CIML Summer Institute: CPU Computing - Hardware, architecture, and, software infrastructure

Mary Thomas SDSC

EXPANSE COMPUTING WITHOUT BOUNDARIES

SAN DIEGO SUPERCOMPUTER CENTER



NSF Award 1928224



Outline

- Expanse Overview & Innovative Features
- Getting Started
- Modules
- Account Management
- Compiling and Linking Code
- Running Jobs
- Hands-on Examples
 - MPI Jobs
 - OpenMP Jobs
 - GPU/CUDA Jobs
 - Hybrid MPI-OpenMP Jobs
- Final Comments

- See Expanse 101 tutorial

Basic Information

- Expanse User Guide:
 - https://www.sdsc.edu/support/user_guides/expanse.html
- You need to have an Expanse account in order to access the system. There are a few ways to do this:
 - Submit a proposal through the <u>XSEDE Allocation Request System</u>
 - PI on an active allocation can add you to their allocation (if you are collaborators working on the same project).
 - Request a trial account, instructions @ https://portal.xsede.org/allocations/startup.
- Online repo and information:
 - https://github.com/sdsc-hpc-training-org/expanse-101
 - https://hpc-training.sdsc.edu/expanse-101/



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- Data and Storage
- Final Comments



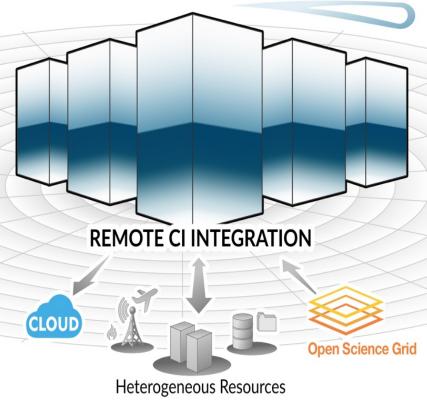
EXPANSE COMPUTING WITHOUT BOUNDARIES 5 PETAFLOP/S HPC and DATA RESOURCE

HPC RESOURCE

13 Scalable Compute Units728 Standard Compute Nodes52 GPU Nodes: 208 GPUs4 Large Memory Nodes

DATA CENTRIC ARCHITECTURE

12PB Perf. Storage: 140GB/s, 200k IOPS
Fast I/O Node-Local NVMe Storage
7PB Ceph Object Storage
High-Performance R&E Networking



LONG-TAIL SCIENCE

Multi-Messenger Astronomy
Genomics
Earth Science
Social Science

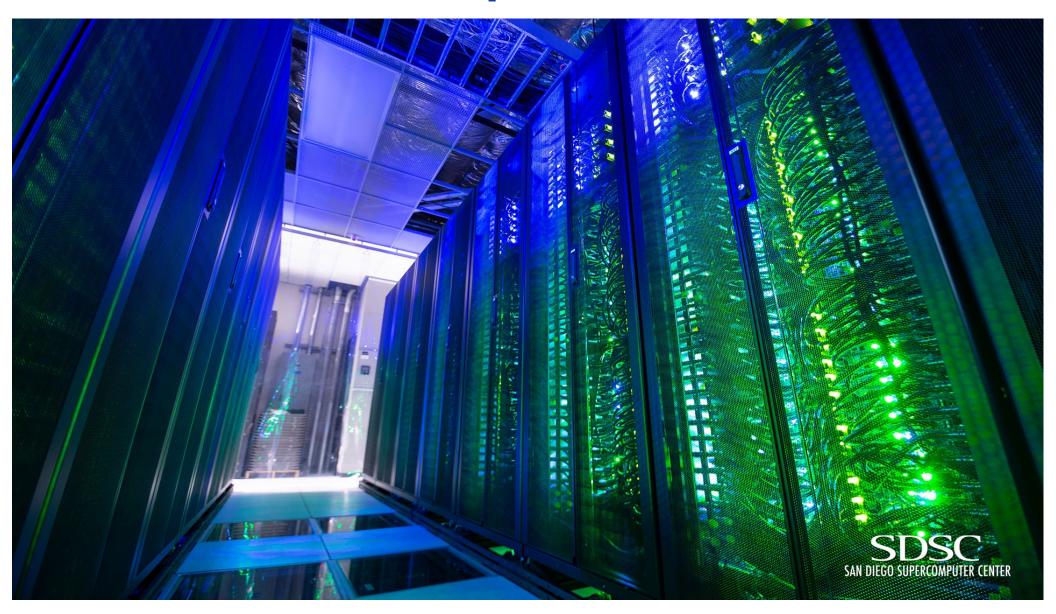
INNOVATIVE OPERATIONS

Composable Systems
High-Throughput Computing
Science Gateways
Interactive Computing
Containerized Computing
Cloud Bursting

For more details see the Expanse user guide @ https://www.sdsc.edu/support/user_guides/expanse.htm| and the "Introduction to Expanse" webinar @ https://www.sdsc.edu/event_items/202006_Introduction_to_Expanse.htm|



Expanse

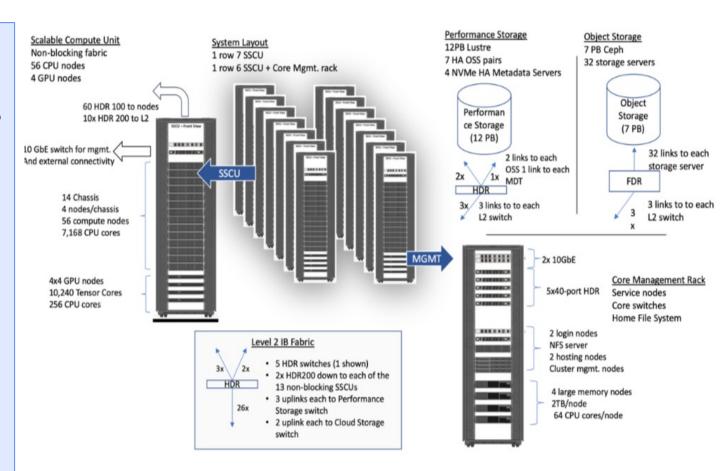




Expanse Heterogeneous Architecture

System Summary

- 13 SDSC Scalable Compute Units (SSCU)
- 728 x 2s Standard Compute Nodes
- 93,184 Compute Cores
- 200 TB DDR4 Memory
- 52x 4-way GPU Nodes w/NVLINK
- 208 V100s
- 4x 2TB Large Memory Nodes
- HDR 100 non-blocking Fabric
- 12 PB Lustre High Performance
- Storage
- 7 PB Ceph Object Storage
- 1.2 PB on-node NVMe
- Dell EMC PowerEdge
- Direct Liquid Cooled





SCCU

The SSCU is Designed for the Long Tail Job Mix, Maximum Performance, Efficient Systems Support, and Efficient Power and Cooling

Standard Compute Nodes

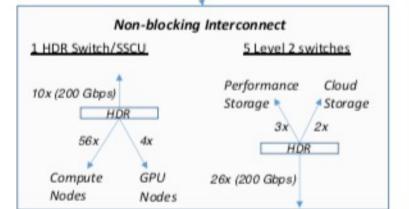
- 2x AMD EPYC 7742 @2.25 GHz
- 128 Zen2 CPU cores
- · PCle Gen4
- 256 GB DDR4
- 1.6 TB NVME

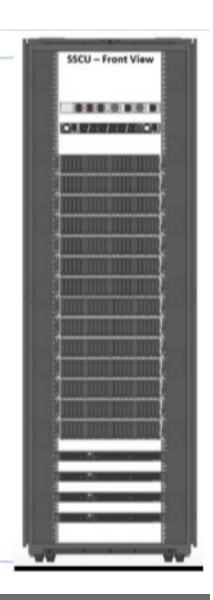
GPU Nodes

- 4x NVIDIA V100/follow-on
- 10,240 Tensor Cores
- 32 GB GDDR
- 1.6 TB NVMe
- Intel CPUs

