# Introduction to High Performance Computing (HPC) – Session 1

using the "Computational Shared Facility" (CSF)

Course materials / slides available from:

https://ri.itservices.manchester.ac.uk/course/rcsf/

Research Platforms, IT Services

its-ri-team@manchester.ac.uk

https://ri.itservices.manchester.ac.uk/csf3/

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## Housekeeping

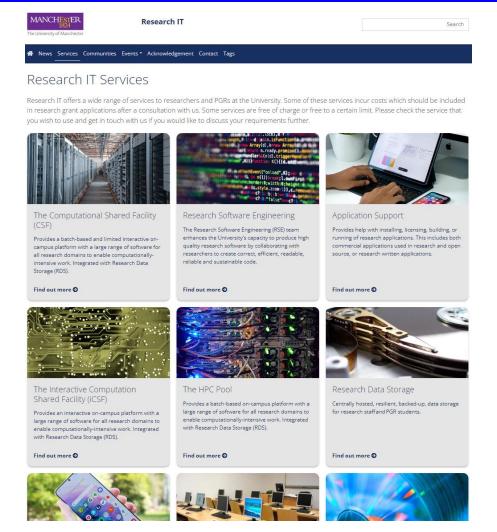
- Please let me know if you're leaving
  - 10am 12:30pm (practical sessions 1, 2, & 3)
  - Lunch approx 12:30pm 1:30pm
  - 1:30pm 4pm (practical sessions 4 & 5)
- 1-to-1 help is available if needed during exercises.
- Power adapters are available for the purposes of charging laptops, please be considerate of other users and be careful of any trailing leads.
- Got a question at any point? PLEASE ASK!!

#### Feedback

- Your feedback is important to us!
- Please give feedback on this course
  - Quick form at <u>https://goo.gl/forms/zfZyTLw4DDaySnCF3</u>
     (choose "Introduction to HPC (Using CSF)")
  - Feedback is important to help us improve our courses
  - Records your attendance on the course

#### Who we are - Research IT

https://research-it.manchester.ac.uk/services/



## Who the course is for - everyone

- Those new to HPC / research computing / who want to try the CSF out?
  - We'll introduce you to these topics and you'll try it out today
- Maybe your supervisor asked you to get a CSF account
  - We'll teach you how to use it
- Those who have used CSF already but want to know more?
  - Parallel jobs, job arrays, the batch system, ...
- CSF *usage* is today's focus so you can use it effectively in your work
  - not theoretical aspects of HPC
    - but we'll explain some of the basics to help you make good use of the resources
  - not parallel software development or version control
    - But we'll show you how to run high-end parallel applications
  - not Linux installation / administration
    - but we'll cover the basic Linux commands needed to use the CSF
  - not the specifics of the software you plan on using
    - ask your PI/supervisor for help with that

Invest a little time now, get results much faster!

Course materials at <a href="https://ri.itservices.manchester.ac.uk/course/rcsf/">https://ri.itservices.manchester.ac.uk/course/rcsf/</a>

#### WHY & WHAT ...

High Performance Computing: why use it & what is it

### Motivation: Why use HPC (and the CSF)?

- Some (most?) research computation not suitable for your desktop/laptop
  - Takes too long to run
  - Needs more memory
  - Uses too much disk/storage space
- Use advanced centrally-run UoM hardware
- Eventually use regional / national supercomputers

Do not let the size/capacity/power of your computer dictate the size and complexity of the models/simulations/systems/problems you are solving!

#### What is HPC?

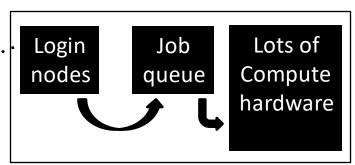
- insideHPC.com
  - High Performance Computing most generally refers to the practice of aggregating computing power in a way that delivers much higher performance than one could get out of a typical desktop computer or workstation in order to solve large problems in science, engineering, or business.
- HPC systems are usually a cluster of compute nodes (with some extra items such as login nodes, storage, networking)
  - The CSF matches this description!

