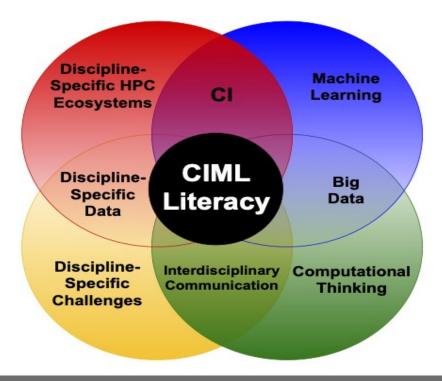
# CIML Summer Institute, Day 1 Day 1: Running Jobs

June 27, 2022

Mary Thomas

San Diego Supercomputer Center







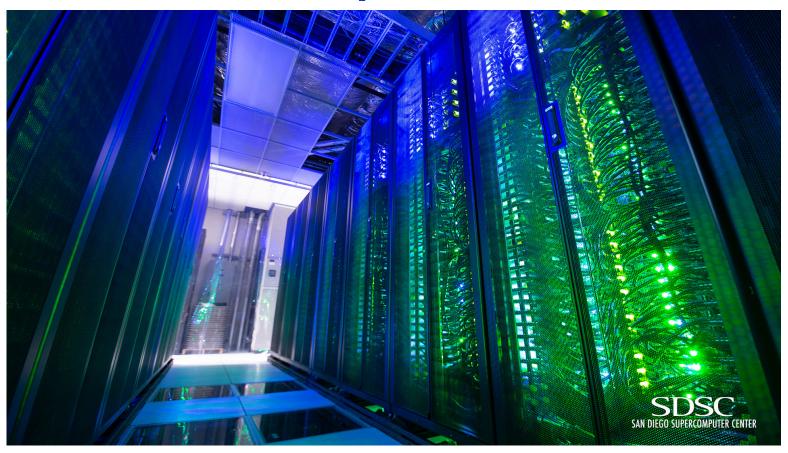
#### **Outline**

- Introduction
- Compiling and Linking Code
- Running Jobs
- Additional Examples
  - MPI Jobs
  - OpenMP Jobs
  - Hybrid MPI-OpenMP Jobs
- Final Comments





# **Expanse**







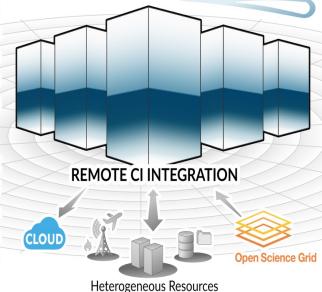
# EXPANSE COMPUTING WITHOUT BOUNDARIES 5 PETAFLOP/S HPC and DATA RESOURCE

#### **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 @ <a href="https://www.sdsc.edu/support/user\_quides/expanse.html">https://www.sdsc.edu/support/user\_quides/expanse.html</a> and the "Introduction to Expanse" webinar @ <a href="https://www.sdsc.edu/event\_items/202006\_Introduction\_to\_Expanse.html">https://www.sdsc.edu/event\_items/202006\_Introduction\_to\_Expanse.html</a>