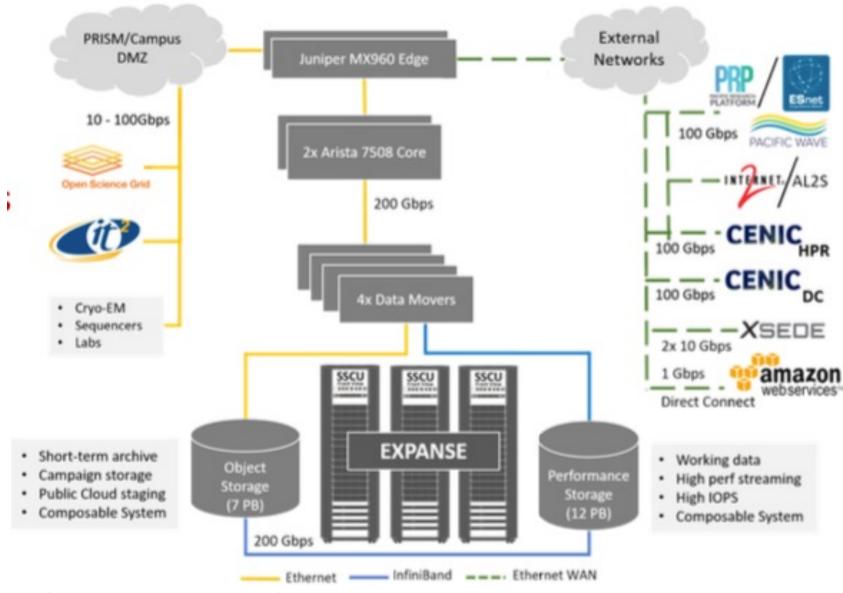
SSCU Components

- 56x CPU nodes
- 7,168 Compute Cores
 - 4x GPU nodes
- 1x HDR Switch
- 1x 10GbE Switch
- HDR 100 non-blocking fabric
- · Wide rack for serviceability
- Direct Liquid Cooling to CPU nodes





Expanse Connectivity Fabric

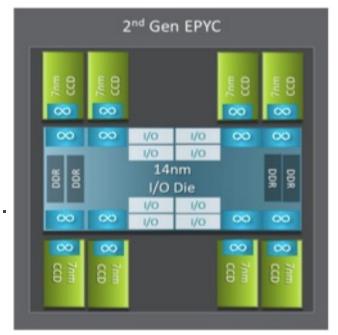


Facilitates Compute and Data Workflows



AMD EPYC 7742 Processor Architecture

- 8 Core Complex Dies (CCDs).
- CCDs connect to memory, I/O, and each other through the I/O Die.
- 8 memory channels per socket.
- DDR4 memory at 3200MHz.
- PCI Gen4, up to 128 lanes of high speed I/O.
- Memory and I/O can be abstracted into separate quadrants each with 2 DIMM channels and 32 I/O lanes.
- 2 Core Complexes (CCXs) per CCD
- 4 Zen2 cores in each CCX share a16ML3 cache. Total of 16x16= 256MB L3 cache.
- Each core includes a private 512KB L2 cache.



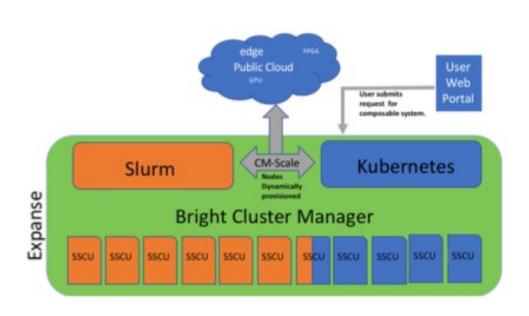


→ EPYC Architecture has impact on compiling and batch script configuration



Composable Systems will support complex, distributed, workflows – making Expanse part of a larger CI ecosystem

- Bright Cluster Manager + Kubernetes
- Core components developed via NSF- funded CHASE-CI (NSF Award # 1730158), and the Pacific Research Platform (NSF Award # 1541349)
- Requests for a composable system will be part of an XRAC request
- Advanced User Support resources available to assist with projects - this is part of our operations funding.





Integration with Public Cloud *

- Supports projects that share data, need access to novel technologies, and integrate cloud resources into workflows
- Slurm + in-house developed software + Terraform (Hashicorp)
- Early work funded internally and via NSF E-CAS/Internet2 project for CIPRES (Exploring Cloud for the Acceleration of Science, Award #1904444).
- Approach is cloud-agnostic and will support the major cloud providers.
- Users submit directly via Slurm, or as part of a composed system.
- Options for data movement: data in the cloud; remote mounting of file systems; cached filesystems (e.g., StashCache), and data transfer during the job.
- Training Events Planned for early 2021

^{*} Funding for user cloud resources is not part of the Expanse award. Researcher must have access to these via other NSF awards and funding.



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Logging into Expanse

- Expanse supports Single Sign-On through the XSEDE User Portal
- From the command line using an XSEDE-wide password,
 - Coming soon the Expanse User Portal.
- CPU and GPU resources are allocated separately, the login nodes are the same.
- To log in to Expanse from the command line, use the hostname:
 - login.expanse.sdsc.edu
- Secure shell (SSH) command examples:

```
ssh <your_username>@login.expanse.sdsc.edu
ssh -l <your_username> login.expanse.sdsc.edu
```

• When you log in to *login.expanse.sdsc.edu*, you will be assigned one of the two login nodes login0[1-2]-expanse.sdsc.edu. Both systems are identical.



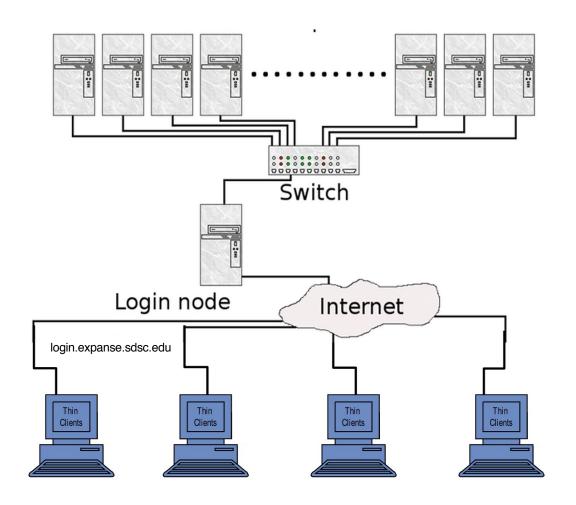
Using SSH Keys

- You can append your public key (e.g.from your laptop) to your ~/.ssh/authorized_keys file to enable access from authorized hosts without having to enter your password.
- RSA, ECDSA and ed25519 keys are accepted.
- Make sure you have a strong passphrase on the private key on your local machine.
- You can use ssh-agent or keychain to avoid repeatedly typing the private key password.
- Hosts which connect to SSH more frequently than ten times per minute may get blocked for a short period of time
- See the SDSC "Indispensable Security: Tips to Use SDSC's HPC Resources Securely"
 - https://www.sdsc.edu/event_items/202007_CometWebinar.html



System Access: Clients

- Linux/Mac
 - use terminal + installed ssh app
- Windows:
 - Win10 terminal app + installed ssh app
 - Older Windows OS's: ssh clients apps Putty, Cygwin
- Login hostname for SDSC Expanse:
 - login.expanse.sdsc.edu
 - 198.202.113.252



SDSC tutorial on HPC security: https://www.sdsc.edu/event_items/202007 CometWebinar.html



Example of a terminal connection:

```
(base) quantum: mthomas$ ssh -l mthomas login.expanse.sdsc.edu
Welcome to Bright release
                                       9.0
                                                                Based on CentOS Linux 8
                                                                              ID: #000002
                                    WELCOME TO
Use the following commands to adjust your environment:
'module avail'

    show available modules

'module add <module>' - adds a module to your environment for this session 'module initadd <module>' - configure module to be loaded at every login
Last login: Mon Jun 21 18:21:08 2021 from 76.176.117.51
[mthomas@login02 ~]$ whoami
mthomas
[mthomas@login02 ~]$ pwd
/home/mthomas
[mthomas@login02 ~]$
[mthomas@login02 ~]$
```

Typically you would also see a logon message – often called the MOTD (message of the day, located in /etc/motd). This has not been implemented at this point on Expanse



Using Login Nodes Properly

- The login nodes are meant for file editing, simple data analysis, & tasks that use minimal compute resources.
- All computationally demanding jobs should be submitted and run through the batch queuing system.
- Do not use the login nodes for:
 - computationally intensive processes,
 - as hosts for running workflow management tools
 - as primary data transfer nodes for large or numerous data transfers
 - as servers providing other services accessible to the Internet.
 - running Jupyter notebooks
- Login nodes are not the same as the batch nodes.
 - Users should request an interactive sessions to compile ;arge programs.



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Expanse Environment Modules

- Expanse uses Lmod, a Lua based module system.
 - https://lmod.readthedocs.io/en/latest/010_user.html
- Users setup custom environments by loading available modules into the shell environment, including needed compilers and libraries and the batch scheduler.
- What modules let you do:
 - Dynamic modification of your shell environment
 - User can set, change, or delete environment variables
 - User chooses between different versions of the same software or different combinations of related codes.