## A new way of working!

VS







#### Running on a desktop

- You can fire up a GUI, run an app immediately. BUT:
- Got enough memory, cores, storage?
- Need to keep the PC to yourself (public cluster PC?)
- For several days?!
- Only one "job" (simulation, analysis) at a time?

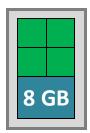
#### Logging in to the HPC system

- Submit "jobs" to the queue
- Jobs wait until selected to be run
- Jobs run on high-end hardware (lots of cores, memory, disk)
- Jobs run safely for days
- Many jobs can run at once
- Can log out any time (jobs still run)
- Log in to check on progress, get results

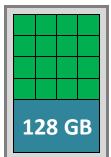
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#### Job Example: Finite Element Analysis

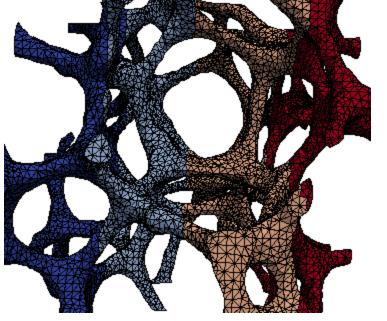
- Perform stress analysis on 3D mesh
  - App s/w splits mesh into chunks
  - It performs calculations on chunks, in parallel
    - Faster and/or larger problem size

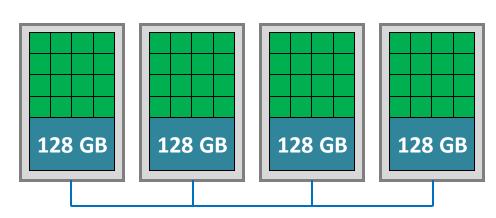


Desktop: 4 cores, 8GB RAM. 8 days to complete



Single HPC compute node: 16 cores, 128GB RAM ~2 days to complete



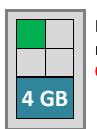


Source: Professor Paul Mummery (MACE) using ParaFEM software http://parafem.org.uk

Multiple HPC compute nodes: 64 cores, 512GB RAM ~0.5 days to complete

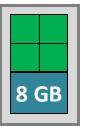
#### Job Example: Image Analysis

- High Throughput Computing
  - Not all apps do "HPC" / parallel
  - But you have lots of data
  - Each image takes 1hr to process
     (and are independent can be processed in any order)



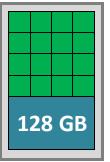
Laptop: 1 copy of software running on 4-core CPU.

Over 1 year to complete!!



Desktop: 4 cores, 4 copies of software running.

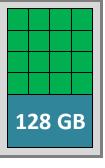
~100 days to complete!

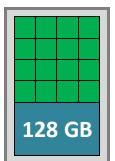


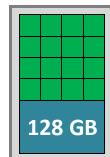
Single HPC compute node: 16 copies of software running.

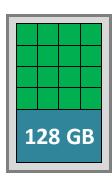
~26 days to complete











Multiple HPC compute nodes: 64 copies of software running.

~6 days to complete

Example: 10,000 image scans to be analysed by an image processing application. Each image takes *1 hour* to process.

## What we'll be using today - the CSF

- Q: who has used the Computational Shared Facility (CSF) before?
- Currently: CSF3
  - A large Linux cluster system
  - 8,644 CPU cores (all Intel Xeon CPUs)
  - 100 GPUS (68 Nvidia v100s + 32 Nvidia A100s)
  - Got big datasets to process? Can run large-memory jobs
  - (we'll cover all of these details throughout the course)

BUT, you <u>do not</u> have to be running huge parallel jobs, or be a Linux / HPC expert, to use our systems and to benefit from "HPC" and the CSF

#### Who can use the CSF?

The following info is mainly for people who may want to "buy in" to the CSF. Your PI/supervisor or School may well have already done this! If interested, ask us at the end of the course.

