## Introduction to Talon

**UND Computational Research Center** 

#### Overview

- About Talon
- Login to Talon (Open OnDemand)
- Modulefiles
- How to Run Programs/slurm

### How to get a Talon account

- Visit https://crc.und.edu
- Click Support in the menu
- Click Account Request
- Request type: Account request
- Account type: Check Globus, Talon, and Talon GPU
- You should receive a request confirmation and an email when the account has been created



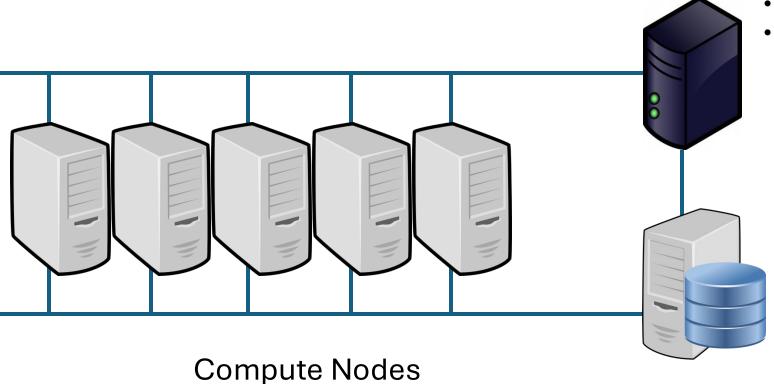
# Talon HPC/Supercomputer

System overview

#### What is Talon?

- A High-Performance Computing (HPC) cluster
  - A small supercomputer
- Compute nodes
  - Grouped into Partitions (aka Queues)
- Resource management
  - Slurm software ensures fair and efficient usage of cores, memory, and GPUs

### Computer Cluster



Research programs run here

#### Login Node

- Login to login node
- Submit jobs from here
- Interface
  - Command line or Browser

#### Network Storage

- Data files
- Program files

## Talon Compute Resources

Partition/Queue	Number Nodes	Total Cores	GPUs	Total Memory (GB)
talon	18	648	-	192
talon-large	3	216	-	3000 (3 TB)
talon-short	2	72	-	192
talon-gpu32	2	144	8	1500 (1.5 TB)
gpu-code-test	1	36	8	1500
6 Private Partitions				

## Open OnDemand

Using Talon with your Browser

### Open OnDemand

- A "one-stop shop" for accessing Talon
  - Web browser provides a graphical interface
- File manager
  - Upload, download, create, edit, copy, move, delete
- Shell access to login node
- Certain applications with graphical user interfaces

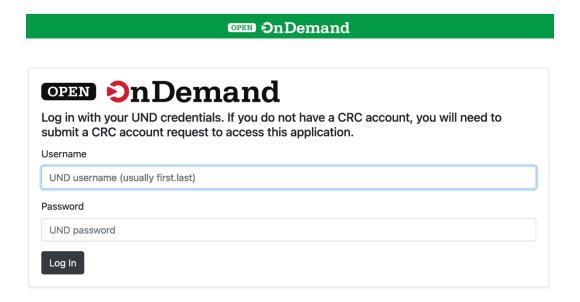
#### **UND VPN**

• If you are off-campus, you must connect to the UND VPN before connecting to Talon.

https://und.teamdynamix.com/TDClient/2048/IT/KB/ArticleDet?ID=145487

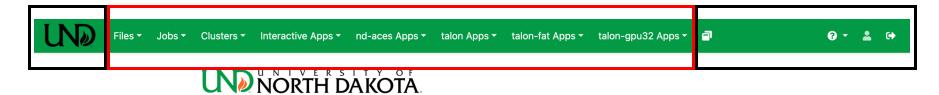
## Log in to Open OnDemand (OOD)

Browse to <a href="https://apps.talon.und.edu">https://apps.talon.und.edu</a>



- Enter your Talon/UND username and password
- Click "Log In" button

#### OOD Dashboard



OnDemand provides an integrated, single access point for all of your HPC resources.

