

# INTRODUCTION TO HPC IN COMPUTATIONAL MODELLING

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Short course on HPC-based Computational Bio-Medicine  
BSC – Barcelona – 12/02/2020



**SURF SARA**



# High Performance Computing at SURFsara

Amsterdam Science Park





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Amsterdam Science Park

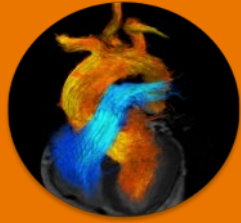




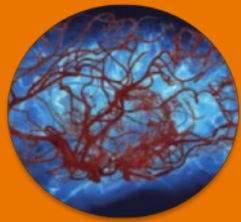
# High Performance Computing at SURFsara

- Maintain Cartesius and Lisa
- User support
- Training
- Development and Innovation
- Involvement in EU projects
- After July 2020: officially SURF (!!!)

# High Performance Computing in CompBioMed



**Cardiovascular  
medicine**



**Neuro  
Musculoskeletal  
medicine**



**Molecularly  
based  
medicine**



**CompBioMed**

A Centre of Excellence in Computational Biomedicine

- Support and facilitate modeling and simulation activities within biomedical community.
- Development and sustainability of software tools and services.
- Enhance industries in the healthcare sector (pharmaceuticals & medical device manufacture).



# SURFsara and HPC in Europe



- HPC resources access, and benchmarking
- Best practices in HPC and trainings

[www.prace-ri.eu](http://www.prace-ri.eu)



- EU Exascale machine based on EU chip (2023)
- Codesign HW/SW

[www.european-processor-initiative.eu](http://www.european-processor-initiative.eu)



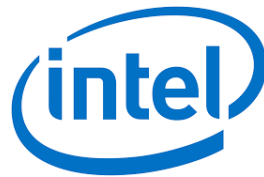
- Simplify access to IT services
- Improving service quality

[www.eosc-hub.eu](http://www.eosc-hub.eu)

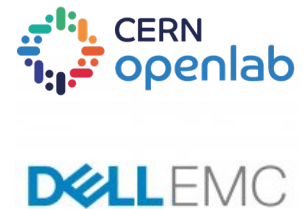
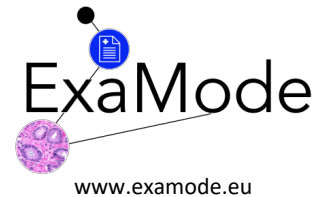
# SURFsara and HPC in Europe



- Convergence of AI and HPC



Intel Parallel Compute Centre



- Quantum computing





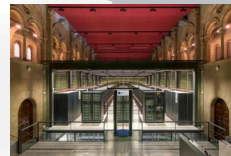
# High Performance Computing in CompBioMed