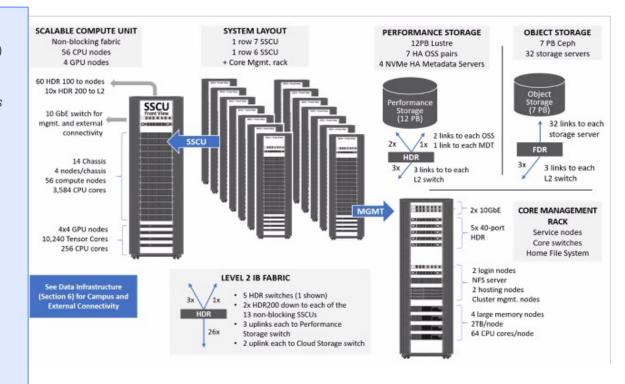




#### **Expanse Heterogeneous Architecture**

#### 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







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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 change your environment by swapping modules or executing the module purge & load commands at the Linux prompt
- Place the load commands in your startup file (~/.cshrc or ~/.bashrc) or batch script

```
[user@login02 ~]$ module list
Currently Loaded Modules:
1) slurm/expanse/21.08.8 2) cpu/0.15.4 3) intel/19.1.1.217 4) openmpi/4.0.4
[user@login02 ~]$ module purge
[user@login02 ~]$ module list
No modules loaded
[user@login02 ~]$ module load slurm
[user@login02 ~]$ module load cpu
[user@login02 ~]$ module load intel
[user@login02 ~]$ module load openmpi/4.0.4
[user@login02 ~]$ module list
Currently Loaded Modules:
1) slurm/expanse/21.08.8 2) cpu/0.15.4 3) intel/19.1.1.217 4) openmpi/4.0.4
[user@login02 MPI]$ module swap intel aocc
Due to MODULEPATH changes, the following have been reloaded:
                                                                                      Loaded the wrong module.
1) openmpi/4.0.4
                                                                                      Use swap
[user@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
[user@login02 ~]$ which mpirun
[mthomas@login01 env info]$ which mpirun
/cm/shared/apps/spack/cpu/opt/spack/linux-centos8-zen/clang-10.0.0/openmpi-4.0.4-nz2g566opuddnwt5ymjorz2cdgb4spfm/bin/mpirun
```





### Intel Compilers: CPU 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

The Intel compilers and the MVAPICH2 MPI compiler wrappers can be loaded by executing the following commands at the Linux prompt:

\$module load intel mvapich2

For more information on the Intel compilers: \$[ifort | icc | icpc] -help





## **Using the Intel Compilers**

[user@login02 ~]\$ module list [user@login02 MPI]\$ 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 [user@login02 ~]\$ module purge [user@login02 ~]\$ module list No modules loaded [mthomas@login01 env\_info]\$ which mpirun /usr/bin/which: no mpirun in (/usr/local/bin:/usr/bin:/usr/local/sbin:/usr/sbin) [user@login02 ~]\$ module load slurm [user@login02 ~]\$ module load cpu [user@login02 ~]\$ module load intel [user@login02 ~]\$ module load openmpi/4.0.4 [user@login02 ~]\$ module list **Currently Loaded Modules:** 1) slurm/expanse/20.02.3 2) cpu/0.15.4 3) intel/19.1.1.217 4) openmpi/4.0.4 [user@login02 ~]\$ which mpirun /cm/shared/apps/spack/cpu/opt/spack/linux-centos8-zen2/intel-19.1.1.217/openmpi-4.0.4-f5mc6sg2jtrw7qqdksf6tru4vo4tawrv/bin/mpirun



[mthomas@login01 env info]\$



### **Accessing Intel Compiler Features**

- Advanced Vector Extensions (AVX2): to enable 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-line-advisor.html





## **GNU Compilers: CPU & GPU**

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

- For AVX support, compile with -march=core-avx2.
- AVX support only available in version 4.7 or later --> explicitly load the gnu/4.9.2 module
- For more information on the GNU compilers:

\$man [gfortran | gcc | g++]





### **GNU Compilers**

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

[user@login02]\$ module purge

[user@login02]\$ module load slurm

[user@login02]\$ module load cpu

[user@login02]\$ module load gcc/10.2.0

[user@login02]\$ module load openmpi/4.0.4

[user@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





#### **Outline**

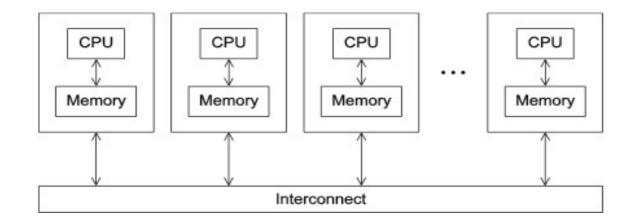
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### Parallel Models: Distributed Memory

- Programs run
   asynchronously, pass
   messages for
   communication and
   coordination between
   resources.
- Examples include: SOAbased systems, massively multiplayer online games, peer-to-peer apps.



