MPI on Cirrus and ARCHER













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Access

- ARCHER: ssh -xx user@login.archer.ac.uk
- Cirrus: ssh -XY user@cirrus-msc.epcc.ed.ac.uk
 - you must use this dedicated MSc login node
- You can access systems using ssh from anywhere
 - Trivial for Linux
 - Mac: enable the X server (xquartz) to display any graphics
 - Windows: need to install an X server program, e.g. MobaXterm





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





Setting up Cirrus environment

- Load the Message-Passing Toolkit
 - module load mpt
- Load the Intel Compilers
 - module load intel-compilers-17
- To automate, add these lines to your ".bash_profile" file

```
[user@cirrus] gedit ~/.bash_profile
```





Compiling MPI Programs on Cirrus

• C programmers use: mpicc -cc=icc

• C++ programmers use: mpicxx -cxx=icpc

Fortran programmers use: mpif90

- There is nothing magic about these MPI compilers!
 - simply wrappers which automatically include various libraries etc
 - compilation done by standard (e.g. Intel) compilers
 - · icc, icpc and ifort
- You can use the supplied Makefiles for convenience

```
- make -f Makefile_c
```

- make -f Makefile cc
- make -f Makefile_f90
- Easiest to make a copy of one of these called "Makefile"
 - also need to change the line "MF=" in the Makefile itself





Running interactively on Cirrus

- Timings will not be reliable
 - shared with other users, many more processes than processors
 - but **very useful** during development and for debugging
- mpirun -n 4 ./mpiprog.exe
 - runs your code on 4 processes
- NOTE
 - output might be buffered
 - if your program crashes, you may see no output at all
- May need to explicitly flush prints to screen
 - FLUSH(6)
 - fflush(stdout);





Running batch jobs on Cirrus

- Run via a batch system
 - Cirrus uses Portable Batch System (PBS); submit script that launches your program
- In MPP-templates/ is a standard batch script: cirrusmpi.pbs
 - make a copy of this file with a name that matches your executable, e.g.
 - user@cirrus\$ cp cirrusmpi.pbs hello.pbs
- To run on 4 processors: qsub hello.pbs
 - use reserved queue during lab sessions, e.g. qsub –q R12345 hello.pbs
 - automatically runs executable called "hello"
 - output will appear in a file called hello.pbs.oxxxxx
 - can follow job progress using qstat or qstat -u \$USER
 - script also times your program using the Unix "time" command
 - full instructions included as comments in the template
 - no need to alter the script just rename it as appropriate
 - e.g. to run a program "pingpong" make another copy called "pingpong.pbs"
- MSc students should alter charging: d167 -> d167-UUN
 - e.g. #PBS -A d167-s1234567



Cirrus idiosyncrasies

 By default, MPI wrappers are not in your path user@cirrus\$ mpicc
 -bash: mpicc: command not found

- To access correct version: module load mpt
 -defaults to GNU compilers: gcc, g++ and gfortran
 -in batch system, job launcher is called mpiexec_mpt
- Intel compilers: module load intel-compilers-17
 -add these to end of your .bash_profile file in home directory
 -to check you have the right version (similarly for mpif90)

```
user@cirrus$ which mpicc
/opt/hpe/hpc/mpt/mpt-2.16/bin/mpicc
```

-mpif90 automatically picks up the Intel Fortran compiler
 -to use Intel C [C++] compilers: mpicc -cc=icc [-cc=icpc]





Compiling MPI Programs on ARCHER

- Fortran programmers use ftn
- C programmers use cc
- There is nothing magic about these MPI compilers!
 - simply wrappers which automatically include various libraries etc
 - compilation done by standard (Cray) compilers
 - crayftn and craycc
- 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"





ARCHER idiosyncrasies

- Not possible to run directly on front-end
- Can be a substantial delay in batch queues
 - we may have dedicated queues for the course for more rapid turnaround!
- Cannot run from the home file system
 - back-end nodes can only see the work file system
- Recommendation
 - do everything in /work/
 - i.e. change directory to /work/y14/y14/username/





Running on ARCHER back-end

- Run via a batch system
 - on ARCHER we use the Portable Batch System (PBS)
 - submit a script that then launches your program
- In MPP-templates/ is a standard batch script: archermpi.pbs
 - make a copy of this file with a name that matches your executable, e.g.
 - user@archer\$ cp archermpi.pbs hello.pbs
- Submit: qsub -q RXXXXXX hello.pbs
 - you will need to alter NPROCS (the argument to "aprun") by hand
 - ... and select more than one node for more than 24 processes
 - output will appear in a file called hello.pbs.oxxxxx
 - can follow job progress using qstat command
 - script also times your program using the Unix "time" command
 - full instructions included as comments in the template
- If there is no reserved queue
 - qsub -q short hello.pbs
 - short queue is for small jobs less than 20 minutes during working hours





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 mpicxx
 - 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 3.1



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

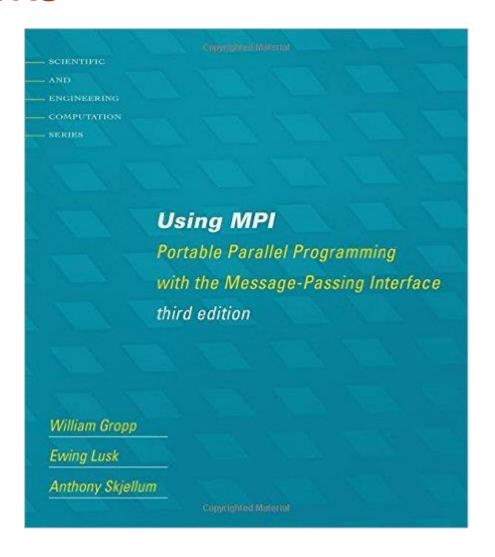


- Man pages available on Cirrus and ARCHER
 - must use the C style of naming: man MPI_Routine_name, e.g.:
 - user@computer\$ man MPI_Init





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 ness
 - 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