**Cartesius, Lisa**  
SURFsara (NL)

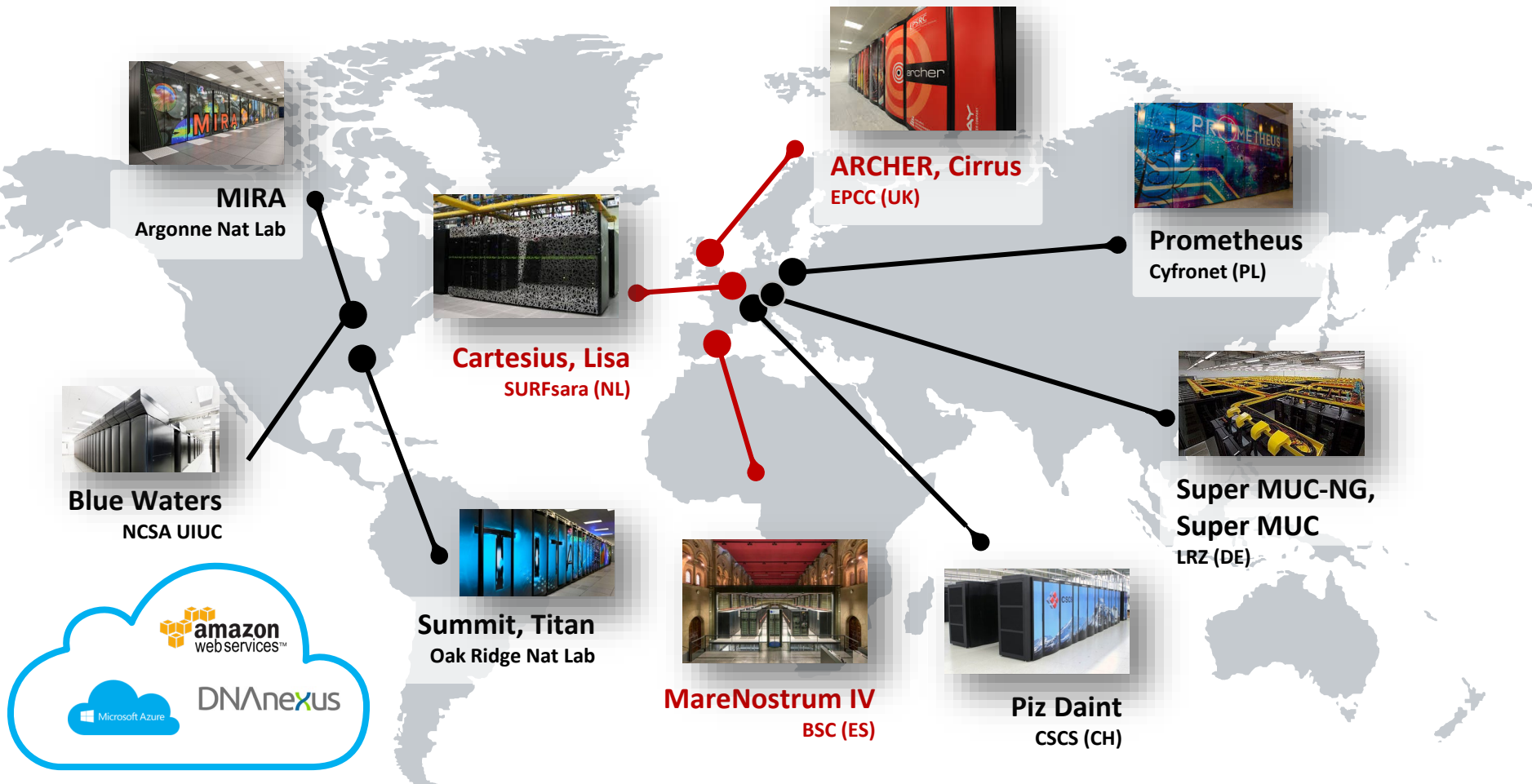


**ARCHER, Cirrus**  
EPCC (UK)



**MareNostrum IV**  
BSC (ES)

# High Performance Computing in CompBioMed







# Outline

## What is a Supercomputer?

- Working with an HPC system
- HPC access and usage

## More Linux tools

- Hands-on with cheat sheet
- Linux scripts

## Running jobs on a Supercomputer

- Use the batch system
- Running jobs

# What is a Supercomputer?

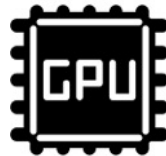
## User Experience

- Multiuser system
- Unix OS
- Optimized software



## Compute power

- Many CPUs system
- Specialized hardware
- Low-latency & high bandwidth connections



## Storage

- Efficient I/O
- Large Memories





# What is a Supercomputer?

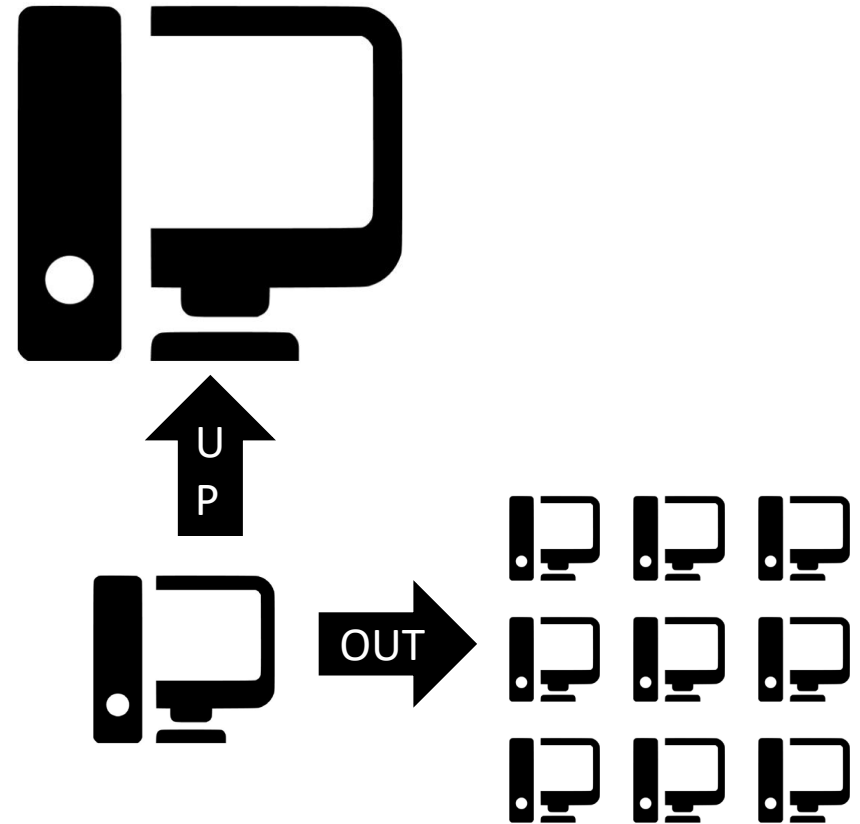
## Why you need a Supercomputer?