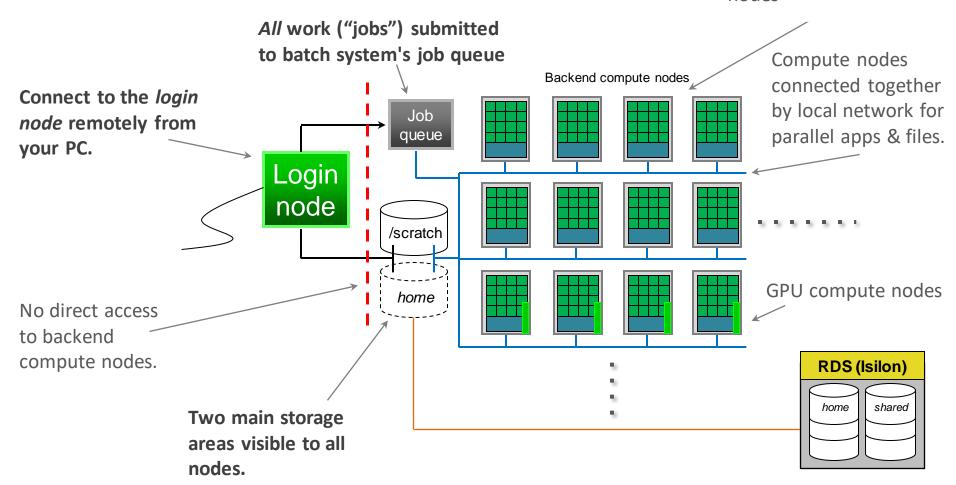
- CSF uses a shared funding model
  - Researchers/academics/schools contribute financially to buy compute hardware
  - All h/w pooled so that all users can access the h/w
  - H/w not associated with individuals so it can always be in use as long as there are jobs to run!
- The time it takes to run all of your jobs depends on the size of the contribution with which you are associated
  - A research group that contributes more will be able to get more jobs done sooner
  - Managed automatically by the batch system you just submit jobs!
- Some limited 'free at the point of use' access for noncontributors

#### **CSF: THE BASICS...**

Hardware, OS, logging in, security, home filesystem, copying files, Linux, GUIs

## What is the CSF?(more details)

- Computational Shared Facility
- A batch compute cluster to run your "jobs" (simulations, analysis,...)
- Here are the main components you'll learn about: 100s of compute nodes



## **Compute Nodes**

#### The CSF is a cluster of 100s of "compute nodes"

Think of a compute node as **high end** PC (the CSF provides 100s of these!):

- Multi-core CPUs (e.g., 16, 28, 32 cores)
- Lots of RAM (e.g., 128GB, ..., 2TB!)
- Network (possibly fast InfiniBand n/w)
- OS (Linux)
- Local disk (for temp files)
- Maybe a GPU

Not all nodes have the same h/w. But the CSF hides these details – let it choose which nodes you use, or you can specify certain requirements.



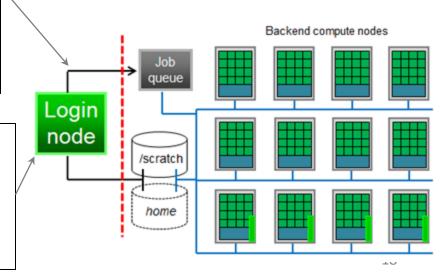
- Some applications can use more than one node! (e.g., for big data / simulations)
- All compute nodes see the same files
- You don't need to transfer files to a compute node!

## Login Nodes

- CSF: More than 400 backend compute nodes
  - Too many for you to find a free compute node to use
  - Simpler: connect to a login node, let job-queue find a node

#### **Key concept!**

- Do: Submit work ("jobs") to be run
  - No direct access to compute nodes
  - Submit jobs to the job queue.
  - The system will run your work on available compute nodes meeting your requirements.
- Do not: run applications on the login node
  - Shared by all users, not much memory
  - For lightweight tasks (job submission, file transfer, ...)



