Practical sessions HPC

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2 Job Submission and Control using SLURM





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- Download the file Private.pem in your files
- ② chmod 600 Private.pem in order to change the permissions
- Issh -i Private.pem ubuntu@185.127.66.38 to access the cluster

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Oreate a directory for your work in the students directory

• sinfo -o "%P"

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- sinfo -o "%P"
 - This command fetches information about the partitions (also known as queues) available on the cluster.
 - -o "%P" specifies the output format to only display the names of the partitions.

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● sinfo -o "%N"

- sinfo -o "%P"
 - This command fetches information about the partitions (also known as queues) available on the cluster.
 - -o "%P" specifies the output format to only display the names of the partitions.
- sinfo -o "%N"
 - This command retrieves information about the number of nodes available in each partition.
 - \bullet -o " % N " specifies the output format to display only the number of nodes.

Job Submission and Control using SLURM Task 1

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• sinfo -o "%P %T"

- sinfo -o "%P %T"
 - This command provides information about the state of each partition.
 - -o "%P %T" sets the output format to display both the partition name and its state.

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- sinfo -o "%P %T"
 - This command provides information about the state of each partition.
 - -o "%P %T" sets the output format to display both the partition name and its state.

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• sinfo -o "%c"

- sinfo -o "%P %T"
 - This command provides information about the state of each partition.
 - -o "%P %T" sets the output format to display both the partition name and its state.
- sinfo -o "%c"
 - This command retrieves information about the number of CPUs in each partition.
 - -o "%c" specifies the output format to display the total number of CPUs in each partition.

• sinfo -o "%m"

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- sinfo -o "%m"
 - This command fetches information about the available memory resources in each partition.
 - -o "%m" sets the output format to display only memory-related information.
 - -MEMORY indicates that we're specifically querying memory resources.

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sinfo --Node

- sinfo -o "%m"
 - This command fetches information about the available memory resources in each partition.
 - -o "%m" sets the output format to display only memory-related information.
 - -MEMORY indicates that we're specifically querying memory resources.
- sinfo --Node
 - This command fetches detailed information about the nodes in the cluster.
 - -Node is an option specifying that we're interested in node-related information.
 - -long requests a long format output, providing detailed information about the nodes.

• sinfo -N -h -O NODELIST

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- sinfo -N -h -O NODELIST
 - This command retrieves a list of nodes in the cluster.
 - -N specifies that we're only interested in node-related information.
 - -h omits the header from the output.
 - O NODELIST specifies the output format to display only the list of nodes.

Job Submission and Control using SLURM Task 1

scontrol show nodes :

- used to display detailed information about the nodes in the Slurm cluster
- provides information such as the node name, state, CPUs, sockets, cores per socket, threads per core, memory, node features, and other relevant attributes.

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scontrol show job followed by the job ID :

- used to display detailed information about a specific job in the Slurm workload manager
- provides information about the job's state, resources requested, job ID, job name, submission time, queue name, user ID, etc...

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srun --pty --nodes=1 --cpus-per-task=4 --time=1:00:00 bash
-i

 \Rightarrow Slurm will schedule the job, find an available node that matches the resource request

- Resource Allocation: It requests one node with 4 CPU cores.
- Time Limit: It sets a maximum runtime of 1 hour for the session.
- Interactive Shell: It opens an *interactive bash shell* on the allocated node.

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```
srun -n1 sleep 1
```

 \Rightarrow Slurm will allocate resources for $one\ task,$ which is to simply pause for $1\ second$

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 squeue : view information about jobs located in the Slurm scheduling queue

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- sacct : displays accounting data for all jobs and job steps in the Slurm job accounting log or Slurm database
- scontrol : view or modify Slurm configuration and state.

OpenMP

• an application programming interface (API) that supports multi-platform shared-memory multiprocessing programming



Figure 1: Shared memory programming \mathbb{P} , we have \mathbb{P} and \mathbb{P} and \mathbb{P}

OpenMP Task 1 : bubble sort



Figure 2: Bubble sort algorithm

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OpenMP

Task 1 : bubble sort odd-even



Figure 3: Bubble sort odd-even algorithm , and the set of \mathcal{A}

OpenMP Useful commands

#include <omp.h>

int main(){

```
// Start measuring time
double start_time = omp_get_wtime();
```

// To get the number of threads used int num_threads = omp_get_max_threads();

// End measuring time
double end time = omp get wtime();

// Print the time taken
printf("Time taken: %f seconds, Number of threads : %d \n", end time - start time, num threads);

Figure 4: Some useful commands

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- Choose how many threads you want with the command in the shell : export OMP_NUM_THREADS=n
- Ocompile your file with gcc -fopenmp -o exec_name file_name.c

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Execute it with ./exec_name

OR with a bash script (better because you won't have to use slurm command to allocate the resources you need)

```
#!/bin/bash -l
#SBATCH --job-name=openMP-job
#SBATCH --time=2:0:0
#SBATCH --nodes=1
#SBATCH --ntasks-per-node=4
# File name (without the .c extension)
filename="prime_para"
# Compile and execution if the compilation worked
gcc -fopenmp -o $filename $filename.c && ./$filename
```

Figure 5: Bash script for OpenMP

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MPI

Message Passing Interface (MPI) is a standardized and portable message-passing standard

- \Rightarrow Contrary to OpenMP, the memory is not shared between the threads
- \Rightarrow the threads communicate with each others by sending <code>messages</code>



Figure 6: Distributed memory programming

MPI Load



3 threads : 20 % 3 = 6,666666666... → :(

20 = 3 * **6** + **2**



```
void load(int me, int size, int Np, int *iBeg, int *iEnd)
    int r = size % Np;
    <u>if (me < r) {</u>
        *iBeg = me * (size / Np + 1);
        *iEnd = *iBeg + (size / Np + 1) - 1;
    } else {
        *iBeg = r + me * (size / Np);
        *iEnd = *iBeg + (size / Np) - 1;
```

Figure 8: Load code

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- Compile your file with mpicc -o filename filename.c
- Secure it with mpirun -np <number of process> ./filename

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OR with a bash script (better because you won't have to use slurm command to allocate the resources you need)

```
#!/bin/bash -l
#SBATCH --job-name=mpi-job
#SBATCH --time=2:0:0
#SBATCH --nodes=1
#SBATCH --nodes=1
#SBATCH --ntasks-per-node=4
# File name (without the .c extension)
filename="prime_para"
# Compile and execution if the compilation worked
mpicc -o $filename $filename.c && mpirun -np 4 ./$filename
```

Figure 9: Bash script for MPI

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MPI Useful commands

```
#include <mpi.h>
int main(int argc, char *argv[]) {
    int me, Np;
   MPI Init(&argc, &argv);
   MPI Comm rank(MPI COMM WORLD, &me);
    MPI Comm size(MPI COMM WORLD, &Np);
    double start time = MPI Wtime();
    int iBeg, iEnd;
    load(me, N, Np, &iBeg, &iEnd);
    double end time = MPI Wtime();
    if (me == 0) {
        printf("Time taken : %f seconds\n", end time - start time);
    MPI Finalize();
    return 0;
```

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