- Scale up

- Faster CPUs
- Large memories
- Specialized Hardware/Software

- Scale out

- Large parallel applications
- Many small- to medium- size jobs



# What is a Supercomputer?

Is NOT like this...





# What is a Supercomputer?

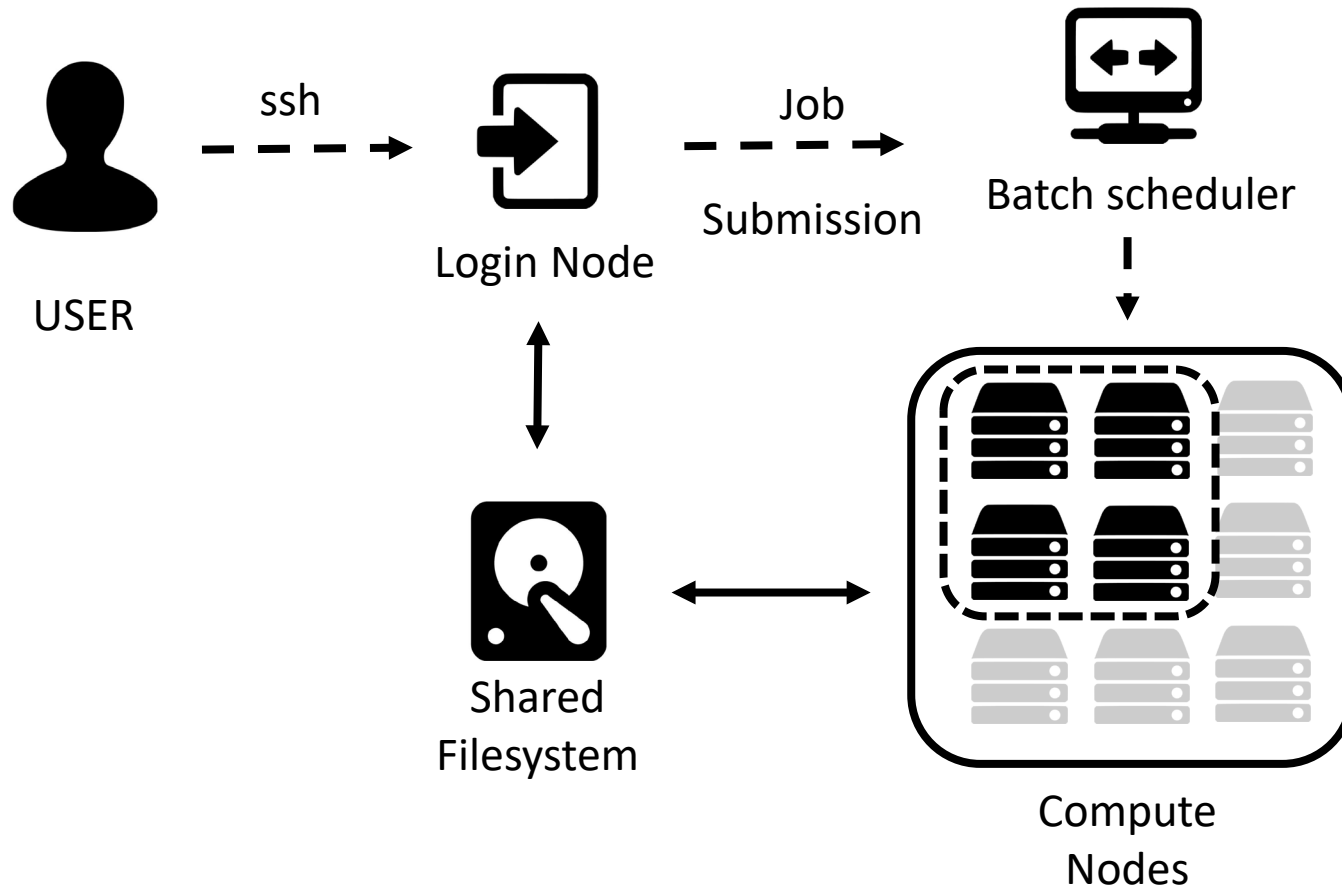
```

2. nct00013@login1:/home/nct00/nct00013 (ssh)
There were 2 failed login attempts since the last successful login.
Last login: Fri Jan 25 10:02:12 2019 from 84.88.53.31
+-----+
|                                             |
|               BSC                           |
|                                             |
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|
| - Please read the User's Guide:
|
|      https://www.bsc.es/user-support/mn4.php
|
| Please contact support@bsc.es for further questions
|
+-----+

/usr/bin/manpath: can't set the locale; make sure $LC_* and $LANG are correct
Set INTEL compilers as MPI wrappers backend
load impi/2017.4 (PATH, MANPATH, LD_LIBRARY_PATH)
load mkL/2017.4 (LD_LIBRARY_PATH)
nct00013@login1::~~>
nct00013@login1::~~>
nct00013@login1::~~>
nct00013@login1::~~>
nct00013@login1::~~>
```