## Some pictures of the CSF



Compute nodes Local storage —





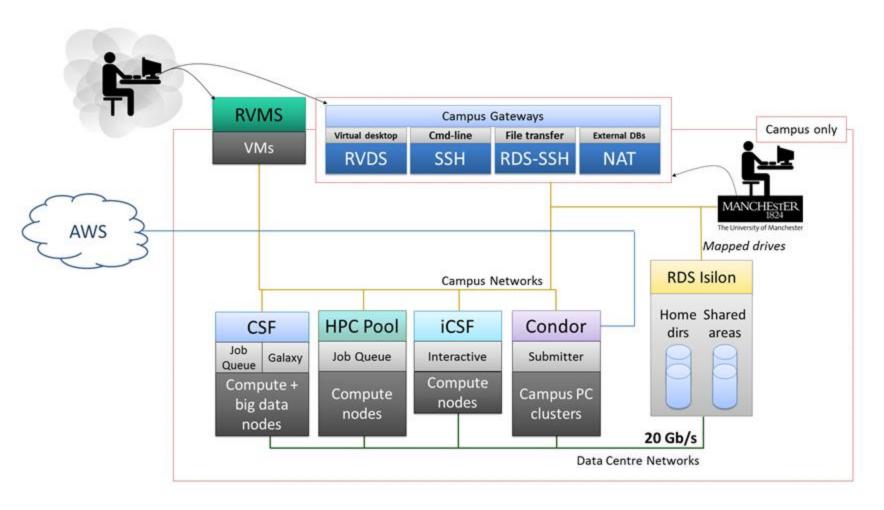
Networking connects everything



#### Features of the CSF

- Variety of software applications (120+) & compilers
  - Gromacs, MATLAB, Abaqus, Bowtie2, Gaussian, Fluent, many more
- 1-core (serial) jobs or many-core (parallel) jobs
  - Do more work. Get it done sooner (see later!)
- Lots of RAM: 64GB, 128GB, 192GB, 256GB, 512GB, 1TB, 1.5TB, 2TB, 4TB (!!) compute nodes
- Backed-up file storage (no more USB disks!!!!!!!)
- Hardware failures don't stop your research
- Leave computational work running (for days)
- 8,644 cores currently in the system
- Dedicated support team

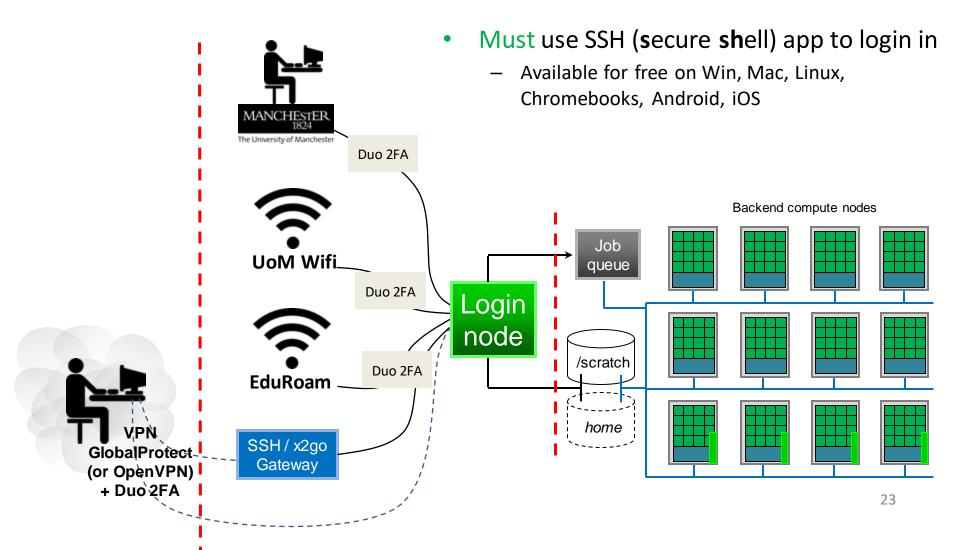
## Part of the *Computationally Intensive Research* Ecosystem