Modules on Expanse

- Users will need to load the scheduler (e.g. slurm)
- Users will not see all available modules when they run command "module available" without loading a compiler.
- Use the command "module spider" option to see if a particular package exists and can be loaded, run command

```
module spider <package> module keywords <term>
```

 For additional details, and to identify module dependencies modules, use the command

```
module spider <application_name>
```

 The module paths are different for the CPU and GPU nodes. Users can enable the paths by loading the following modules:

```
module load cpu (for cpu nodes) module load gpu (for gpu nodes)
```

Avoid loading both modules



Module Command Examples

[mahidhar test@login01 ~]\$ module reset Resetting modules to system default. Reseting \$MODULEPATH back to system default. All extra directories will be removed from **\$MODULEPATH.** [mahidhar_test@login01 ~]\$ module list Currently Loaded Modules:
1) shared 2) slurm/expanse/20.02.3 3) cpu/0.15.4 4) DefaultModules List Current environment: list, li **Currently Loaded Modules:** [mahidhar test@login01 ~]\$ module avail ------/cm/shared/apps/spack/cpu/lmod/linux-centos8-x86 64/Core -----------bzip2/1.0.8 qcc/7.5.0 intel/19.1.1.217 parallel/20200822 abagus/2018 cmake/3.18.2 libtirpc/1.2.6 pciutils/3.7.0 anaconda3/2020.11 acc/9.2.0 aocc/2.2.0 curl/7.72.0 gcc/10.2.0 (D) matlab/2020b aria2/1.35.0 emboss/6.6.0 gmp/6.1.2 mpfr/4.0.2 curl/7.72.0 gcc/10.2.0 (D) matlab/2020b pigz/2.4 subversion/1.14.0 arm-forge/21.0.1-linux-x86 64 gaussian/16.C.01 go/1.15.1 nbo/7.0-openblas zstd/1.4.5 Show bvacc/master gaussian09/09.E.01 idl/8.4 openjdk/11.0.2 available -----/cm/local/modulefiles -----modules boost/1.71.0 cmjob lua/5.3.5 shared (L) singularitypro/3.5 slurm/expanse/20.02.3 (L) ------/cm/shared/apps/xsede/modulefiles ------/cm/shared/apps/xsede/modulefiles cue-login-env xdinfo/1.5-1 xdusage/2.1-1 ------/usr/share/modulefiles ------DefaultModules (L) cpu/0.15.4 (L) qct/6.2 qlobus/6.0 qpu/0.15.4 nostack/0.15.4



Modules: Popular commands

Command	Description	
module list	List the modules that are currently loaded	
module avail	List the modules that are available in environment	
module spider	List of the modules and extensions currently available	
module display <module_name></module_name>	Show the environment variables used by <module name=""> and how they are affected</module>	
module unload <module name=""></module>	Remove <module name=""> from the environment</module>	
module load <module name=""></module>	Load <module name=""> into the environment</module>	
module swap <module one=""> <module two=""></module></module>	Replace <module one=""> with <module two=""> in the environment</module></module>	
module help	get a list of all the commands that module knows about do:	
Shorthand notation: ml foo	"ml" == module load foo	
ml -bar	"ml -bar" == module unload bar	

SDSC Guidance: add module calls to your environment and batch scripts



Module Command Examples

Use "module show" to find out what a particular module will change in the environment

[[mahidhar test@login01 ~]\$ module show cmake

/cm/shared/apps/spack/cpu/lmod/linux-centos8-x86_64/Core/cmake/3.18.2.lua:

whatis("Name : cmake") whatis("Version : 3.18.2") whatis("Target : zen")

whatis("Short description : A cross-platform, open-source build system. CMake is a family of tools designed to build, test and package

software. ")

help([[A cross-platform, open-source build system. CMake is a family of tools

designed to build, test and package software.]])

prepend_path("PATH","/cm/shared/apps/spack/cpu/opt/spack/linux-centos8-zen/gcc-8.3.1/cmake-3.18.2-rfzatdti4qlsrf2zezwad75fnccy4f7d/bin")

prepend_path("ACLOCAL_PATH","/cm/shared/apps/spack/cpu/opt/spack/linux-centos8-zen/gcc-8.3.1/cmake-3.18.2-

rfzatdti4qlsrf2zezwad75fnccy4f7d/share/aclocal")

prepend_path("CMAKE_PREFIX_PATH","/cm/shared/apps/spack/cpu/opt/spack/linux-centos8-zen/gcc-8.3.1/cmake-3.18.2-

rfzatdti4qlsrf2zezwad75fnccy4f7d/")

setenv("CMAKEHOME","/cm/shared/apps/spack/cpu/opt/spack/linux-centos8-zen/gcc-8.3.1/cmake-3.18.2-rfzatdti4qlsrf2zezwad75fnccy4f7d")

[mahidhar_test@login01 ~]\$



Module: check Environment

Once you have loaded the modules, you can check the system variables that are available for you to use.

[mahidhar_test@login01 ~]\$ module load cmake
[mahidhar_test@login01 ~]\$ echo \$PATH
/cm/shared/apps/spack/cpu/opt/spack/linux-centos8-zen/gcc-8.3.1/cmake-3.18.2rfzatdti4qlsrf2zezwad75fnccy4f7d/bin:/cm/shared/apps/slurm/current/sbin:/cm/shared/apps/slurm/
current/bin:/home/mahidhar_test/.local/bin:/home/mahidhar_test/bin:/usr/local/bin:/usr/local/sbin:/usr/sbin:/opt/dell/srvadmin/bin
[mahidhar_test@login01 ~]\$
[mahidhar_test@login01 ~]\$ echo \$LD_LIBRARY_PATH
/cm/shared/apps/slurm/current/lib64/slurm:/cm/shared/apps/slurm/current/lib64
[mahidhar_test@login01 ~]\$
[mahidhar_test@login01 ~]\$ echo \$CMAKEHOME
/cm/shared/apps/spack/cpu/opt/spack/linux-centos8-zen/gcc-8.3.1/cmake-3.18.2rfzatdti4qlsrf2zezwad75fnccy4f7d
[mahidhar_test@login01 ~]\$



Module: command not found

- Sometimes encountered when switching from one shell to another or attempting to run the module command from within a shell script or batch job.
- Module command may not be inherited to the shell
- To keep this from happening, execute the following command:
 - command line (interactive shells)
 - source /etc/profile.d/modules.sh
 - OR add to your shell script (including Slurm batch scripts)



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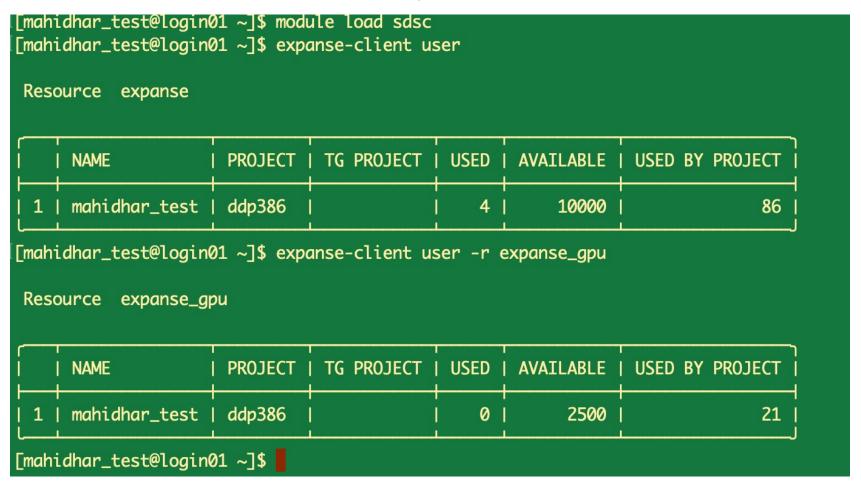
Multiple Allocations

- Many users will have access to multiple accounts and hence projects:
 - an allocation for a research project and a separate allocation for classroom or educational use
- Users should verify that the correct project is designated for all batch jobs.
- Awards are granted for a specific purposes and should not be used for other *projects*.
- To charge your job to one of your projects, replace << project
 >> with one from your list and put this PBS directive in your job script:
 - #SBATCH -A << project >>



Allocation Information

module load sdsc expanse-client user expanse-client user -r expanse_gpu





Charging

- Charge unit for all SDSC machines, including Expanse, is the Service Unit (SU).
- Your charges are based on the resources that are tied up by your job, may not reflect how resources are used.
- Charges are based on either:
 - Number of cores
 - Fraction of the memory requested, whichever is larger.
- The minimum charge for any job is 1 SU. This is important to note - can quickly use up SUs if you run a lot of very short jobs.
- More details in Expanse user guide:

https://www.sdsc.edu/support/user_guides/expanse.html#charging



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Supported Compilers on Expanse

CPU nodes

- GNU, Intel, AOCC (AMD) compilers
- multiple MPI implementations (OpenMPI, MVAPICH2, and IntelMPI).
- A majority of applications have been built using *gcc/10.2.0* which *features AMD Rome* specific optimization flags (-march=znver2).
- Intel, and AOCC compilers all have flags to support Advanced Vector Extensions 2 (AVX2).
- GPU Compiling:
 - Expanse GPU nodes have GNU, Intel, and PGI compilers.
 - Note: Expanse login nodes are not the same as the GPU nodes →
 all GPU codes must be compiled by requesting an interactive
 session on the GPU nodes.