# What is a Supercomputer?

## Working with a Supercomputer





# What is a Supercomputer?

## Working with a Supercomputer



### Login node(s)

- Editing and transferring files
- Compile programs
- Prepare simulations



### Compute nodes

- Multicore nodes
- Large memories
- High-speed interconnections



### Batch scheduler

- Resource allocation
- Job queueing
- Accounts and budget



### File system

- Parallel FS
- Efficient I/O
- Node local disks

# MORE LINUX TOOLS

Introduction to HPC in Computational  
Modelling



# Getting started with HPC systems



## What is UNIX?

- Operating System
  - Program that controls all other parts of a computer system
  - Allocates computer's resources and schedules tasks
  - Allows the user to use the facilities provided by the system
  - Essential to all computer systems
- Multi-User and Multi-Tasking
  - Multiple users have multiple tasks running simultaneously
- Two essential principles
  - Everything is a file
  - KISS



# Getting started with HPC systems

## Working with a Supercomputer

**1**

### Login and transfer files to remote machine

- ssh, scp/ftp
- Command line, GUI

**2**

### Prepare your jobs

- Input + Software
- Job submission script

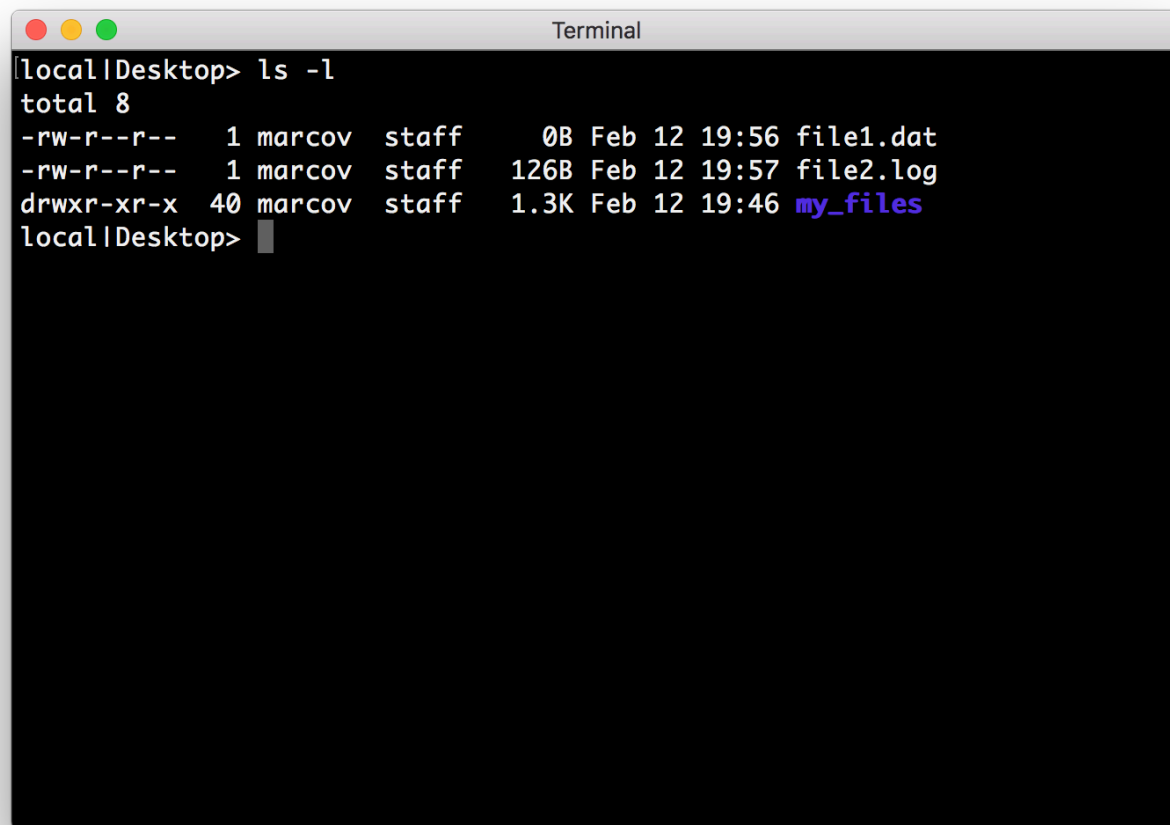
**3**

### Submit your job and retrieve output

- Submit job to the batch system
- Monitor job, retrieve output (remote visualization)