## Logging in ...

Let's get access to the system

## Where can I log in from?



## Security (1)

- The CSF has a private (campus only) IP address
  - Firewall also controls connections to and from the system
- When you are on campus
  - Connect from any PC/laptop with a wired connection, or University WiFi, or Eduroam WiFi
  - Does not matter if using VPN or not, but you will always be asked to authenticate using your 2FA (DUO) device, e.g.

```
@i-uliit636838l ~ $ ssh -X @csf.itservices.manchester.ac.uk
Password:
Duo two-factor login for

Enter a passcode or select one of the following options:

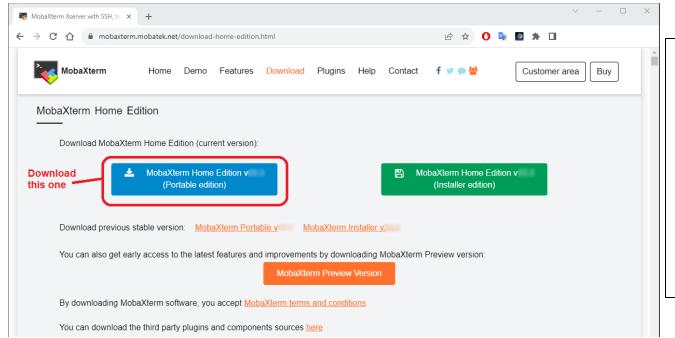
1. Duo Push to +XX XXXX XX6353

Passcode or option (1-1): 1
```

- When you are off campus
  - May not be available to all UoM account types today
  - Sign-in to University Global Protect VPN + DUO 2FA
  - Allows you to make a direct connection to CSF
- Further documentation: <a href="https://ri.itservices.manchester.ac.uk/csf3/getting-started/connecting/">https://ri.itservices.manchester.ac.uk/csf3/getting-started/connecting/</a>

#### Connect to CSF from Windows

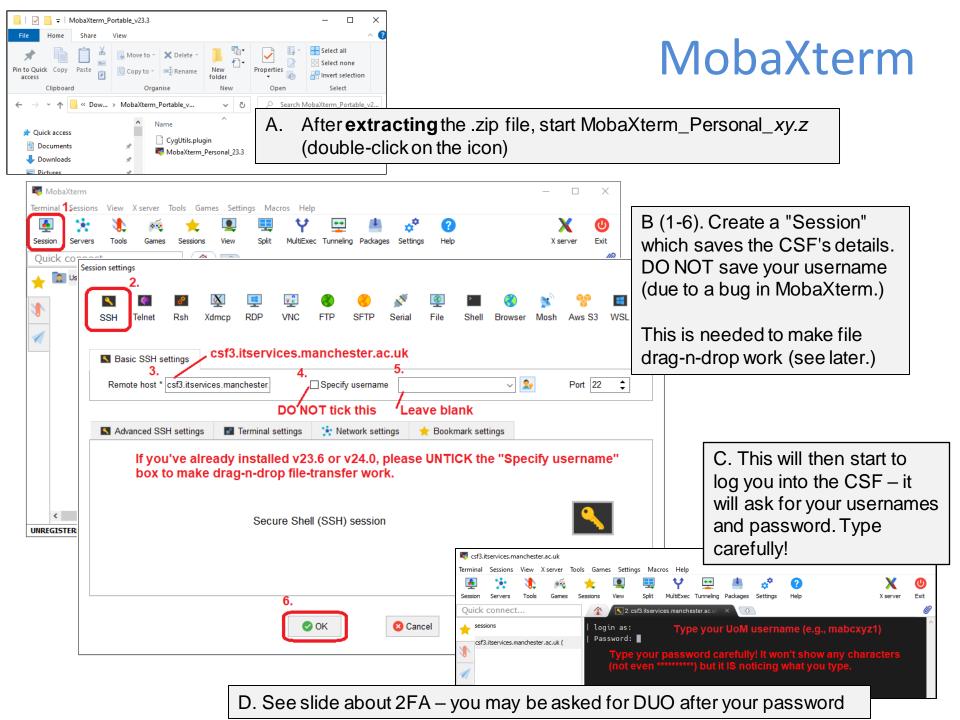
- Job queue //scratch
- Access the CSF from a PC / laptop using an SSH (Secure Shell) app
  - Sometimes called a "terminal".
  - There's no web-site or other fancy GUI on the CSF use the "command-line".
- Windows users need to install a free terminal app called MobaXterm
- https://mobaxterm.mobatek.net/download-home-edition.html
   the Home edition (portable edition) does not require Administrator rights just extract the small .zip file in your P-Drive or USB stick for example.