AMD AOCC Compilers: CPU Only

Language	Serial	MPI	OpenMP	MPI + OpenMP
Fortran	flang	mpif90	ifort -openmp	mpif90 -openmp
С	clang	mpiclang	icc -openmp	mpicc -openmp
C++	clang++	mpiclang	icpc -openmp	mpicxx -openm

The AMD Optimizing C/C++ Compiler (AOCC) is only available on CPU nodes. AMD compilers can be loaded using the module load command:

\$ module load aocc

For more information on the AMD compilers:

\$ [flang | clang] -help



Using the AOCC Compilers

- If you have modified your environment, you can reload by executing the module purge & load commands at the Linux prompt, or placing the load command in your startup file (~/.cshrc or ~/.bashrc)
- Note: The examples below are for the simple "hellompi" examples shown below

```
[mthomas@login02 ~]$ module list
Currently Loaded Modules:
 1) shared 2) cpu/1.0 3) DefaultModules 4) hdf5/1.10.1 5) intel/ 19.1.1.217
## need to change multiple modules
[mthomas@login02 ~]$ module purge
[mthomas@login02 ~]$ module list
No modules loaded
[mthomas@login02 ~]$ module load slurm
[mthomas@login02 ~]$ module load cpu
[mthomas@login02 ~]$ module load gcc
[mthomas@login02 ~]$ module load openmpi/4.0.4
[mthomas@login02 ~]$ module list
Currently Loaded Modules:
 1) slurm/expanse/20.02.3 2) cpu/1.0 3) gcc/10.2.0 4) openmpi/4.0.4
[mthomas@login02 MPI]$ module swap intel aocc
Due to MODULEPATH changes, the following have been reloaded:
 1) openmpi/4.0.4
[mthomas@login02 ~]$ module list
Currently Loaded Modules:
 1) slurm/expanse/20.02.3 2) cpu/1.0 3) aocc/2.2.0 4) openmpi/4.0.4
[mthomas@login02 ~]$
```



Intel Compilers: GPU and GPU

 Default/Suggested Compilers to used based on programming model and languages:

	Serial	MPI	OpenMP	MPI + OpenMP
Fortran	ifort	mpif90	ifort -openmp	mpif90 -openmp
С	icc	mpicc	icc -openmp	mpicc -openmp
C++	icpc	mpicxx	icpc -openmp	mpicxx -openmp

- In this tutorial, we include hands-on examples that cover many of the cases in the table:
 - (1) MPI
 - (2) OpenMP
 - (3) HYBRID



Using the Intel Compilers

• If you have modified your environment, you can reload by executing the module purge & load commands at the Linux prompt, or placing the load command in your startup file (~/.cshrc or ~/.bashrc)

[mthomas@login02 ~]\$ module list

Currently Loaded Modules:

1) slurm/expanse/20.02.3 2) cpu/1.0 3) aocc/2.2.0 4) openmpi/4.0.4

[mthomas@login02 ~]\$ module purge

[mthomas@login02 ~]\$ module list

No modules loaded

[mthomas@login02 ~]\$ module load slurm

[mthomas@login02 ~]\$ module load cpu

[mthomas@login02 ~]\$ module load intel

[mthomas@login02 ~]\$ module load openmpi/4.0.4

[mthomas@login02 ~]\$ module list

Currently Loaded Modules:

1) slurm/expanse/20.02.3 2) cpu/1.0 3) aocc/2.2.0 4) openmpi/4.0.4

[mthomas@login02 ~]\$



Using the Intel Compilers

- For Intel compilers, to enable Advanced Vector Extensions (AVX2) support, compile with the -march=core-avx2 option.
 - https://en.wikipedia.org/wiki/Advanced_Vector_Extensions (128/256bit SIMD, Vector ops (MPI broadcast, gather, ...)
 - Note that -march=core-avx2 alone does not enable aggressive optimization, so compilation with -O3 is also suggested.
- Intel Math Kernal Lib (MKL) libraries are available as part of the "intel" modules on Expanse.
 - Once this module is loaded, the environment variable INTEL_MKLHOME points to the location of the mkl libraries and
 - Use MKL Link Advisor to see what libraries are recommended for your compiler and system configuration:
 - https://software.intel.com/content/www/us/en/develop/articles/intel-mkl-link-lineadvisor.html



GNU Compilers: CPU and GPU

• The GNU compilers can be loaded by executing the following commands at the Linux prompt or placing in your startup files (~/.cshrc or ~/.bashrc)

```
[mthomas@login01 MPI]$ module purge
[mthomas@login01 MPI]$ module load slurm
[mthomas@login01 MPI]$ module load cpu
[mthomas@login01 MPI]$ module load gcc/10.2.0
[mthomas@login01 MPI]$ module load openmpi/4.0.4
[mthomas@login01 MPI]$ module list
Currently Loaded Modules:

1) slurm/expanse/20.02.3 2) cpu/1.0 3) gcc/10.2.0 4) openmpi/4.0.4
```

- For AVX support, compile with -mavx.
- Note that AVX support is only available in version 4.7 or later, so it is necessary to explicitly load the gnu/4.9.2 module until such time that it becomes the default.
- For more information on the GNU compilers: man [gfortran | gcc | g++]



Using the GNU Compilers

Table of recommended GNU compilers:

	Serial	MPI	OpenMP	MPI+OpenMP
Fortran	gfortran	mpif90	gfortran -fopenmp	mpif90 -fopenmp
С	gcc	mpicc	gcc -fopenmp	mpicc -fopenmp
C++	g++	mpicxx	g++ -fopenmp	mpicxx -fopenmp



PGI Compilers: GPU Only

- PGI (formerly The Portland Group, Inc.), was a company that produced a set of commercially available Fortran, C and C++ compilers for highperformance computing systems.
- It is now owned by NVIDIA.
- To compile code, you need to obtain an interactive node.
- For AVX support, compile with -fast

```
[mahidhar_test@login01 ~]$ module reset
[mahidhar_test@login01 ~]$ module load gpu
[mahidhar_test@login01 ~]$ module load pgi
[mahidhar_test@login01 ~]$
[mahidhar_test@login01 ~]$ which pgcc
/cm/shared/apps/spack/gpu/opt/spack/linux-centos8-skylake_avx512/gcc-8.3.1/pgi-20.4-
2tsjnv2icisxmgdy4mijl4t5mkbr32ea/linux86-64/20.4/bin/pgcc
[mahidhar_test@login01 ~]$ which mpicc
/cm/shared/apps/spack/gpu/opt/spack/linux-centos8-skylake_avx512/gcc-8.3.1/pgi-20.4-
2tsjnv2icisxmgdy4mijl4t5mkbr32ea/linux86-64/20.4/mpi/openmpi-3.1.3/bin/mpicc
```

For more information on the PGI compilers run: man [pgf90 | pgcc | pgCC]



Recommended PGI Compilers

	Serial	MPI	OpenMP	MPI+OpenMP
Fortran	pgf90	mpif90	pgf90 -mp	mpif90 -mp
С	pgcc	mpicc	pgcc -mp	mpicc -mp
C++	pgCC	mpicxx	pgCC -mp	mpicxx -mp

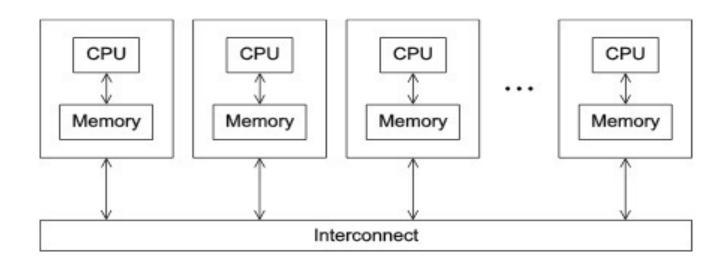
- PGI supports the following high-level languages:
 - Fortran 77, 90/95/2003, 2008 (partial)
 - High Performance Fortran (HPF)
 - ANSI C99 with K&R extensions
 - ANSI/ISO C++
 - CUDA Fortran
 - OpenCL
 - OpenACC
 - OpenMP



