Parallel Computing Exercise session 5

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Outline

HPC@KUL

Working on the Cluster

Example: Hello, World!

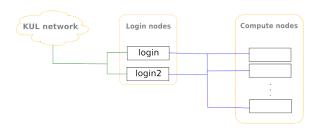
VSC

- VSC is a collaboration between U Antwerpen, VUB, U Gent U Hasselt and KU Leuven
- Rotating investment in new hardware at different institutes
- All institutes can access all hardware
- Two kinds of cluster:
 - ▶ Tier-1: Latest, fastest hardware; access only after project is approved
 - ► Tier-2: Not as new (but still a lot of compute power); Anyone can access if they have/purchase credits
- ▶ When a new Tier-1 system is built, the old one becomes a Tier-2 system
- A full overview can be found here.

ThinKing

- Older tier-2 system at KU Leuven/U Hasselt
- Newer system is called Genius
- ► ThinKing consists of:
 - ► 160 ivybridge nodes
 - ▶ 2 Xeon E5-2680 v2 CPUs@2.8 GHz, 10 cores each
 - ▶ 64 GB RAM (130 nodes) / 128 GB RAM (30 nodes)
 - ▶ 144 haswell nodes
 - 2 Xeon E5-2680 v3 CPUs@2.5 GHz, 12 cores each
 - ▶ 64 GB RAM (32 nodes) / 128 GB RAM (96 nodes)
 - 5 GPGPU nodes
 - 2 Xeon E5-2650 v3 CPUs@2.3 GHz, 10 cores each
 - ► 64 GB RAM
 - 2 NVIDIA K40c@750 MHz, 12 GB GDDR

Getting access



- VSC clusters consist of login nodes and compute nodes
- You can log in to both using ssh, but in general you will work on a login node
- Access to login node: ssh vsc01234@login-thinking.hpc.kuleuven.be
- Windows devices can use a third party tool such as Putty

Overhead on the cluster

- To run a job on the cluster, you need to be allocated a node
 ⇒ Queuing system takes care of fair distribution of resources
- The queuing system needs to be able to schedule jobs
 ⇒ Provide a .pbs file with resource estimates for your job
- The cluster contains many conflicting software packages
 ⇒ Load the ones you want with a module system
- ▶ Jobs are scheduled and run in a batch fashion
 ⇒ Output is available in files once the job completes

Useful commands

- Copying files to cluster: scp local_file vsc01234@login-thinking.hpc.kuleuven.be:remote_folder
- Copying directory from cluster: scp -r vsc01234@login-th inking.hpc.kuleuven.be:remote_folder local_folder
- ▶ Loading modules: module load {intel, GCC, ...}
- Submitting a job: qsub -A lp_edu_alg_parallel_comp_2021 job_name.pbs
- Checking your jobs: qstat
- Estimate a job's start time: showstart jobID
- Check directory contents: 1s
- Copy file/directory: cp
- Change directory: cd
- Find out more about a command: man command
- ► Edit file: nano file_name
- ▶ Other useful commands: pwd, mv, mkdir, touch, rm,...



An example PBS file

► This file defines a job on 4 nodes for 1 minute, requests 10 processors per node and loads the most recent Intel toolkit:

```
#!/bin/bash -I
#PBS -I nodes=4:ppn=10
#PBS -I walltime=00:01:00
module load intel

cd $PBS_O_WORKDIR
./myprog
```

OpenMP

- Download omp_hello.c and hello_omp.pbs
- 2. Copy these files to the cluster using scp
- Compile the program using g++ -o hello omp_hello.c -fopenmp
- 4. Check that you understand the file hello_omp.pbs
- 5. Submit the job using qsub -A lp_edu_alg_parallel_comp_2021 hello_omp.pbs
- 6. Check your results

MPI

- Download mpi_hello.c and hello_mpi.pbs
- 2. Copy these files to the cluster using scp
- Load the Intel MPI toolkit using module load intel
- Compile the program using mpiicpc -o hello mpi_hello.c
- 5. Check that you understand the file hello_mpi.pbs
- 6. Submit the job using qsub -A lp_edu_alg_parallel_comp_2021 hello_mpi.pbs
- 7. Check your results