HiPerGator 2.0:

Moving your research to the next generation of super computing

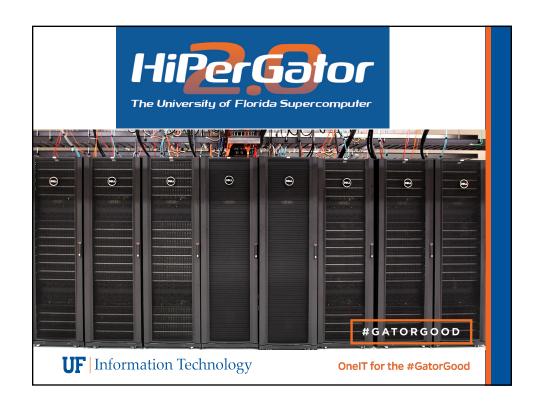
Matt Gitzendanner

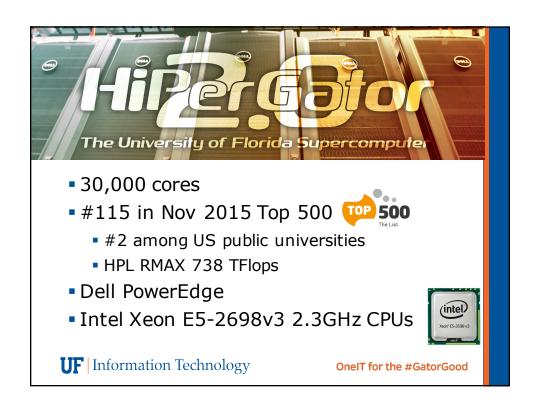
magitz@ufl.edu

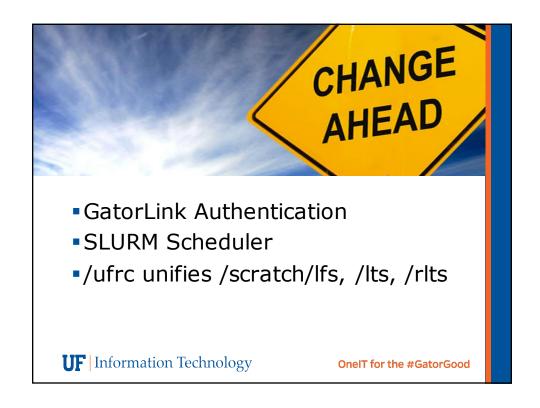






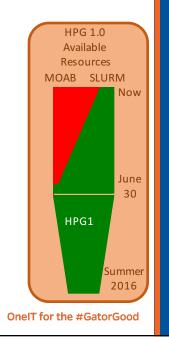






Transition Period

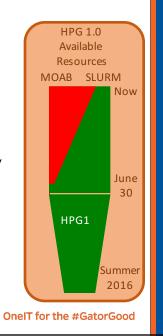
- Three places to send jobs:
 - HiPerGator 1.0:
 - MOAB/Torque
 - SLURM
 - HiPerGator 2.0:
 - SLURM



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Transition Period

- Three places to send jobs:
 - HiPerGator 1.0:
 - MOAB/Torque—No changes
 - Finishing project by 6/30
 - MPI applications not yet ready
 - SLURM
 - HiPerGator 2.0:
 - SLURM

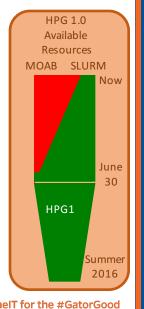


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Transition Period

- Three places to send jobs:
 - HiPerGator 1.0:
 - MOAB/Torque—No changes
 - SLURM
 - module load slurm
 - Rewrite job scripts to use **SLURM**
 - MPI applications need to be recompiled
 - HiPerGator 2.0:
 - SLURM

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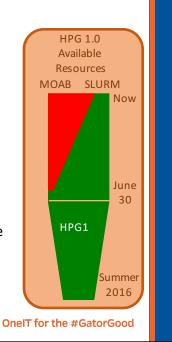


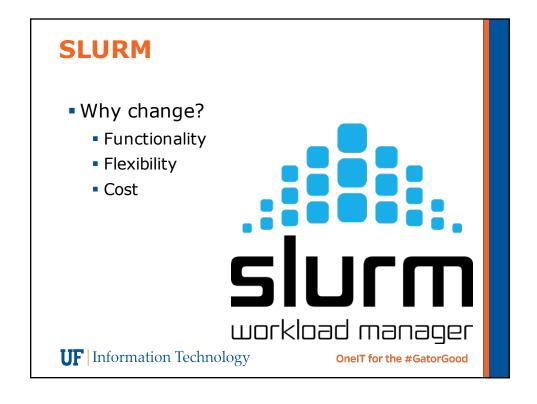
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Transition Period

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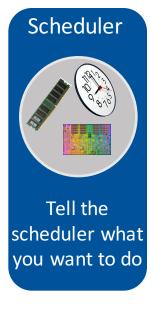
Moab to SLURM

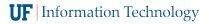
- Documentation
 - PBS2Slurm Command Reference wiki page
 - Other Wiki pages being developed