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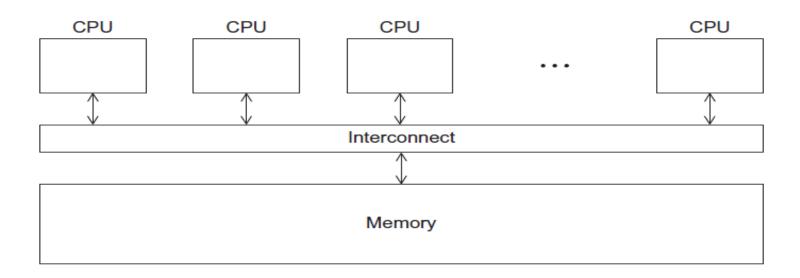
Parallel Models: Distributed Memory



- Programs that run asynchronously, pass messages for communication and coordination between resources.
- Examples include: SOA-based systems, massively multiplayer online games, peer-to-peer apps.
- Different types of implementations for the message passing mechanism: HTTP, RPC-like connectors, message queues
- HPC historically uses the Message Passing Interface (MPI)



Parallel Models: Shared Memory



- CPUs all share same localized memory (SHMEM);
 - Coordination and communication between tasks via interprocessor communication (IPC) or virtual memory mappings.
- May use: uniform or non-uniform memory access (UMA or NUMA); cache-only memory architecture (COMA).
- Most common HPC API's for using SHMEM:
 - Portable Operating System Interface (POSIX); Open Multi-Processing (OpenMP) designed for parallel computing best for multi-core computing.



Running Jobs on Expanse

- When you run in the batch mode, you submit jobs to be run on the compute nodes using the sbatch command as described below.
- Remember that computationally intensive jobs should be run only on the compute nodes and not the login nodes.
- Expanse places limits on the number of jobs queued and running on a per group (allocation) and partition basis.
- Please note that submitting a large number of jobs (especially very short ones) can impact the overall scheduler response for all users.



Methods for Runing Jobs on Expanse

- Expanse uses the Simple Linux Utility for Resource Management (SLURM) batch environment.
 - Batch Jobs: Submit batch scripts to Slurm from the login nodes:
 - Partition (queue)
 - Time limit for the run (maximum of 48 hours)
 - Number of nodes, tasks per node; Memory requirements (if any)
 - Job name, output file location; Email info, configuration
- Interactive Jobs: Use the srun command to obtain nodes for 'live,' command line interactive access:
 - CPU:

```
srun --partition=debug --account=XYZ123 --pty --nodes=1 --ntasks-per-node=128 --mem=248 -t 00:30:00 --wait=0 --export=ALL /bin/bash
```

GPU:

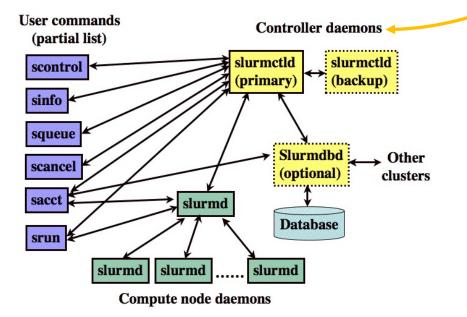
```
srun --pty --account=XYZ123 --nodes=1 --ntasks-per-node=1 --cpus-per-task=10 -p gpu-debug --gpus=1 -t 00:10:00 /bin/bash
```

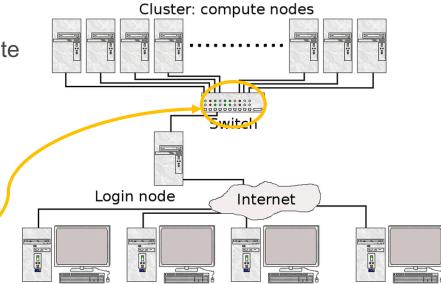


Slurm Resource Manager

Simple Linux Utility for Resource Management

- "Glue" for parallel computer to schedule and execute jobs
- Role: Allocate resources within a cluster
 - Nodes (unique IP address)
 - Interconnect/switches
 - Generic resources (e.g. GPUs)
 - Launch and otherwise manage jobs





Users, submitting jobs

- Functionality:
 - Prioritize queue(s) of jobs;
 - decide when and where to start jobs;
 - terminate job when done;
 - Appropriate resources;
 - manage accounts for jobs

Slurm Partitions on Expanse

Partition limits subject to change based on Early User Period evaluation

Partition Name	QOS	Max Walltime	Max Nodes/Job	Max RunningJobs	Max Running + Queued Jobs	Charge Factor	Comments
compute	normal	48 hrs	32	64	128	1	Used for exclusive access to regular compute nodes
shared	shared- normal	48 hrs	1	4096	4096	1	Single-node jobs using fewer then 128 cores
gpu	gpu-normal	48 hrs	4	16	24	1	Used for exclusive access to the GPU nodes
gpu-shared	gpu-shared- normal	48 hrs	1	16	24	1	Single-node job using fewer then 4 GPUs
large-shared	large-shared- normal	48 hrs	1	1	4	1	Single-node jobs using large memory up to 2 TB (minimum memory required 256G)
debug	debug- normal	15 min	2	1	2	1	Priority access to compute nodes set aside for testing of jobs with short walltime and limited resources
gpu-debug	gpu-debug- normal	15 min	2	1	2	1	** Priority access to gpu nodes set aside for testing of jobs with short walltime and limited resources
preempt	preempt- normal	7 days	32		128	.8	Discounted jobs to run on free nodes that can be pre-empted by jobs submited to any other queue (NO REFUNDS)
preempt-gpu	preempt- gpu-normal	7 days	1			.8	Discounted jobs to run on unallocated nodes that can be pre-empted by jobs submitted to higher priority queues (NO REFUNDS)

Common Slurm Commands

Submit jobs using the sbatch command:

```
$ sbatch mycode-slurm.sb
```

Submitted batch job 8718049

Check job status using the squeue command:

```
$ squeue -u $USER

JOBID PARTITION NAME USER ST TIME NODES NODELIST(REASON)

8718049 compute mycode user PD 0:00 1 (Priority)
```

Once the job is running:

```
$ squeue -u $USER

JOBID PARTITION NAME USER ST TIME NODES NODELIST(REASON)

8718049 debug mycode user R 0:02 1 expanse-14-01
```

Cancel a running job:

\$ scancel 8718049



Example Batch Script

```
[mthomas@login01 ENV INFO]$ cat env-slurm.sb
#!/bin/bash
#SBATCH --job-name="envinfo"
#SBATCH --output="envinfo.%j.%N.out"
#SBATCH --partition=compute
#SBATCH --nodes=1
#SBATCH --ntasks-per-node=1
#SBATCH --export=ALL
#SBATCH -t 00:01:00
## Environment
module purge
module load slurm
module load cpu
## perform some basic unix commands
echo "-----"
echo "hostname= " `hostname`
echo "date= " `date`
echo "whoami= " `whoami`
echo "pwd= " `pwd`
echo "module list= " `module list`
echo "-----"
echo "env= " `env`
echo "-----"
```

Simple batch script showing environment, date, etc.

```
[mthomas@login01 ENV INFO]$ cat envinfo.108867.exp-6-56.out
hostname= exp-6-56
date= Wed Oct 7 23:45:43 PDT 2020
whoami= mthomas
pwd= /home/mthomas/DEMO/ENV INFO
Currently Loaded Modules:
1) slurm/expanse/20.02.3 2) cpu/1.0
_____
env= SLURM MEM PER CPU=1024
LD LIBRARY PATH=/cm/shared/apps/slurm/current/lib64/slurm:/cm/shared/ap
ps/slurm/current/lib64 LS COLORS=rs=0
[SNIP]
MODULESHOME=/usr/share/Imod/Imod LMOD SETTARG FULL SUPPORT=no
HISTSIZE=5000 LMOD_PKG=/usr/share/Imod/Imod
LMOD CMD=/usr/share/Imod/Imod/libexec/Imod SLURM LOCALID=0
LESSOPEN=||/usr/bin/lesspipe.sh %s LMOD_FULL_SETTARG_SUPPORT=no
LMOD DIR=/usr/share/Imod/Imod/libexec BASH FUNC module%%=() { eval
$($LMOD CMD bash "$@") && eval $(${LMOD SETTARG CMD:-:} -s sh) }
BASH_FUNC_ml%%=() { eval $($LMOD_DIR/ml_cmd "$@") } _=/usr/bin/env
```



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Hands On Examples

- Clone examples:
 - https://github.com/sdsc-hpc-training-org/expanse-101.git
- CPU:
 - OpenMP
 - MPI
 - HYBRID
- GPU
- Large Memory Nodes