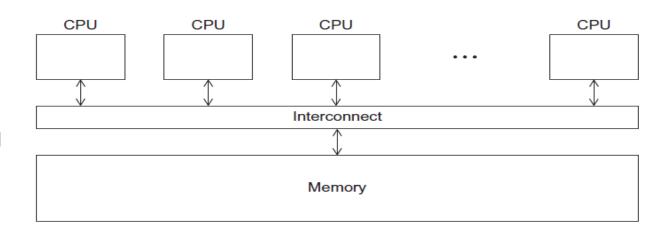
- Different types of implementations for the message passing mechanism:
   HTTP, RPC-like connectors, message queues
- HPC historically uses the Message Passing Interface (MPI)





### **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 nonuniform 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

- "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 Running Jobs on Expanse

- 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
  - Expanse uses the Simple Linux Utility for Resource Management (SLURM) batch environment.
- Interactive Jobs: Use the srun command to request 'live' nodes from Slurm for command line, interactive access
  - See below

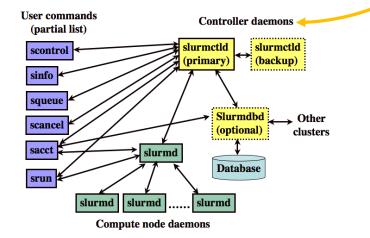


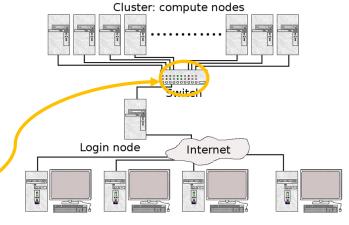


### Slurm Resource Manager

#### 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





#### **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, monitor it's state:

```
$ 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

https://slurm.schedmd.com/sbatch.html





# **Slurm Partitions on Expanse**

https://www.sdsc.edu/support/user\_guides/expanse.html#running

Partition Name	Max Walltim e	Max Nodes/Job	Max Running Jobs	Max Running + Queued Jobs	Charge Factor	Notes
compute	48 hrs	32	32	64	1 1	Used for exclusive access to regular compute nodes; <i>limit</i> applies per group
shared	48 hrs	1	4096	4096	1	Single-node jobs using fewer than 128 cores
gpu	48 hrs	4	4	8 (32 Tres GPU)	1	Used for exclusive access to the GPU nodes
gpu-shared	48 hrs	1	24	24 (24 Tres GPU)	1	Single-node job using fewer than 4 GPUs
large-shared	48 hrs	1	1	4	1	Single-node jobs using large memory up to 2 TB (minimum memory required 256G)
debug	30 min	2	1	2	1	Priority access to shared nodes set aside for testing of jobs with short walltime and limited resources
gpu-debug	30 min	2	1	2	1	Priority access to gpu-shared nodes set aside for testing of jobs with short walltime and limited resources; max two gpus per job
preempt	7 days	32		128	ı ×	Non-refundable discounted jobs to run on free nodes that can be pre-empted by jobs submitted to any other queue
gpu-preempt	7 days	1		24 (24 Tres GPU)		Non-refundable discounted jobs to run on unallocated nodes that can be pre-empted by higher priority queues

https://slurm.schedmd.com/sbatch.html





## **Example Batch Script: Show ENV**

```
[uswername@login02 calc-prime]$ cat mpi-prime-slurm.sb
#!/bin/bash
#SBATCH --job-name="mpi prime"
#SBATCH --output="mpi_prime.%j.%N.out"
####SBATCH --partition=compute
#SBATCH --partition=debug
#SBATCH --nodes=1
#SBATCH --ntasks-per-node=1
#SBATCH --export=ALL
#SBATCH -t 00:10:00
#SBATCH -A use300
## Environment
module purae
module load slurm
module load cpu
module load qcc/10.2.0
module load openmpi/4.0.4
## echo job name and id:
echo "SLURM JOB NAME: $SLURM JOB NAME"
echo "SLURM JOB ID: $SLURM JOB ID"
d=`date`
echo "DATE: $d"
```