# Getting started with HPC systems

## Terminal – Working with a Unix system



```
Terminal
[local@Desktop> ls -l
total 8
-rw-r--r--  1 marcov  staff    0B Feb 12 19:56 file1.dat
-rw-r--r--  1 marcov  staff  126B Feb 12 19:57 file2.log
drwxr-xr-x 40 marcov  staff  1.3K Feb 12 19:46 my_files
local@Desktop>
```

# Getting started with HPC systems

## Install UNIX tools on your local machine

- **Windows**

- Putty
- MobaXterm (<http://mobaxterm.mobatek.net>)

- **Mac OSX**

- Terminal (pre-installed)
- XQuartz (<http://www.xquartz.org>)

- **Linux**

- You are already well equipped!



# Getting started with HPC systems

- Keep the Linux cheat sheet open:

<https://edu.nl/f4qaf>

- Check the material for future hands-on exercises:

<https://edu.nl/mtbvf>

- Can you download and transfer the zip file to MareNostrum4 and extract the zip in your home?

- Otherwise just copy the files to your home directly:

```
nct00008@login1:~> cp -r  
/gpfs/projects/nct00/nct00008/linux-hands-on ~/
```

# Getting started with HPC systems

## Basic scripting

- A Bash script is a plain text file which contains a series of commands.
- Any command you can run on the command line can be put into a script (v.v.)
- It will be executed like a normal program:  
`./script.sh`

```
nct00013@login1:~> cat unix/simple.sh
```

```
#!/bin/bash
```

```
echo "Hi, I'm your first script."  
echo
```

```
lscpu --help > cpu.log  
lscpu >> cpu.log
```

```
echo "I've left something for you."  
echo "Ciao"
```

# Getting started with HPC systems

## Basic scripting

- A script need execution permissions

```
chmod +x script.sh
```

- A script (command) produce output and error messages

- Stdin: your keyboard
- Stdout: your screen
- Stderr: your screen

- Out/Err can be redirected to files or commands

```
echo "3/0" | bc > calc.log
```

```
nct00013@login1:~> cat unix/simple.sh
```

```
#!/bin/bash
```

```
echo "Hi, I'm your first script."  
echo
```

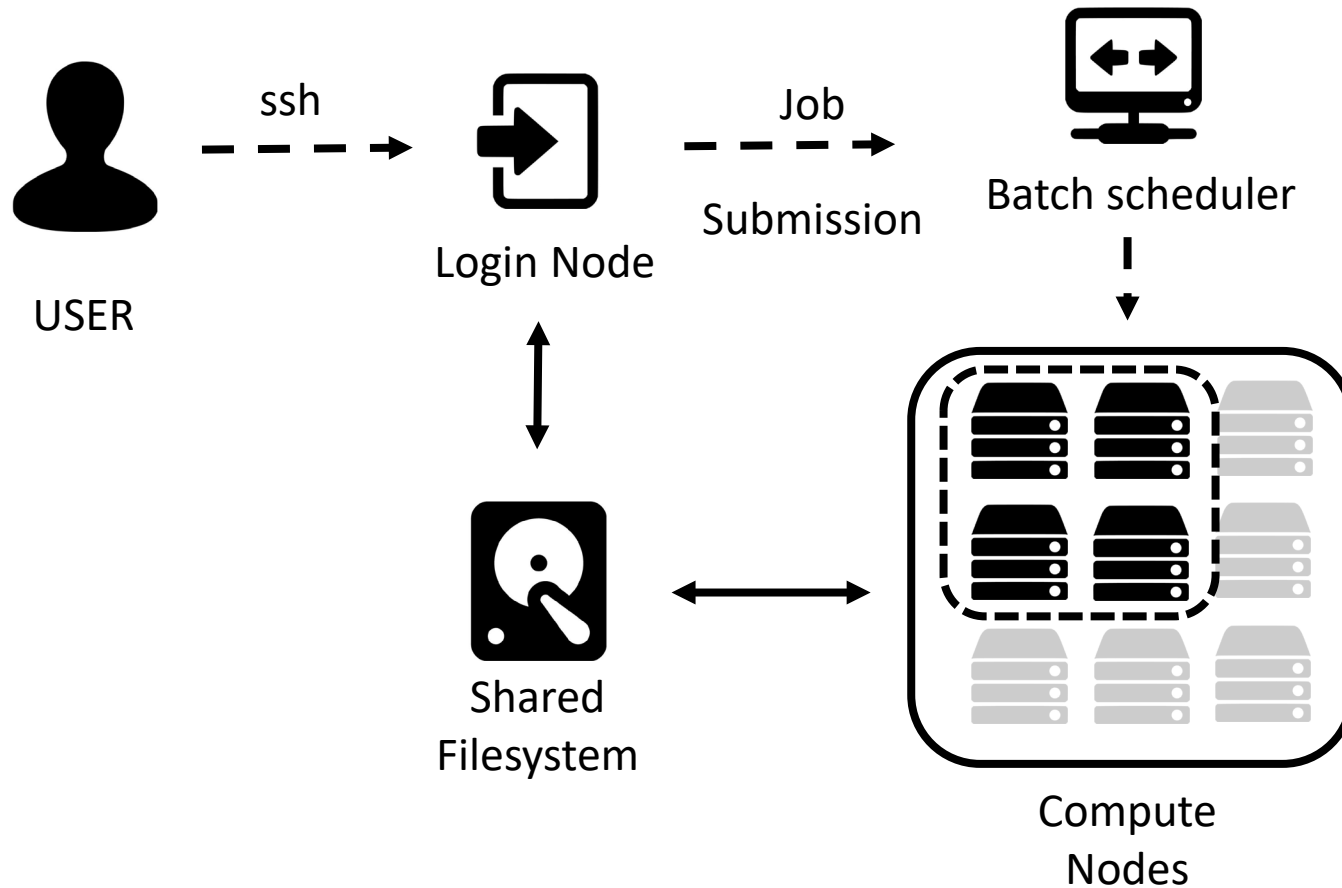
```
lscpu --help > cpu.log  
lscpu >> cpu.log
```

```
echo "I've left something for you."  
echo "Ciao"
```