General Steps: Compiling/Running Jobs

- Change to a working directory (for example the expanse101 directory):
 cd /home/\$USER/expanse101/MPI
- Verify that the correct modules loaded:

```
module list
```

Currently Loaded Modulefiles:

- 1) slurm/expanse/20.02.3 2) cpu/1.0 3) gcc/10.2.0 4) openmpi/4.0.4
- Compile the MPI hello world code: mpif90 -o hello mpi hello mpi.f90
- Verify executable has been created (check that date):
 Is -It hello_mpi
 -rwxr-xr-x 1 user sdsc 721912 Mar 25 14:53 hello_mpi
- Submit job sbatch hello mpi slurm.sb



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MPI Hello World

Change to the MPI examples directory:

```
[mthomas@login01 MPI]$ cat hello_mpi.f90
! Fortran example
program hello
include 'mpif.h'
integer rank, size, ierror, tag, status(MPI_STATUS_SIZE)

call MPI_INIT(ierror)
call MPI_COMM_SIZE(MPI_COMM_WORLD, size, ierror)
call MPI_COMM_RANK(MPI_COMM_WORLD, rank, ierror)
print*, 'node', rank, ': Hello world!'
call MPI_FINALIZE(ierror)
end
[mthomas@login01 MPI]$
```

MPI Hello World: Compile

Set the environment and then compile the code

[mthomas@login01 MPI]\$ cat README.txt [1] Compile:

Load module environment

module purge module load slurm module load cpu module load gcc/10.2.0 module load openmpi/4.0.4

mpif90 -o hello_mpi hello_mpi.f90

[2a] Run using Slurm:

sbatch hellompi-slurm.sb

[2b] Run using Interactive CPU Node

```
srun --partition=debug --account=sds184 --pty --nodes=1 --ntasks-per-
node=128 --mem=248 -t 00:30:00 --wait=0 --export=ALL /bin/bash
```

[mthomas@login01 MPI]\$ module list

Currently Loaded Modules:

1) cpu/1.0 2) slurm/expanse/20.02.3

[mthomas@login01 MPI]\$ module purge [mthomas@login01 MPI]\$ module load slurm [mthomas@login01 MPI]\$ module load cpu [mthomas@login01 MPI]\$ module load gcc/10.2.0 [mthomas@login01 MPI]\$ module load openmpi/4.0.4 [mthomas@login01 MPI]\$ module list

Currently Loaded Modules:

1) slurm/expanse/20.02.3 2) cpu/1.0 3) gcc/10.2.0 4) openmpi/4.0.4

[mthomas@login01 MPI]\$ mpif90 -o hello_mpi hello_mpi.f90 [mthomas@login01 MPI]\$

MPI Hello World: Batch Script

- To run the job, use the batch script submission command.
- Monitor the job until it is finished using the squeue command.

```
[mthomas@login01 MPI]$ cat hellompi-slurm-gnu.sb
#!/bin/bash
#SBATCH --job-name="hellompi-gnu"
#SBATCH --output="hellompi-gnu.%j.%N.out"
#SBATCH --partition=compute
#SBATCH --nodes=2
#SBATCH --ntasks-per-node=128
#SBATCH --export=ALL
#SBATCH -t 00:10:00
#This job runs with 2 nodes,
128 cores per node for a total of 256 cores.
## Environment
module purge
module load slurm
module load cpu
module load gcc/10.2.0
module load openmpi/4.0.4
## Use srun to run the job
srun --mpi=pmi2 -n 256 --cpu-bind=rank ./hello_mpi_gnu
[mthomas@login01 MPI]$
```

```
[mthomas@login01 MPI]$ sbatch hellompi-slurm-gnu.sb; squeue -u mthomas
Submitted batch job 108910
      JOBID PARTITION NAME USER ST
                                           TIME NODES NODELIST(REASON)
      108910 compute hellompi mthomas PD
                                              0:00 2 (None)
[mthomas@login01 MPI]$ cat hellompi-gnu.108910.exp-12-54.out
          4: Hello world!
node
node
          5: Hello world!
          7 : Hello world!
node
node
          0: Hello world!
node
          2: Hello world!
          3: Hello world!
node
          9: Hello world!
node
node
          10: Hello world!
[SNIP]
node
         247: Hello world!
         248: Hello world!
node
         249: Hello world!
node
node
         186: Hello world!
         220: Hello world!
node
         203: Hello world!
node
         135 : Hello world!
node
```



Using An Interactive mode

```
[mthomas@login01 MPI]$ module purge
[mthomas@login01 MPI]$ module load slurm
[mthomas@login01 MPI]$ module load cpu
[mthomas@login01 MPI]$ module load gcc/10.2.0

[mthomas@login01 MPI]$ module load openmpi/4.0.4

[mthomas@login01 MPI]$ srun --partition=debug --account=sds184 --pty --nodes=1 --ntasks-per-node=128 --mem=248 -t

00:30:00 --wait=0 --export=ALL /bin/bash
[mthomas@exp-9-55 MPI]$
```

```
[mthomas@exp-9-55 MPI]$ mpirun -np 16 ./hello mpi
           1 : Hello world!
node
          15: Hello world!
node
node
          7 : Hello world!
node
          14: Hello world!
          11: Hello world!
node
node
          6: Hello world!
          4: Hello world!
node
          5: Hello world!
node
          12: Hello world!
node
          13: Hello world!
node
node
          0 : Hello world!
          8: Hello world!
node
node
          9: Hello world!
node
          10: Hello world!
          2: Hello world!
node
           3: Hello world!
node
```

- Exit interactive session when your work is done or you will be charged CPU time.
- Beware of oversubscribing your job: asking for more cores than you have. Intel compiler allows this, but your performance will be degraded.

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OpenMP Hello World

Change to the OPENMP examples directory:

```
[mthomas@login01 examples]$ cd OPENMP/
[mthomas@login01 OPENMP]$ II
total 89
drwxr-xr-x 2 mthomas use300 7 Oct 7 11:28.
drwxr-xr-x 7 mthomas use300 7 Oct 8 00:03 ...
-rwxr-xr-x 1 mthomas use300 19640 Oct 7 11:28 hello openmp
-rw-r--r-- 1 mthomas use300 236 Oct 7 11:28 hello_openmp.f90
-rw-r--r-- 1 mthomas use300 672 Oct 7 11:28 hello openmp shared.108737.exp-6-56.out
-rw-r--r-- 1 mthomas use300 442 Oct 7 11:28 openmp-slurm-shared.sb
-rw-r--r-- 1 mthomas use300 168 Oct 7 11:28 README.txt
[mthomas@login01 OPENMP]$ cat hello_openmp.f90
   PROGRAM OMPHELLO
  INTEGER TNUMBER
  INTEGER OMP GET THREAD NUM
!$OMP PARALLEL DEFAULT(PRIVATE)
  TNUMBER = OMP_GET_THREAD_NUM()
  PRINT *, 'HELLO FROM THREAD NUMBER = ', TNUMBER
!$OMP END PARALLEL
   END
```



OpenMP Hello World: Compile (using aocc compiler)

Set the environment and then compile the code

[mthomas@login01 OPENMP]\$ cat README.txt [1] Compile:

#load module environmentmodule module purge module load slurm module load cpu module load aocc

flang -fopenmp -o hello_openmp hello_openmp.f90

[2] Run:

sbatch openmp-slurm-shared.sb

[mthomas@login01 OPENMP]\$

[mthomas@login01 OPENMP]\$ module list

[mthomas@login01 OPENMP]\$ module purge [mthomas@login01 OPENMP]\$ module load slurm [mthomas@login01 OPENMP]\$ module load cpu [mthomas@login01 OPENMP]\$ module load aocc

Currently Loaded Modules:

1) slurm/expanse/20.02.3 2) cpu/1.0 3) aocc/2.2.0

[mthomas@login01 OPENMP]\$

[mthomas@login01 MPI]\$ mpif90 -o hello_mpi hello_mpi.f90



OpenMP Hello World: Controlling #Threads

A key issue when running OpenMP code is controlling thread behavior. If you run from command line, it will work, but it is not recommended because you will be using Pthreads, which automatically picks the number of threads - in this case 24.