```
[mthomas@login02 calc-prime]$ sbatch --export=NLO=1000 mpi-prime-slurm.sb
Submitted batch job 14126259
[mthomas@login02 calc-prime]$ !sq
squeue -u mthomas
JOBID PARTITION NAME USER ST
                                       TIME NODES NODELIST(REASON)
14126259 debug mpi prim mthomas R
                                        0:04
                                              1 exp-9-55
[mthomas@login01 env info]$ cat envinfo.14126259.exp-4-35.out
SLURM JOB NAME: envinfo
SLURM JOB ID: 14126259
hostname= exp-4-35
date= Sun Jun 26 22:05:15 PDT 2022
whoami= mthomas
pwd= /home/mthomas/hpctr-examples/env info
Currently Loaded Modules:
                          1) slurm/expanse/21.08.8 2) cpu/0.15.4
env= LD LIBRARY PATH=/cm/shared/apps/slurm/current/lib64/slurm:
[SNIP]
/cm/shared/apps/slurm/current/lib64
SLURM SUBMIT DIR=/home/mthomas/hpctr-examples/env info
HISTCONTROL=ignoredups
DISPLAY=localhost:16.0
HOSTNAME=exp-4-35
[SNIP]
```





#### **SLURM Environment Variables**

https://slurm.schedmd.com/sbatch.html#lbAJ

Internal ENV vars that exist when job submitted:

#### INPUT ENVIRONMENT VARS

https://slurm.schedmd.com/sbatch.html#lbAJ

- Upon startup, sbatch will read and handle the options set in the following environment variables.
- SBATCH\_JOB\_NAME
  - Same as -J, --job-name
- SBATCH ACCOUNT
  - Same as -A, --account
- SBATCH\_TIMELIMIT
  - Same as -t, --time

#### **OUTPUT ENVIRONMENT VARS**

https://slurm.schedmd.com/sbatch.html#lbAK

- The Slurm controller will set the following variables in the environment of the batch script.
- SLURM\_EXPORT\_ENV
  - Same as --export.
- SLURM\_JOB\_ID
  - The ID of the job allocation.
- SLURM JOB NAME
  - Name of the job.





## Passing values into the batch script

- For SLURM: use the --export flag.
  - For other schedulers check documentation
- Example: pass the value of two variables x and B into the job script named jobscript.sbatch
  - sbatch --export=x=7,B='mystring' jobscript.sb
  - OR: sbatch --export=ALL,x=7,B= mystring 'jobscript.sbatch
- The first example will replace the user's environment with a new environment containing only values for x and B and the SLURM\_\* environment variables. The second will add the values for A and b to the existing environment.





### **Example: Passing Vars to Batch Script**

```
[uswername@login02 calc-prime]$ cat mpi-prime-slurm.sb
     #!/bin/bash
     #SBATCH --job-name="mpi prime"
     #SBATCH --output="mpi prime.%j.%N.out"
     ####SBATCH --partition=compute
     #SBATCH --partition=debug
     #SBATCH --nodes=1
     #SBATCH --ntasks-per-node=24
     #SBATCH --export=ALL
     #SBATCH -t 00:10:00
     #SBATCH -A use300
     ## Environment
     module purge
     module load slurm
     module load cpu
     module load qcc/10.2.0
     module load openmpi/4.0.4
     ## echo job name and id:
     echo "SLURM JOB NAME: $SLURM JOB NAME"
     echo "SLURM JOB ID: $SLURM JOB ID"
     d=`date`
     echo "DATE: $d"
Bat echo "Var NLO: $NLO"
     ## Use srun to run the job, pass variable to code
     srun --mpi=pmi2 -n 24 --cpu-bind=rank ./mpi prime Y
```

