## **Practical MPI**

## HelloWorld

Login to MareNostrum with your student account and find the source files in the professor account:

/home/nct00/nct00002/hellloWorld/

```
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#include <stdio.h>
#include <mpi.h>

int main (int argc, char argv[]) {

int rank, size;

MPI_Init(NULL, NULL);

MPI_Comm_size(MPI_COMM_WORLD, &size);

MPI_Comm_rank(MPI_COMM_WORLD, &rank);

char processor_name(MPI_MAX_PROCESSOR_NAME];
int name_len;

MPI_Get_processor_name(processor_name, &name_len);

printf("Hello world from %s, rank %d"
    " out of %d processors\n",
    processor_name, rank, size);

MPI_Finalize();

return(0);
}

int mans:

helloworld:bash
```

HelloWorld MPI Source code

Copy the files to your student accounts, compile the program and submit it to the execution queue.

- cp /home/nct00...
- make
- bsub < submit-mpi.sh

View the output files for messages from the batch system and the actual output from your program.

How many processes can you execute on one node and how can you influence this?

Run the application on a different number of processors and observe the name of the processor in the output file.

The MareNostrum Userguide can be downloaded at

http://www.bsc.es/support/MareNostrum3-ug.pdf