To control thread behavior, there are several key environment variables: OMP_NUM_THREADS controls the number of threads allowed, and OMP_PROC_BIND binds threads to "places" (e.g. cores) and keeps them from moving around (between cores).

```
[expanse-ln2:~/expanse1010PE/NMP] export OMP_NUM_THREADS=4; ./hello_openmp
HELLO FROM THREAD NUMBER = 3
HELLO FROM THREAD NUMBER = 1
HELLO FROM THREAD NUMBER = 2
HELLO FROM THREAD NUMBER = 0
```

See: https://www.ibm.com/support/knowledgecenter/SSGH2K_13.1.3/com.ibm.xlc1313.aix.doc/compiler_ref/ruomprun.html



OpenMP Hello World: Batch Script

```
[mthomas@login01 OPENMP]$ cat openmp-slurm-shared.sb
#!/bin/bash
#SBATCH --job-name="hell openmp shared"
#SBATCH --output="hello openmp shared.%j.%N.out"
#SBATCH --partition=shared
#SBATCH --nodes=1
                                                 [expanse-ln2:~/expanse101/OPENMP] cat openmp-slurm-shared.sb
#SBATCH --ntasks-per-node=1
                                                 #!/bin/bash
#SBATCH --cpus-per-task=16
                                                 #SBATCH --job-name="hell_openmp_shared"
#SBATCH --mem=32G
                                                 #SBATCH --output="hello_openmp_shared.%j.%N.out"
#SBATCH --export=ALL
                                                 #SBATCH --partition=shared
#SBATCH -t 01:30:00
                                                 #SBATCH --share
                                                 #SBATCH --nodes=1
                                                 #SBATCH --ntasks-per-node=16
# AOCC environment
                                                 #SBATCH --mem=80G
                                                 #SBATCH --export=ALL
module purge
                                                 #SBATCH -t 01:30:00
module load slurm
module load cpu
                                                 #SET the number of openmp threads
module load aocc
                                                 export OMP_NUM_THREADS=16
#SET the number of openmp threads
                                                 #Run the openmp job
export OMP NUM THREADS=16
                                                 ./hello_openmp
#Run the openmp job
./hello openmp
[mthomas@login01 OPENMP]$
```

- Note: Expanse supports shared-node jobs (more than one job on a single node).
- Many applications are serial or can only scale to a few cores.
- Shared nodes improve job throughput, provide higher overall system utilization, and allow more users to run on nodes.



OpenMP Hello World: submit job & monitor

To run the job, type the batch script submission command:

```
[mthomas@login01 OPENMP]$ sbatch openmp-slurm-shared.sb; squeue -u mthomas
Submitted batch job 108911
                                              TIME NODES NODELIST(REASON)
       JOBID PARTITION NAME USER ST
      108911 shared hell ope mthomas PD
                                            0:00
                                                   1 (None)
[mthomas@login01 OPENMP]$ II
total 98
drwxr-xr-x 2 mthomas use300
                            8 Oct 8 03:37.
drwxr-xr-x 7 mthomas use300
                          7 Oct 8 00:03 ..
-rwxr-xr-x 1 mthomas use300 19640 Oct 7 11:28 hello openmp
-rw-r--r-- 1 mthomas use300 236 Oct 7 11:28 hello_openmp.f90
-rw-r--r-- 1 mthomas use300 672 Oct 8 03:37 hello openmp shared.108911.exp-6-56 out
-rw-r--r-- 1 mthomas use300 442 Oct 7 11:28 openmp-slurm-shared.sb
                                                                         [mthomas@login01 OPENMP]$ cat
                                                                         hello_openmp_shared.108911.exp-6-56.out
-rw-r--r-- 1 mthomas use300 168 Oct 7 11:28 README.txt
                                                                         HELLO FROM THREAD NUMBER =
                                                                                                            7
                                                                         HELLO FROM THREAD NUMBER =
                                                                                                            6
                                                                                                            12
                                                                         HELLO FROM THREAD NUMBER =
                                                                         HELLO FROM THREAD NUMBER =
                                                                                                            10
                                                                         HELLO FROM THREAD NUMBER =
                                                                                                            1
                                                                         HELLO FROM THREAD NUMBER =
                                                                         HELLO FROM THREAD NUMBER =
                                                                                                            0
                                                                         HELLO FROM THREAD NUMBER =
                                                                         HELLO FROM THREAD NUMBER =
                                                                                                            4
                                                                         HELLO FROM THREAD NUMBER =
                                                                         HELLO FROM THREAD NUMBER =
                                                                                                            13
                                                                         HELLO FROM THREAD NUMBER =
                                                                                                            15
                                                                         HELLO FROM THREAD NUMBER =
                                                                                                           14
                                                                         HELLO FROM THREAD NUMBER =
                                                                                                           11
                                                                         HELLO FROM THREAD NUMBER =
```



8

HELLO FROM THREAD NUMBER =

[mthomas@login01 OPENMP]\$

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Hybrid MPI + OpenMP Hello World

```
#include <stdio.h>
#include "mpi.h"
#include <omp.h>
int main(int argc, char *argv[]) {
  int numprocs, rank, namelen;
 char processor name[MPI MAX PROCESSOR NAME];
  int iam = 0, np = 1;
 MPI Init(&argc, &argv);
 MPI Comm size(MPI COMM WORLD, &numprocs);
 MPI Comm rank(MPI COMM WORLD, &rank);
 MPI Get processor name(processor name, &namelen);
 #pragma omp parallel default(shared) private(iam, np)
   np = omp get num threads();
    iam = omp get thread num();
    printf("Hello Webinar particpants from thread %d out of %d from process %d out of
%d on %s\n",
           iam, np, rank, numprocs, processor name);
 MPI Finalize();
```

Hybrid MPI + OpenMP Jobs

```
[mthomas@login01 HYBRID]$ cat README.txt

[1] Compile:

# Load module environment
module purge
module load slurm
module load cpu
module load intel
module load intel-mpi

export I_MPI_CC=icc
mpicc -qopenmp -o hello_hybrid hello_hybrid.c

[2] Run:
sbatch hybrid-slurm.sb
```

```
[mthomas@login01 HYBRID]$ cat hybrid-slurm.sb
#!/bin/bash
#SBATCH --job-name="hellohybrid"
#SBATCH --output="hellohybrid.%j.%N.out"
#SBATCH --partition=shared
#SBATCH --nodes=1
#SBATCH --ntasks-per-node=2
#SBATCH --cpus-per-task=16
#SBATCH -B 2:16:1
#SBATCH --export=ALL
#SBATCH -t 01:30:00
# Load Module Environment
module purge
module load slurm
module load cpu
module load intel
module load intel-mpi
#Run
export OMP NUM THREADS=16
mpirun -genv I MPI PIN DOMAIN=omp:compact ./hello hybrid
```

```
[mthomas@login01 HYBRID]$ mpicc -qopenmp -o hello_hybrid hello_hybrid.c
[mthomas@login01 HYBRID]$ sbatch hybrid-slurm.sb

Submitted batch job 108875
[mthomas@login01 HYBRID]$ squeue -u mthomas

JOBID PARTITION NAME USER ST TIME NODES NODELIST(REASON)
108875 shared hellohyb mthomas PD 0:00 1 (None)
[mthomas@login01 HYBRID]$
```



Hybrid Hello World: Output

Code ran on:

- 1 node,
- 2 cores per node,
- 16 threads per core

```
[expanse-ln2:~/expanse101/HYBRID] cat hellohybrid. 108875.expanse-06-48.out | sort
Hello from thread 0 out of 16 from process 0 out of 2 on exp-6-56
Hello from thread 0 out of 16 from process 1 out of 2 on exp-6-56
Hello from thread 1 out of 16 from process 0 out of 2 on exp-6-56
Hello from thread 1 out of 16 from process 1 out of 2 on exp-6-56
Hello from thread 2 out of 16 from process 0 out of 2 on exp-6-56
Hello from thread 2 out of 16 from process 1 out of 2 on exp-6-56
Hello from thread 3 out of 16 from process 0 out of 2 on exp-6-56
Hello from thread 3 out of 16 from process 1 out of 2 on exp-6-56
Hello from thread 4 out of 16 from process 0 out of 2 on exp-6-56
Hello from thread 4 out of 16 from process 1 out of 2 on exp-6-56
Hello from thread 5 out of 16 from process 0 out of 2 on exp-6-56
Hello from thread 5 out of 16 from process 1 out of 2 on exp-6-56
Hello from thread 6 out of 16 from process 0 out of 2 on exp-6-56
Hello from thread 6 out of 16 from process 1 out of 2 on exp-6-56
Hello from thread 7 out of 16 from process 0 out of 2 on exp-6-56
Hello from thread 7 out of 16 from process 1 out of 2 on exp-6-56
Hello from thread 8 out of 16 from process 0 out of 2 on exp-6-56
Hello from thread 8 out of 16 from process 1 out of 2 on exp-6-56
Hello from thread 9 out of 16 from process 0 out of 2 on exp-6-56
Hello from thread 9 out of 16 from process 1 out of 2 on exp-6-56
Hello from thread 10 out of 16 from process 0 out of 2 on exp-6-56
Hello from thread 10 out of 16 from process 1 out of 2 on exp-6-56
Hello from thread 11 out of 16 from process 0 out of 2 on exp-6-56
Hello from thread 11 out of 16 from process 1 out of 2 on exp-6-56
Hello from thread 12 out of 16 from process 0 out of 2 on exp-6-56
Hello from thread 12 out of 16 from process 1 out of 2 on exp-6-56
Hello from thread 13 out of 16 from process 0 out of 2 on exp-6-56
Hello from thread 13 out of 16 from process 1 out of 2 on exp-6-56
Hello from thread 14 out of 16 from process 0 out of 2 on exp-6-56
Hello from thread 14 out of 16 from process 1 out of 2 on exp-6-56
Hello from thread 15 out of 16 from process 0 out of 2 on exp-6-56
Hello from thread 15 out of 16 from process 1 out of 2 on exp-6-56
```