# Getting started with HPC systems

## Submitting jobs in an HPC system



# Getting started with HPC systems

- Supercomputers use batch systems to distribute computational tasks over the available nodes.
- Instead of executing commands interactively, you prepare a job script
  - Script containing the commands to execute
  - Resource characteristics (specific)
- The batch system is responsible for allocating cores, processors or nodes to a job.



# Getting started with HPC systems

## Advantages of a batch system are:

- It allows to run MANY jobs at the same time
  - The system takes care that they are run efficiently on the available resources.
- Multiusers, queue system
  - A batch system allows users to always submit jobs, even if a lot of people are using the system at the same time. In addition take care of budgeting and fair resource usage.
- System load balance
  - The system takes care of balancing the load across nodes and during time. In a batch system, most jobs may be submitted during office hours, but the scheduler will continue to start jobs at night as nodes become available.





# RUNNING ON A SUPERCOMPUTER

Introduction to HPC in Computational  
Modelling



# Running on a Supercomputer

## Job submission script

### ■ SLURM Job directives

A job script must contain directives to inform the batch system about the characteristics of the job. These directives appear as comments (#SBATCH) in the job script and have to conform with the sbatch syntax.

- number of compute nodes
- number of processes to start
- total wall clock time of the job
- requesting a specific queue
- name of the file where std out is printed

<https://slurm.schedmd.com/sbatch.html>

[www.bsc.es/support/MareNostrum4-ug.pdf](http://www.bsc.es/support/MareNostrum4-ug.pdf)

```
nct00008@login1:~> head batch/slurm2.sub
```

```
#!/bin/bash
#SBATCH --job-name="test_multinode"
#SBATCH --nodes=2
#SBATCH --tasks-per-node=3
#SBATCH --time=00:02:00
#SBATCH --qos=training
#SBATCH --workdir=.
#SBATCH --output=multinode_%j.out
#SBATCH --error=multinode_%j.err
```

# Running on a Supercomputer

## Submitting and monitoring jobs

Jobs are submitted and controlled using SLURM commands

- submits a “job script” to the queue system:

```
nct00008@login1:~> sbatch <job script>
```

- shows all the submitted jobs and their status:

```
nct00008@login1:~> squeue
```

- remove the job from the queue system:

```
nct00008@login1:~> scancel <jobid>
```

- run a command on the cluster

```
nct00008@login1:~> srun <command>
```

# Running on a Supercomputer

## Queues

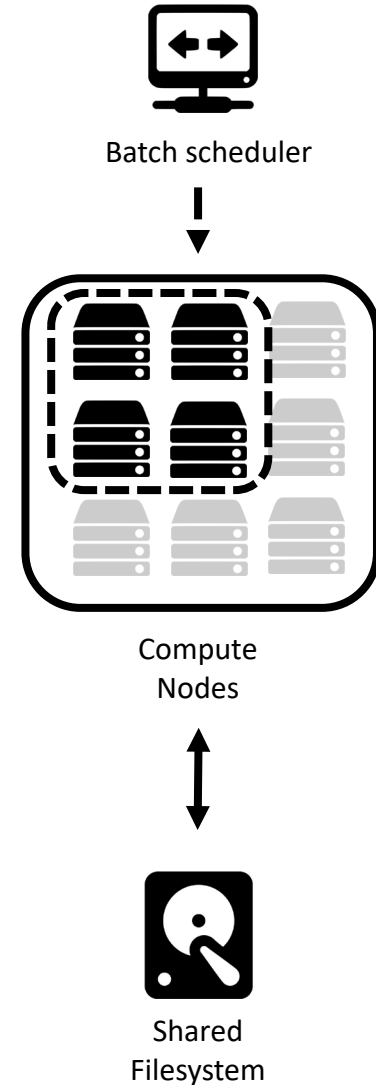
- There are several queues present in the machines and different users may access different queues.
- `bsc_queue` shows information of the available queues