```
[mthomas@login02 calc-prime]$ sbatch --export=NLO=1000 mpi-prime-slurm.sb
Submitted batch job 9113467
[mthomas@login02 calc-prime]$ !sq
squeue -u mthomas
JOBID PARTITION NAME USER ST TIME NODES NODELIST(
9113467 debug mpi prim mthomas R
                                      0:04
                                            1 exp-9-55
[mthomas@login02 calc-prime]$ cat mpi prime. 9113467.exp-9-55.out
SLURM JOB NAME: mpi prime SLURM JOB ID: 9113467
DATE: Fri Jan 28 14:54:54 PST 2022
Var NHI: 250000
The argument supplied is 250000
PRIME_MPI. n hi= 250000 C/MPI version
    Ν
          Ρi
                 Time
    1
          0
               0.000361
               0.000004
    4
               0.000768
    8
               0.000003
    16
                0.000003
    32
          11
                0.000003
    64
                0.000003
          18
   128
                 0.000004
           31
   256
                 0.000554
[SNIP]
  16384
           1900
                   0.008385
  32768
           3512
                   0.030101
  65536
           6542
                   0.110856
  131072
          12251
                    0.421177
```



and



### What is Interactive HPC-Computing

- In computer science, interactive computing refers to software which accepts input from the user as it runs.
  - commonly used programs, such as word processors or spreadsheet applications.
- Interactive HPC computing involves real-time user inputs to perform tasks on a set of compute node(s) including:
  - Code development, real-time data exploration, and visualizations.
  - Used when applications have large data sets or are too large to download to local device, software is difficult install, etc.
  - User inputs come via command line interface or application GUI (Jupyter Notebooks, Matlab, R-studio).
  - Actions performed on remote compute nodes as a result of user input or program out.
- Expanse supports running applications via command line and Web-based services (Jupyter Notebooks, Matlab, R-studio)





#### **Accessing Interactive Compute Nodes on Expanse**

- Connect via terminal using SSH → secure connections
- Use the *srun* command to obtain nodes for 'live,' command line interactive access:

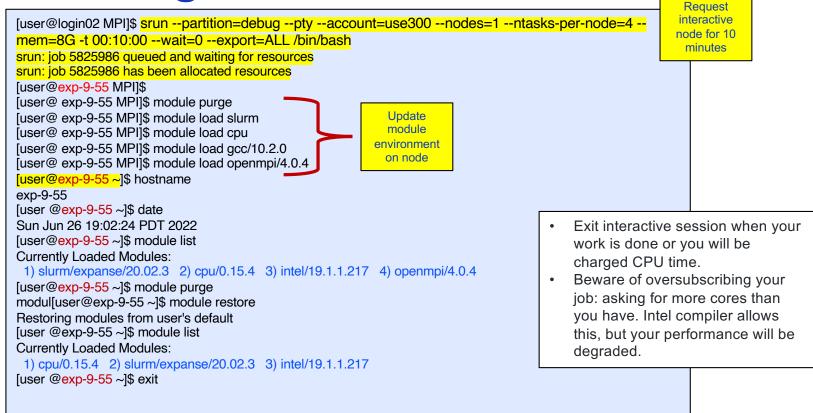
CPU	srunpartition=debugptyaccount=use300nodes=1ntasks-per-node=4mem=8G -t 00:30:00wait=0export=ALL /bin/bash
GPU	srunpartition=gpu-debugptyaccount=use300ntasks-per-node=10nodes=1mem=96Ggpus=1 -t 00:30:00wait=0export=ALL /bin/bash

 Note: you don't need an interactive node to launch secure Jupyter notebooks. Use the Portal or the SRPS/galyleo





## **Using An Interactive mode**







### **Running Interactive Jobs - CPU**

Obtaining an interactive CPU node with 1 node, 128 cores, and 249GB:

```
[user@login01 ~]$ srun --partition=debug --account=abc123 --pty --nodes=1 --ntasks-per-node=128
--mem=248 -t 00:30:00 --wait=0 --export=ALL /bin/bash
srun: job 5825986 queued and waiting for resources
srun: job 5825986 has been allocated resources
[user@exp-9-55 ~]$ hostname
exp-9-55
[user @exp-9-55 ~]$ date
Sun Jun 26 19:02:24 PDT 2022
[user@exp-9-55 ~]$ module list
Currently Loaded Modules:
1) slurm/expanse/20.02.3 2) cpu/0.15.4 3) intel/19.1.1.217 4) openmpi/4.0.4
[user@exp-9-55 ~]$ module purge
modul[user@exp-9-55 ~]$ module restore
Restoring modules from user's default
[user @exp-9-55 ~]$ module list
Currently Loaded Modules:
1) cpu/0.15.4 2) slurm/expanse/20.02.3 3) intel/19.1.1.217
[user @exp-9-55 ~]$ exit
```





# **Running Interactive Jobs - GPU**

The following example will request a GPU node, 10 cores, 1 GPU and 96G in the debug partition for 30 minutes. To ensure the GPU environment is properly loaded, please be sure run both the module purge and module restore commands.