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Scheduling a job

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





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Basic SLURM job script

```
#!/bin/bash
#SBATCH --job-name=test #A name for your job
#SBATCH -o job_%j.out #Name output file
#SBATCH --mail-type=ALL #What emails you want
#SBATCH --mail-user=<Email address> #Where
#SBATCH --ntasks=1 #Optional-single CPU
#SBATCH --mem-per-cpu=100mb #Per processor memory
#SBATCH -t=00:01:00 #Walltime in hh:mm:ss
#or d-hh:mm:ss
```

hostname module load python python -V

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SLURM CPU Requests

- 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

HiPerGator 2.0 Compute Servers:

• 32 cores (2 X 16-core Intel Xeon CPUs)

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SLURM CPU Requests

- For single processor jobs
 - --ntasks=1 (or omit)
- For parallel jobs on a single node:
 - --ntasks=8

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SLURM CPU Requests

- For MPI jobs
 - --ntasks=32
 - Gets 32 cores for 32 MPI ranks
 - SLURM will determine node layout
- For Hybrid MPI/OpenMP jobs
 - --nodes=4
 - --ntasks=4 (4 MPI ranks)
 - --cpus-per-task=8



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SLURM Memory Requests

- Memory: --mem-per-cpu=1gb
 - Can use mb or gb
 - Like Moab, no decimal values

HiPerGator 2.0 Compute Servers:

- 128 GB total RAM (vs 256 GB on HPG1)
- Diskless servers: OS takes ~8GB RAM

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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)



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SLURM output/error files

- #SBATCH -o output.file
- #SBATCH -e error.file
- #SBATCH -o output.file #W/o -e combined
- Can also use --output and --error
- #SBATCH -o JobFile.%j.out
 - Use %j instead of \$SLURM_JOBID

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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

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Quality of Service (--qos)

- Each group has two QOS options
 - Investment QOS:
 - The NCUs the group has purchased
 - --qos=group (or leave off as this is default)
 - Burst QOS:
 - The burst capacity, available when idle resources are available on the cluster
 - --qos=group-b
- Unlike under MOAB/Torque the burst capacity is not automated

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SLURM Task Arrays

- #SBATCH --array=1-200%10
- Similar to Moab: range with % to limit number of jobs at a time
- •\$SLURM_ARRAY_TASK_ID
- Output file naming:
 - %A: job id%a: task id
 - Output.%A_%a.out

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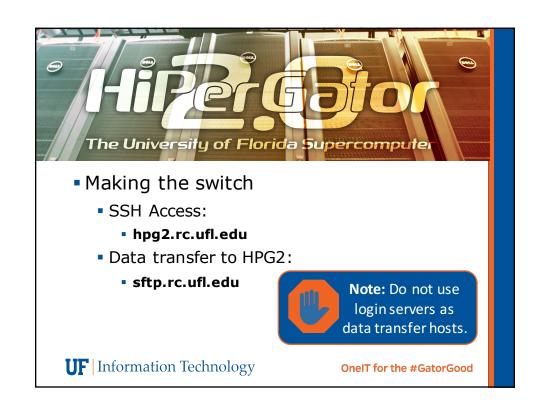
HiPerGator 1.0

- Continue as now until June 30th
 - Dwindling MOAB/Torque resources available
- •module load slurm and use SLURM
 - No need to move data or change login
 - Most applications will work unmodified



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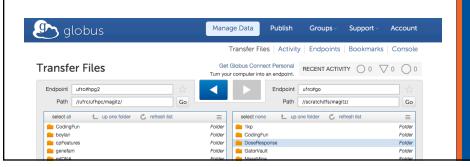




Globus.org



- Fast transfer from /scratch/lfs/ to /ufrc
- Particularly good for large files
- ufrc#hpg2 and ufrc#go



Compilers

- HPG 2.0 and 1.0 under SLURM
 - intel/2016.0.109
 - openmpi/1.10.2
 - gcc/5.2.0
 - openmpi/1.10.2

Note that OpenMPI 1.10.2 on HPG1 is only compatible with SLURM. MOAB cannot run applications compiled with this module.

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Don't look back!

• 30,000 fast cores ready for you!

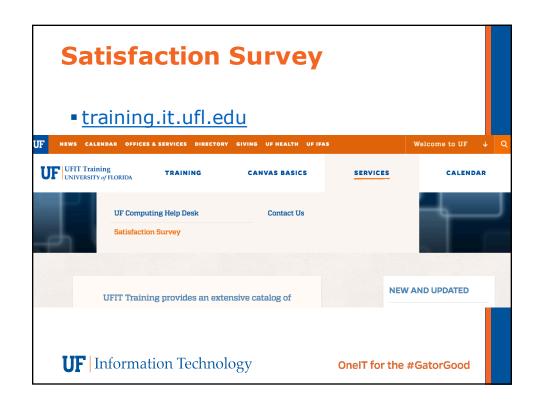


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End of free usage

- Previous policy
 - Up to 8-cores for free
- New policy
 - Research Computing has been told we can no longer offer any free access
 - Try-and-buy loans
 - 1-3 month loan of resources
 - Test the system
 - Verify needs are met
 - Become an investor





Next Week:

- Writing SLURM Submission Scripts
 - 11:00am
 - NPB 2205