### Navigating the Dashboard, Buttons



Clusters ▼

Interactive Apps ▼

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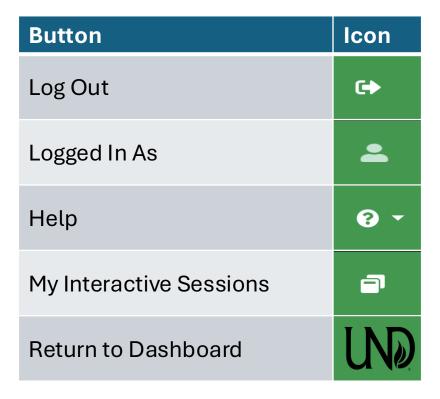
talon Apps ▼

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### Navigating the Dashboard, Menus



Files 🕶

obs ▼

Clusters ▼

Interactive Apps ▼







talon-gpu32 Apps ▼







- Drop-down Menus
  - Files
  - Jobs
  - Clusters
  - Partition-specific Applications

## Module Files

Configure your environment on Talon

#### **Environment Variables**

- Some programs may need specific compilers, libraries, and include files.
- Module files let you configure your environment.

```
$ module avail
$ module list
$ module load [module name]
$ module unload [module name]
$ module help [module name]
Show available modules
Show modules currently loaded
Load a specific module
Unload a module
Short description of a module
```

#### Module Exercise

- Login to Talon
- List the currently loaded modules
- \$echo \$PATH
- \$echo \$LD LIBRARY PATH
- Load the Gnu Compiler module
  - \$ module load gcc
- \$echo \$PATH
- \$ echo \$LD\_LIBRARY\_PATH
- Unload the Gnu Compiler module
- Confirm that the \$PATH and \$LD\_LIBRARY\_PATH are reset

## Slurm

Resource Manager / Job Scheduler

### What is a resource manager?

- A computer cluster is a finite resource
- Users compete for nodes, cores, network, memory...
- Resource manager: special software to provide for fair and efficient cluster usage

### **Key Functions**

- Manage and allocate cluster resources
- Provide accounting for cluster resources and user jobs
- Let users
  - submit jobs to compute nodes
  - monitor jobs running on compute nodes
  - cancel jobs in queue or running on compute nodes

### Helpful Commands

sinfo
 Cluster status

squeue Cluster/job status

squeue --me
 My jobs

squeue -j [job id]

sbatch [script filename] Job submission

scancel [job id]
 Job deletion

# Batch Scripts (Slurm Job Submission Script)

```
#!/bin/bash
  #SBATCH --job-name=hello
  #SBATCH --partition=talon
  #SBATCH --ntasks=1
  #SBATCH --time=1-00:01:00
  #SBATCH --output=%x.%j.txt
  module load gcc
  echo "Job started at $(date)"
  ./a.out [arguments]
11 echo "Job ended at $(date)"
12 echo "Tasks: $SLURM_NTASKS"
```

#### Time to Solution

- All Slurm jobs spend time in queue and running
- Goal is to minimize turnaround time
  - Asking for more resources can increase time in queue (waiting)
  - Asking for more resources can decrease execution time

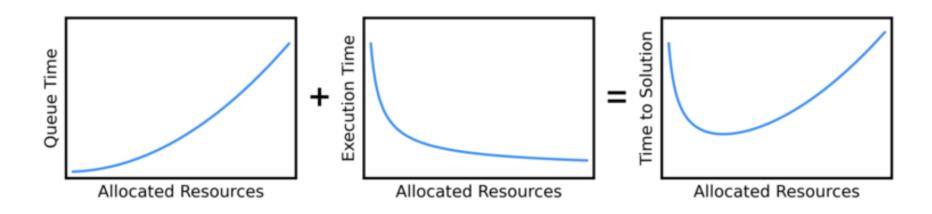


Image:researchcomputing.princeton.edu/support/knowledge\_base/slurm

### Job Length

- Asking for too little time -- Slurm kills the job
- Asking for too much time -- can increase wait time
- The sacct command displays accounting data
  - sacct --format="JobId, JobName, Elapsed, ExitCode" --job=[job id]
- Can help set timelimit on future jobs: Elapsed + 20%

#### Helpful Links

- sbatch
  - https://slurm.schedmd.com/sbatch.html
- squeue
  - https://slurm.schedmd.com/squeue.html
- sinfo
  - https://slurm.schedmd.com/sinfo.html
- Linux commands
  - https://ryanstutorials.net/linuxtutorial/