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Expanse GPU Hardware

GPU Nodes		
GPU Type	NVIDIA V100 SMX2	
Nodes	52	
GPUs/node	4	
CPU Type	Xeon Gold 6248	
Cores/socket	20	
Sockets	2	
Clock speed	2.5 GHz	
Flop speed	34.4 TFlop/s	
Memory capacity	*384 GB DDR4 DRAM	
Local Storage	1.6TB Samsung PM1745b NVMe PCle SSD	
Max CPU Memory bandwidth	281.6 GB/s	



Using GPU Nodes

- GPU nodes are allocated as a separate resource. The conversion rate is (TBD) Expanse Service Units (SUs) to 1 V100 GPU-hour.
- Login nodes are not the same as the GPU nodes:
 - → GPU codes must be compiled by requesting an interactive session on a GPU nodes.
- Batch: GPU nodes can be accessed via either the "gpu" or the "gpu-shared" partitions.
 - #SBATCH -p gpu
 - or #SBATCH -p gpu-shared
- Interactive GPU node:
 - srun --pty --nodes=1 --ntasks-per-node=1 --cpus-per-task=10 -p gpu-debug --gpus=1 -t 00:10:00 /bin/bash



GPU/CUDA: Interactive Node

Change to the OpenACC directory

```
[mthomas@exp-7-59 OpenACC]$ II
total 71
-rw-r--r-- 1 mthomas use300 2136 Oct 7 11:28 laplace2d.c
-rwxr-xr-x 1 mthomas use300 52056 Oct 7 11:28 laplace2d.openacc.exe
-rw-r--r-- 1 mthomas use300 234 Oct 7 11:28 OpenACC.108739.exp-7-57.out
-rw-r--r-- 1 mthomas use300 307 Oct 8 00:21 openacc-gpu-shared.sb
-rw-r--r-- 1 mthomas use300 1634 Oct 7 11:28 README.txt
-rw-r--r-- 1 mthomas use300 1572 Oct 7 11:28 timer.h
```

Obtain an interactive node:

```
[mthomas@login01 OpenACC]$ srun --pty --nodes=1 --ntasks-per-node=1 --cpus-per-task=10 -p gpu-debug --gpus=1 -t 00:10:00 /bin/bash
```



GPU/CUDA: Node Information

Check node configuration:

```
[mthomas@exp-7-59 OpenACC]$ nvidia-smi
Thu Oct 8 03:58:44 2020
 NVIDIA-SMI 450.51.05 Driver Version: 450.51.05 CUDA Version: 11.0
l-----+
            Persistence-M | Bus-Id Disp.A | Volatile Uncorr. ECC |
GPU Name
| Fan Temp Perf Pwr:Usage/Cap| | Memory-Usage | GPU-Util Compute M. |
               | MIG M. |
 0 Tesla V100-SXM2... On | 00000000:18:00.0 Off |
| N/A 32C PO 41W / 300W | OMiB / 32510MiB | 0% Default |
                                N/A |
l Processes:
GPU GI CI PID Type Process name GPU Memory
 No running processes found
[mthomas@exp-7-59 OpenACC]$
```



GPU: Compile on Interactive node

```
[mthomas@login01 OpenACC]$
cat README.txt
[1] Compile Code:
(a) Get an interactive GPU debug node:
module load slurm
srun --pty --nodes=1 --ntasks-per-node=1 --cpus-per-task=10 -p qpu-debug --qpus=1 -t 00:10:00 /bin/bash
(b) On the GPU node:
module purge
module load slurm
module load apu
module load pgi
pgcc -o laplace2d.openacc.exe -fast -Minfo -acc -ta=tesla:cc70 laplace2d.c
Compiler output:
GetTimer:
   20. include "timer.h"
      61, FMA (fused multiply-add) instruction(s) generated
laplace:
   47, Loop not fused: function call before adjacent loop
     Loop unrolled 8 times
     FMA (fused multiply-add) instruction(s) generated
   55, StartTimer inlined, size=2 (inline) file laplace2d.c (37)
[SNIP]
      75, #pragma acc loop gang, vector(4) /* blockldx.y threadldx.y */
     77, #pragma acc loop gang, vector(32) /* blockldx.x threadldx.x */
   88. GetTimer inlined, size=9 (inline) file laplace2d.c (54)
(Exit out of debug node after this)
[2] Run job:
sbatch openacc-gpu-shared.sb
```



GPU: Submit Batch Script on CPU node

```
[mthomas@login01 OpenACC]$ cat openacc-gpu-shared.sb
                                                                               [mthomas@login01 OpenACC]$ cat OpenACC.108915.exp-7-
#!/bin/bash
#SBATCH --job-name="OpenACC"
                                                                               57.out
#SBATCH --output="OpenACC.%j.%N.out"
                                                                               main()
#SBATCH --partition=gpu-shared
                                                                               Jacobi relaxation Calculation: 4096 x 4096 mesh
                                                                                 0.0.250000
#SBATCH --nodes=1
#SBATCH --ntasks-per-node=1
                                                                                100, 0.002397
                                                                                200, 0.001204
#SBATCH --gpus=1
                                                                                300.0.000804
#SBATCH -t 01:00:00
                                                                                400, 0.000603
                                                                                500, 0.000483
#Environment
                                                                                600, 0.000403
module purge
module load slurm
                                                                                700, 0.000345
                                                                                800, 0.000302
module load gpu
                                                                                900, 0.000269
module load pgi
                                                                                total: 1.084470 s
                                                                               [mthomas@login01 OpenACC]$
#Run the job
./laplace2d.openacc.exe
[mthomas@login01 OpenACC]$ sbatch openacc-gpu-shared.sb
```

TIME NODES NODELIST(REASON)

1 (None)

0:00



Submitted batch job 108915

JOBID PARTITION NAME USER ST

108915 gpu-share OpenACC mthomas PD

[mthomas@login01 OpenACC]\$ sbatch openacc-gpu-shared.sb; squeue -u mthomas

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Yes, You are Correct: Running Jobs on HPC Systems is Complex

- Multiple layers of hardware and software affect job performance
- Learn to develop and test in a modular fashion
- Build up a suite of test cases:
 - When things go wrong, make sure you can run simple test cases (HelloWorld).
 - This can eliminate questions about your environment.
- Consider using a code repository
 - When things go wrong, you can get back to a working version
- If you need help/have questions, contact XSEDE help desk:
 - They are very helpful and respond quickly
 - Support users around the world, so they are truly a 7/24 service
 - Avoid wasting your time.



When Things Go Wrong, Check Your User Environment

- Do you have the right modules loaded?
- What software versions do you need?
- Is your code compiled and updated
 - Did you compile it last year? Have the libraries changed?
- Are you running your job from the right location?
 - \$HOME versus \$WORK?



Run jobs from the right location

- Lustre scratch filesystem:
 - /oasis/scratch/expanse/\$USER/temp_project
 - Preferred: Scalable large block I/O)
- Compute/GPU node local SSD storage:
 - /scratch/\$USER/\$SLURM_JOBID
 - Meta-data intensive jobs, high IOPs)
- Lustre projects filesystem:
 - /oasis/projects/nsf
- /home/\$USER:
 - Only for source files, libraries, binaries.
 - Do not use for I/O intensive jobs.



Thank You



Resources

- Expanse User Guide
 - https://www.sdsc.edu/support/user_guides/expanse.html
- GitHub Repo for this webinar: clone code examples for this tutorial – clone example code:
 - https://github.com/sdsc-hpc-training-org/expanse-101
- SDSC Training Resources
 - https://www.sdsc.edu/education and training/training
 - https://github.com/sdsc-hpc-training/webinars
- XSEDE Training Resources
 - https://www.xsede.org/for-users/training
 - https://cvw.cac.cornell.edu/expanse/

