MPI on Cirrus and ARCHER2





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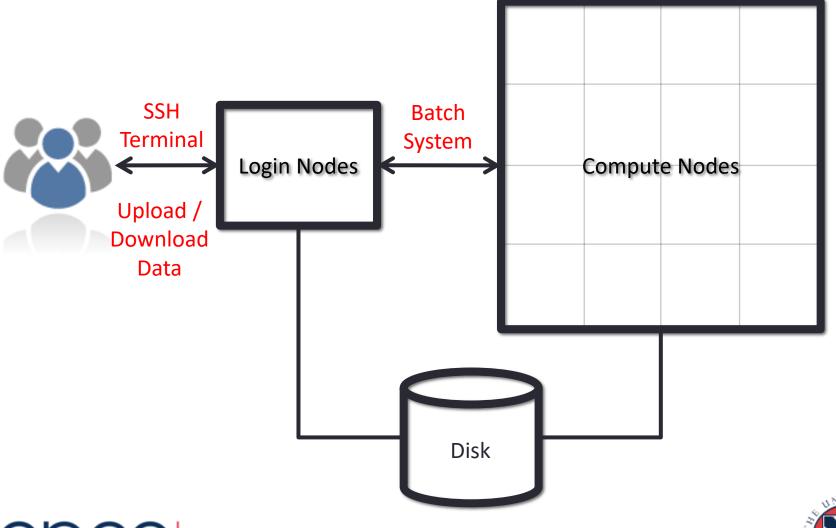


Access

- ARCHER2: ssh -x user@login.archer2.ac.uk
- Cirrus: ssh -X user@login.cirrus.ac.uk
- You can access EPCC systems using ssh from anywhere
 - Linux: terminal + command line
 - Mac: Mac terminal + command line + enable an X server (e.g. xquartz) to display any graphics
 - Windows: need to install an ssh program such as MobaXterm
- This gives you access to a login node
 - a few login nodes, users assigned to different nodes depending on system load
 - login nodes shared by all users
 - the many hundreds of compute nodes are accessed exclusively via the SLURM batch system
 - on ARCHER2, MPI programs can only be run on the compute nodes
- Full instructions on access & login in separate document on course web pages



Typical HPC system layout



Useful files and templates

- Take a copy of MPP-templates.tar
 - see the course web pages
- unpack: tar -xvf MPP-templates.tar
- Crib sheets for MPI programs available on course web pages





Compiling MPI Programs on ARCHER2

- C programmers use cc
- C++ programmers use CC
- Fortran programmers use ftn
- There is nothing magic about these MPI "compilers"!
 - simply wrappers which automatically include various libraries etc
 - compilation done by standard (Cray) compilers
 - clang, clang++ and crayftn
- You can use the supplied Makefiles (C, C++, Fortran) for convenience

```
make -f Makefile_c
make -f Makefile_cc
make -f Makefile f90
```

- Easiest to make a copy of your choice called "Makefile"
 - e.g. cp Makefile_c Makefile
 - also need to change the first line "MF=" in the Makefile itself
 - then you can just type "make"





ARCHER2 idiosyncrasies

- Not possible to run MPI programs directly on login nodes
- Can be a substantial delay in batch queues
 - have dedicated queues for some courses for more rapid turnaround
 - or use the short queue for development
- Cannot run from the home file system
 - on logging in, your working directory is /home/taXXX/taXXX/username/
 - compute nodes can only see the /work/ file system, not /home/
- Recommendation
 - do everything in /work/
 - i.e. change directory to /work/taXXX/taXXX/username/





Running on ARCHER2 back-end

- Run via a batch system
 - on ARCHER2 we use SLURM
 - submit a script that then launches your program
- In MPP-templates/ is a standard batch script: archer2mpi.job
 - set up to run a program called hello in the short queue
- Submit: sbatch archer2mpi.job
 - runs on 4 processes of a single ARCHER2 node (each node has 128 CPU-cores)
 - output will appear in a file called hello-xxxxxx.out
 - can follow job progress using command: squeue -u \$USER
 - full instructions included as comments in the template
- Batch script has instructions on how to use different queues





Running interactively

- Cannot run on ARCHER2 login node
 - must run on compute nodes via SLURM and sbatch
- However, can run "interactively"
 - sbatch command waits until resources are available
 - output appears on the screen in real time as your program runs
 - can be very useful for debugging (e.g. to detect deadlock)

```
srun --partition=standard --qos=short --reservation=shortqos \
--time=00:01:00 --unbuffered --cpu-bind=cores --nodes=1 --tasks=4 ./hello
```





C++ Interface

- MPI is not an OO interface
 - however, can be called from C++
- Originally had different function calls, e.g.

```
- MPI::Intracomm comm;
- ...
- MPI::Init();
- comm = MPI::COMM_WORLD;
- rank = comm.Get_rank();
- size = comm.Get_size();
```

- Compiler is called CC
 - see hello.cc and Makefile_cc

C++ interface is now removed

Must therefore cross-call to C





Documentation

- MPI Standard available online
 - See: http://www.mpi-forum.org/docs/
 - currently version 4.0



http://www.hlrs.de/mpi/mpi31/

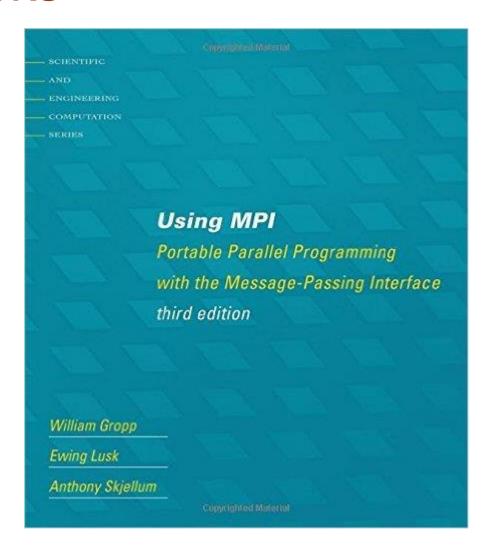


- Man pages available on Cirrus and ARCHER2
 - must use the C style of naming: man MPI_Routine_name, e.g.:
 - user@computer\$ man MPI_Init





MPI Books







Exercise: Hello World

The minimal MPI program

- See Exercise 1 on the exercise sheet
- Write an MPI program that prints a message to the screen
- Main purpose is to get you compiling and running parallel programs on HPC system
 - both interactively and in batch via SLURM and sbatch
 - also illustrates the SPMD model and use of basic MPI calls
- We supply some very basic template code
 - you need to add appropriate calls to compute rank and size