```
[user@login02 ~]$ srun --partition=gpu-debug --pty --account=abc123 --ntasks-per-node=10
--nodes=1 --mem=96G --gpus=1 -t 00:30:00 --wait=0 --export=ALL /bin/bash
srun: job 5826039 queued and waiting for resources
srun: job 5826039 has been allocated resources
[user@exp-7-59 ~]$
[user@exp-7-59 ~1$ hostname
exp-7-59
                                    [user@exp-7-59 ~]$ nvidia-smi
                                    Thu Sep 16 00:09:41 2021
[user@exp-7-59 ~]$ module purge
[user@exp-7-59 ~]$ module restore
                                    | NVIDIA-SMI 460.32.03 | Driver Version: 460.32.03 | CUDA Version: 11.2
Restoring modules from user's default
                                    | GPU Name | Persistence-M| Bus-Id | Disp.A | Volatile Uncorr. ECC |
[user@exp-7-59 ~]$
                                    | Fan Temp Perf Pwr:Usage/Cap| | Memory-Usage | GPU-Util Compute M. | | | MIG M. |
                                       0 Tesla V100-SXM2... On | 00000000:18:00.0 Off |
                                                                                                             0 1
                                    | N/A 32C P0 41W / 300W | 0MiB / 32510MiB |
                                                                                                       Default |
                                    | Processes:
                                      GPU GI CI
                                                           PID Type Process name
                                                                                                    GPU Memory
                                            ID ID
                                                                                                    Usage
```





# General Steps: Compiling/Running Jobs

 Change to your working directory (e.g. hpctr-examples):

 Verify that the correct modules are loaded:

```
[mthomas@login02 ~]$ module list
Currently Loaded Modules:
1) shared 2) cpu/0.15.4 3) slurm/expanse/21.08.8
4) sdsc/1.0 5) DefaultModules
```

Compile MPI hello world code:

```
[mthomas@login02 mpi]$
[mthomas@login02 mpi]$ mpif90 -o hello_mpi hello_mpi.f90
[mthomas@login02 mpi]$
```

Verify executable create date:

```
[mthomas@login02 mpi]$
[mthomas@login02 mpi]$ ls -lt hello_mpi
-rwxr-xr-x 1 mthomas use300 22440 Jun 26 18:45 hello_mpi
[mthomas@login02 mpi]$
```

Submit job

```
[mthomas@login02 mpi]$
[mthomas@login02 mpi]$ sbatch hellompi-slurm.sb
Submitted batch job 14124495
[mthomas@login02 mpi]$
```





#### **Outline**

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### **Additional Examples**

- Copy code examples from:
  - /cm/shared/examples/sdsc/
  - These are the most recent verions
- Clone examples from
  - https://github.com/sdsc-hpc-training-org/expanse-101
- CPU Jobs & GPU Jobs





#### **Outline**

- Expanse Overview & Innovative Features
- Compiling and Linking Code
- Running Jobs
- Additional Examples
  - MPI Jobs
  - OpenMP Jobs
  - Hybrid MPI-OpenMP Jobs
- Final Comments





#### **MPI Hello World**

Change to the MPI examples directory:

```
[user@login02 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
[user@login02 MPI]$
```





# **MPI Hello World: Compile**

#### Set the environment and then compile the code

[user@login02 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-pernode=128 --mem=248 -t 00:30:00 --wait=0 --export=ALL /bin/bash

#### [user@login02 MPI]\$ module list

Currently Loaded Modules:

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

[user@login02 MPI]\$ module purge

[user@login02 MPI]\$ module load slurm

[user@login02 MPI]\$ module load cpu

[user@login02 MPI]\$ module load gcc/10.2.0

[user@login02 MPI]\$ module load openmpi/4.0.4

[user@login02 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

[user@login02 MPI]\$ mpif90 -o hello\_mpi hello\_mpi.f90 [user@login02 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.

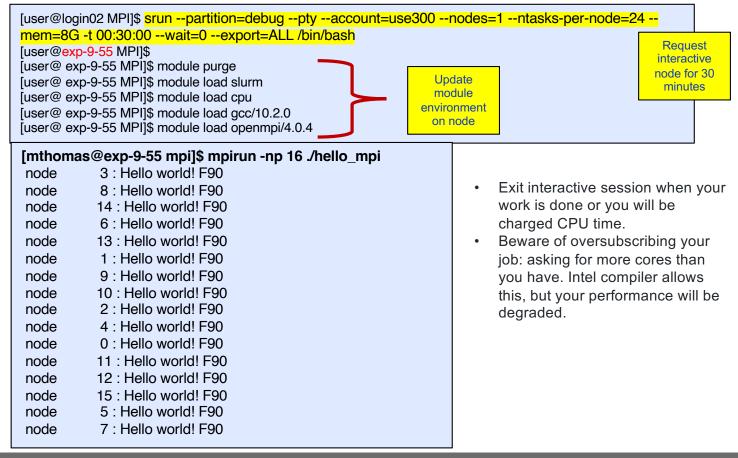
```
[user@login02 MPI]$ cat hellompi-slurm-gnu.sb
#!/bin/bash
#SBATCH --job-name="hellompi-gnu"
#SBATCH --output="hellompi-gnu.%j.%N.out"
#SBATCH --partition=compute
#SBATCH --nodes=2
#SBATCH --ntasks-per-node=128
#SBATCH --export=ALL
#SBATCH -A abc123
#SBATCH -t 00:10:00
                   squeue -u mthomas
#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
[user@login02 MPI]$
```