## Filesystems

- Each user has several areas of disk space for storing files.
- These areas may have size or time limits (`bsc_quota`).
- Choose carefully where to store your data!

## Software stack

- HPC centers usually offer a set of software to their users
- `module` command show available software





# Running on a Supercomputer

## Module environment

- Provide lots of useful software packages
- In many different versions
- Maintained by experts
- Optimized for the architecture
- Available operations:
  - list all / specific software
  - load/unload a specific software
  - list loaded modules

```
nct00008@login1:~> module avail
```

```
nct00008@login1:~> module avail python
```

```
nct00008@login1:~> module load siesta (SUCCESS...?)
```

```
nct00008@login1:~> module load ...
```

```
nct00008@login1:~> module load python/2.7.14
```

```
nct00008@login1:~> module unload siesta
```

```
nct00008@login1:~> module list
```

# Running on a Supercomputer

```
nct00013@login1:~> module list
```

Currently Loaded Modules:

1) intel/2017.4 2) impi/2017.4 3) mkl/2017.4 4) bsc/1.0 5) lzo/2.10

```
nct00013@login1:~> gmx_mpi --version
```

-bash: gmx\_mpi: command not found

```
nct00013@login1:~> module load gromacs
```

load gromacs/2016.4 (PATH, LD\_LIBRARY\_PATH)

```
nct00013@login1:~> which gmx_mpi
```

/apps/GROMACS/2016.4/INTEL/IMPI/bin/gmx\_mpi

```
nct00013@login1:~> gmx_mpi --version
```

:-) GROMACS - gmx\_mpi, 2016.4-dev-20170515-2eb5a6307-unknown (-:

```
nct00013@login1:~> which gmx_mpi
```

/apps/GROMACS/2016.4/INTEL/IMPI/bin/gmx\_mpi

# Running on a Supercomputer

- Play with the material!

<https://edu.nl/mtbvf>

- ... and keep the cheat sheet around for extra help

<https://edu.nl/f4qaf>

- unix
  - bash scripts and basic Unix commands
- batch
  - simple submissions scripts to start using SLURM
- parallel
  - advanced submissions scripts to run parallel applications

# Getting started with HPC

## Working with a Supercomputer

### Login and transfer files to remote machine



- ssh, scp/ftp
- Command line, GUI

### Prepare your jobs



- Input + Software
- Job submission script

### Submit your job and retrieve output

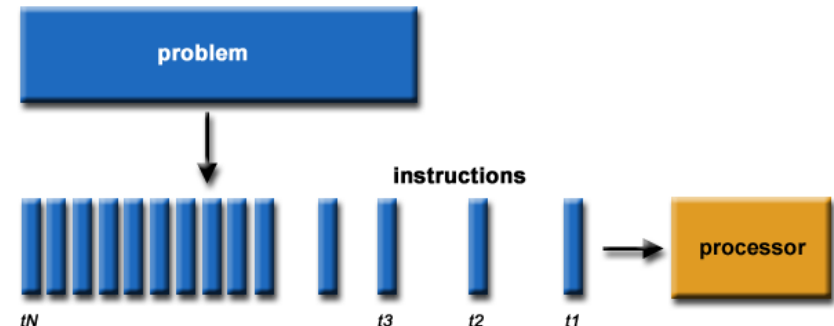


- Submit job to the batch system
- Monitor job, retrieve output (remote visualization)

# Running on a Supercomputer

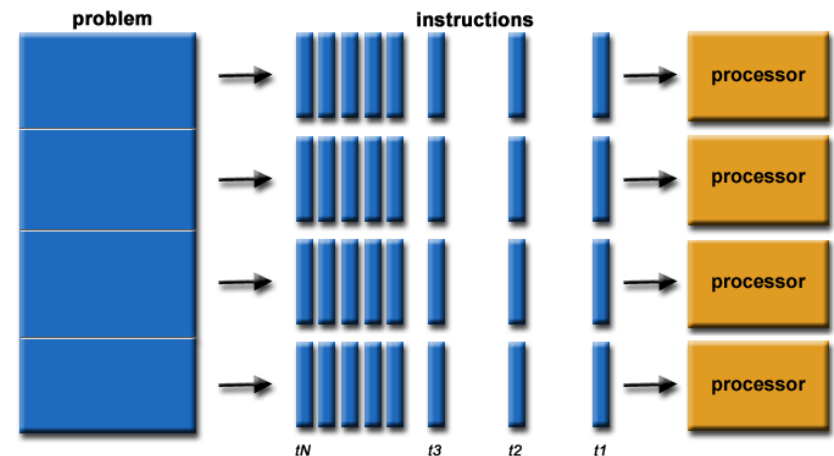
## Serial computing

- A problem is broken into a discrete series of instructions, which are executed sequentially on a single processor.



