - Also need the commands that will be run to do your work

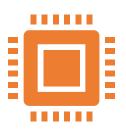


## Basic SLURM job script

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

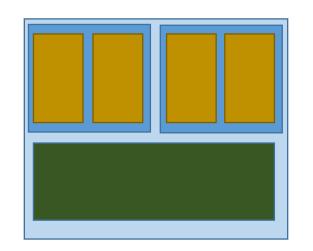


# SLURM CPU Requests



For threaded applications (single node):

```
#SBATCH --nodes=1 # Physical servers
#SBATCH --ntasks=1 # MPI ranks or processes
#SBATCH --cpus-per-task=8
```









## SLURM GPU Requests



```
#SBATCH --partition=gpu # required for GPUs #SBATCH --gpus=1
```

```
#SBATCH --gpus=a100:1 #Specify type
```

See: https://help.rc.ufl.edu/doc/GPU Access

#### Cluster partition (--partition, -p)

gpu

Select a specific cluster partition for job. (default = first available compute partition)

#### Generic Resource Request (--gres).

gpu:a100:1

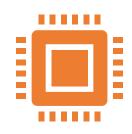
--gres: gpu:a100:1

This is the Generic resource request string to request GPU resources. See also https://help.rc.ufl.edu/doc/GPU\_Access

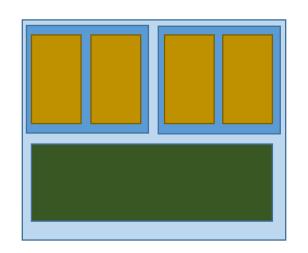


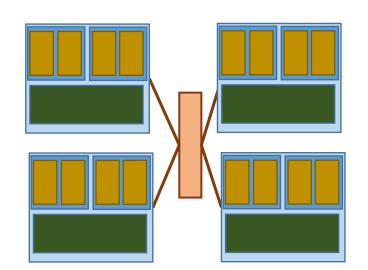


# SLURM CPU Requests



- Parallel applications
  - OpenMP, Threaded, Pthreads
    - All cores on one sever, shared memory
  - MPI—Message Passing Interface
    - Can use multiple servers
    - See: <a href="help.rc.ufl.edu/doc/Sample SLURM Scripts">help.rc.ufl.edu/doc/Sample SLURM Scripts</a>

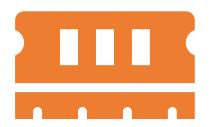








# SLURM Memory Requests



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

HPG 2.0

120

GB RAM

HPG 3.0

1000

GB 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)



# Quality of Service (--qos)

- Each group has two QOS options
  - Investment QOS: --qos=group
  - Burst QOS:
    - The burst capacity, available when idle resources are available on the cluster
    - --qos=group-b
- Users can choose higher priority, or larger pool of resources

### Compute allocation

nvestment

Burst capacity (up to 9X investment)

Lower priority access to idle resources as available

## **SLURM**

Note that multi-letter directives are double-dash:

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

- --mem-per-cpu
- Use either, but not both, space or =
  - --mail-user=magitz@ufl.edu 🗸
  - --mail-user magitz@ufl.edu 🗸
  - not: --mail-user= magitz@ufl.edu

# Submit your job

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

[magitz@login3 SLURM_examples]$ squeue --me

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]$
```



Applications Essentials ▶ Help

Infrastructure Scheduler

Recent changes

Services

Tools

Help

Go

#### **UFRC Help and Documentation**

Welcome to the University of Florida Research Computing Help and Documentation site. General information, announcements, and purchase request forms are on our main web site. The information here is focused on particular applications, services, and usage examples. Provide Feedback.



#### FOR NEW USERS

Guides to get you started with HPG, and best practices!

Getting Started | Training, Events, Videos | New user training | Interactive Development and Testing | Practical Storage Use



#### ACCESS

How to connect to HPG with Duo, authenticate multiple connections, or check account.

Multi-Factor Authentication | SSH Multiplexing | Blocked Accounts



#### HELP

Resources to ask questions, get support, or communicate directly.

How To Get Help | Submit a Support Request | Al Help | FAQ



#### INTERFACES

Guides for GUI/web interfaces you can run on HPG.

Jupyter (Python, R) | Galaxy Genomics Framework | OnDemand (Matlab, RStudio,...) | Conda and Jupyter Kernels | Running GUI Apps



#### SCHEDULER

How to schedule and manage jobs and resources on HPG and more SLURM info.

SLURM Commands | Account/QOS Limits Under SLURM | Big Memory Partition | Available Node Features | SLURM Job Script Examples | SLURM Job Arrays | GPU Access



#### SOFTWARE AND REFERENCE DATA

View installed programs and tools, how to utilize them, and reference data that they can run.

Installed Applications | HiPerGator ufrc Tools | Environment Modules System | Reference Data | Al Reference Datasets



#### TRANSFER DATA

low to move different sizes of data to and from HDG



#### SPECIFIC RESEARCH AREAS

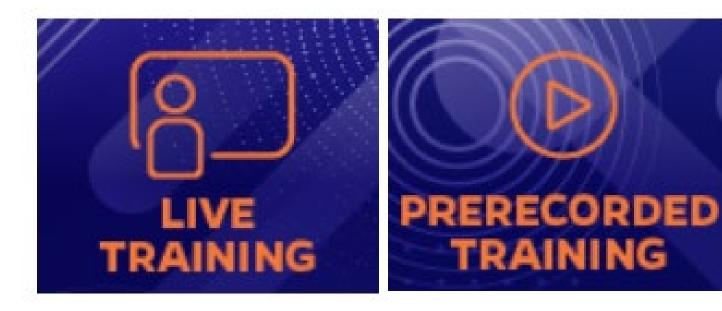
Highlighted areas of specialization commonly seen with HDG

help.rc.ufl.edu





# Training





help.rc.ufl.edu/doc/Training