```
[user@login02 MPI]$ sbatch hellompi-slurm-gnu.sb; squeue -u user
Submitted batch job 108910
      JOBID PARTITION NAME USER ST TIME NODES NODELIST(REASON)
      108910 compute hellompi user PD 0:00 2 (None)
[user@login02 MPI]$ cat hellompi-gnu.108910.exp-12-54.out
          4: Hello world!
node
          5: Hello world!
          7: Hello world!
node
node
          0: Hello world!
          2: Hello world!
node
node
          3: Hello world!
node
          9: Hello world!
node
          10: Hello world!
[SNIP]
         247: Hello world!
node
         248 : Hello world!
node
         249: Hello world!
node
         186: Hello world!
node
node
         220: Hello world!
node
         203: Hello world!
         135: Hello world!
node
```





## **Using An Interactive mode**







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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
!SOMP 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

- 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.

```
[expanse-ln2:~/expanse101/OPENMP] cat openmp-slurm-
shared.sb
#!/bin/bash
#SBATCH --iob-name="hell openmp shared"
#SBATCH --output="hello_openmp_shared.%j.%N.out"
#SBATCH --partition=shared
#SBATCH --share
#SBATCH --nodes=1
#SBATCH --ntasks-per-node=16
#SBATCH --mem=80G
#SBATCH --export=ALL
#SBATCH -t 01:30:00
#SET the number of openmp threads
export OMP_NUM_THREADS=16
#Run the openmp job
./hello_openmp
```



module load cpu

module load aocc

#Run the openmp job

./hello openmp

#SET the number of openmp threads

export OMP NUM THREADS=16

[mthomas@login01 OPENMP]\$



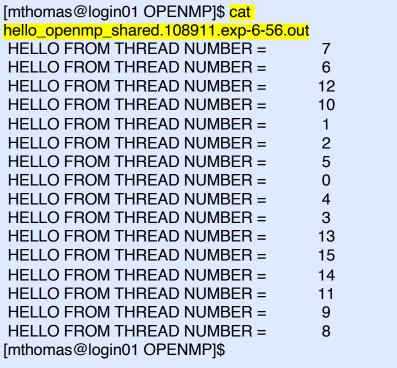
## OpenMP Hello World: submit job & monitor

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

```
[mthomas@login01 OPENMP]$ sbatch openmp-slurm-shared.sb
Submitted batch job 108911
[mthomas@login01 OPENMP]$ squeue -u mthomas
                                        TIME NODES NODELIST(REASO
JOBID PARTITION NAME USER ST
108911 shared hell ope mthomas PD
                                              1 (None)
                                       0:00
[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

-rw-r--r-- 1 mthomas use300 168 Oct 7 11:28 README.txt







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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
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-In2:~/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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## 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.





#### **Useful Links**

- Expanse User Guide:
  - https://www.sdsc.edu/support/user\_guides/expanse.html
- Expanse Overview:
  - https://education.sdsc.edu/training/interactive/202201 parallel computing concepts
- 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>
  - A 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 @ <a href="https://portal.xsede.org/allocations/startup">https://portal.xsede.org/allocations/startup</a>.
- Hands-on/code examples for this tutorial:
  - https://github.com/sdsc-hpc-training-org/hpctr-examples
  - https://github.com/sdsc-hpc-training-org/expanse-101
  - https://hpc-training.sdsc.edu/expanse-101/Expanse\_Tutorial.html





#### Resources

- Expanse User Guide & Tutorial
  - https://www.sdsc.edu/support/user\_guides/expanse.html
  - https://hpc-training.sdsc.edu/expanse-101/
- SDSC Training Resources
  - https://www.sdsc.edu/education and training/training hpc.html
- XSEDE Training Resources
  - https://www.xsede.org/for-users/training
  - https://cvw.cac.cornell.edu/expanse/





# Thank You