## Parallel computing

- A problem is broken into discrete parts that can be solved concurrently using simultaneously multiple resources.



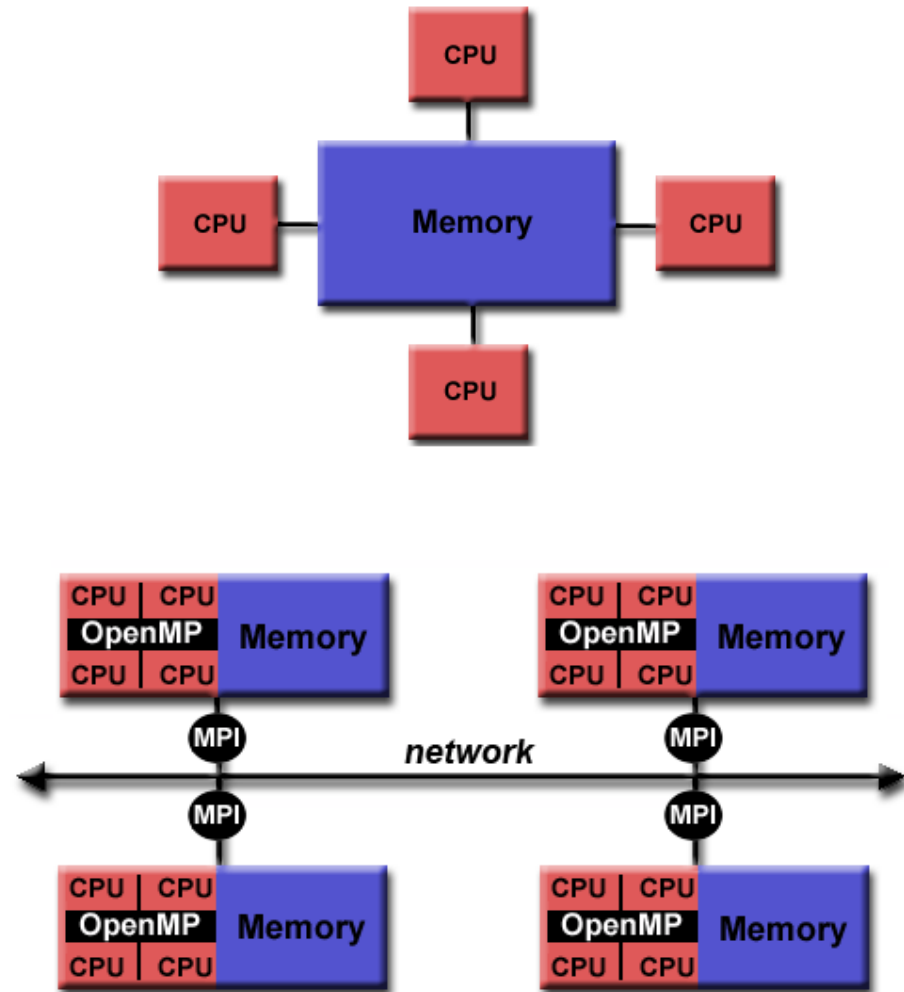
credits: [https://computing.llnl.gov/tutorials/parallel\\_com](https://computing.llnl.gov/tutorials/parallel_com)



# Running on a Supercomputer

## Levels of parallelism

- Task parallel
  - many independent runs
  - needs orchestration
  - for monte-carlo, parameter sweeps
- Shared memory
  - always within one batch node
  - uses threads
  - often implicit
- Distributed memory
  - can use one or more batch nodes
  - uses separate processes
  - almost always using MPI
  - for PDE problems, time stepping



## One last step...

- Get ready for the real hands-on example: hemodynamics with HemoCell!
- Please copy the required files to your home directory

```
nct00008@login1:~> cp
/gpfs/projects/nct00/nct00008/DATA_HEMOCELL/H
emoCell-master.zip ~/.
nct00008@login1:~> cp
/gpfs/projects/nct00/nct00008/DATA_HEMOCELL/pa
labos_dev.tgz ~/.
```

- Enjoy the next sessions!

# Acknowledgements

Marco Verdicchio + SURFsara advisors

BSC support



**SURF SARA**