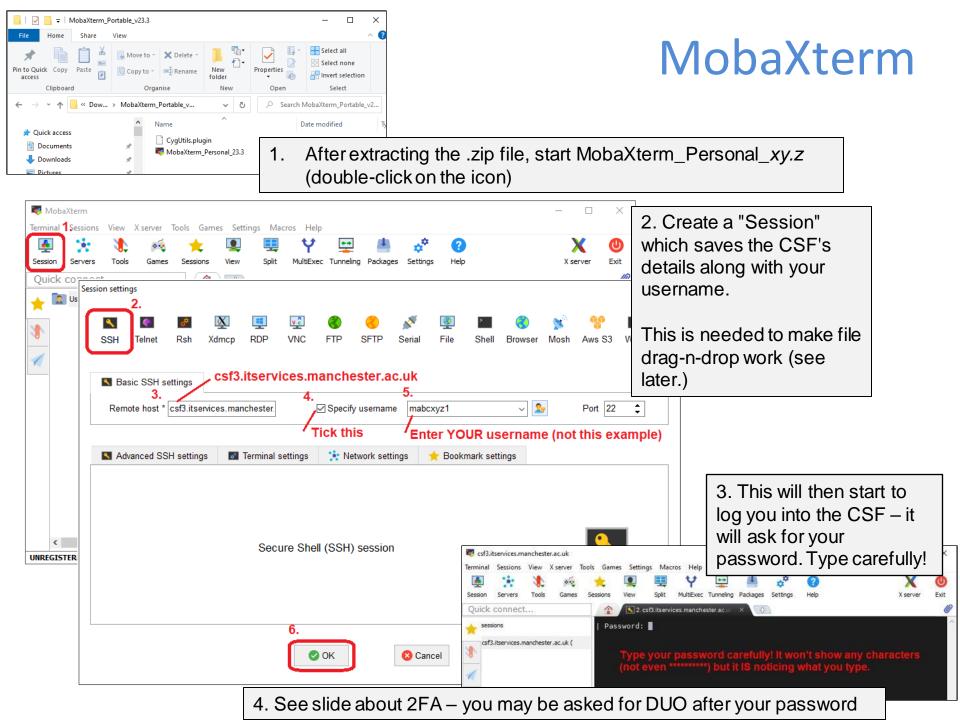


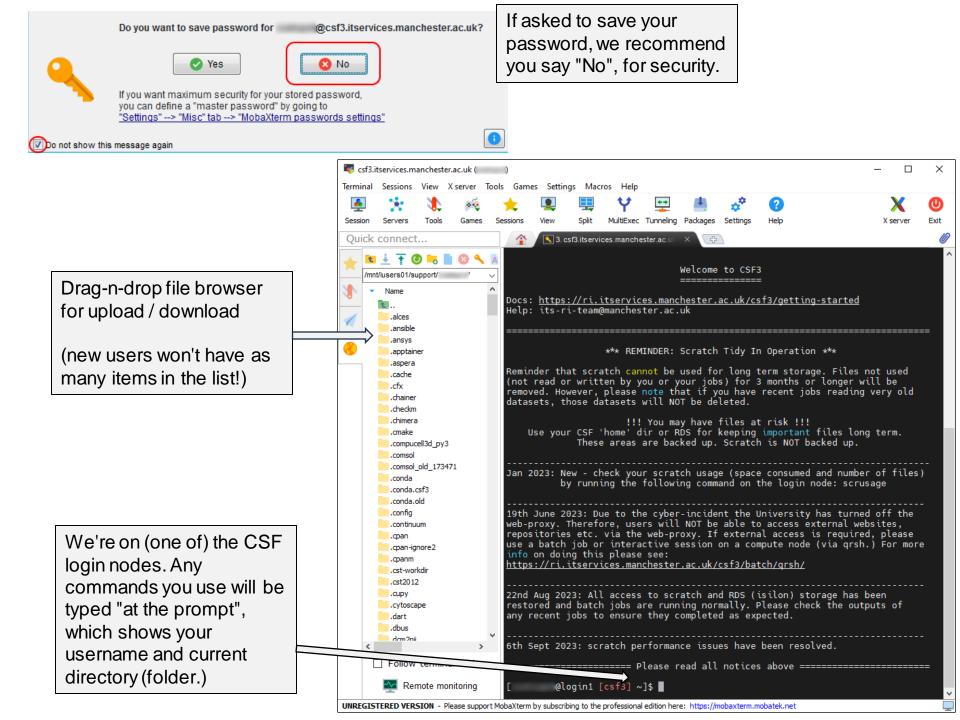
- Download using the blue box.
- Once downloaded, right-click on the .zip file and select:

"Extract all ..."

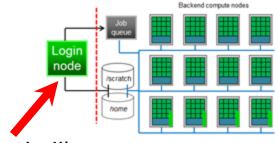
This will *unpack* the .zip file to a folder.



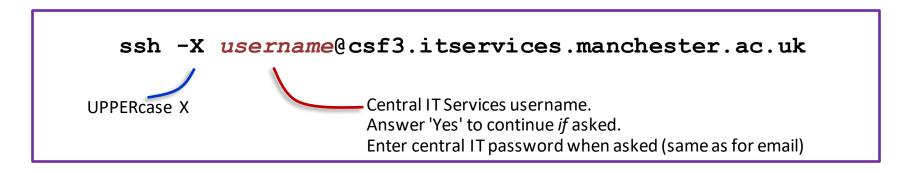




#### Connect to CSF from Linux

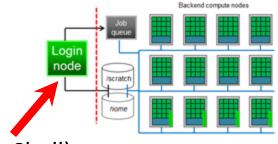


- Access the CSF from a PC / laptop using an SSH (Secure Shell) app, eg a 'terminal'.
  - There's no web-site or other fancy GUI on the CSF command-line for now.
- **Linux** users have a *terminal* application by default
  - Start a Terminal and type the following command:

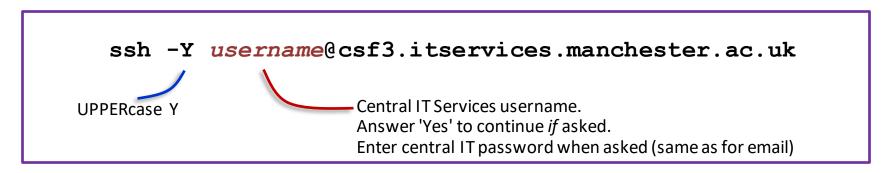


Finished using CSF? Log out with: logout or exit

#### Connect to CSF from a Mac

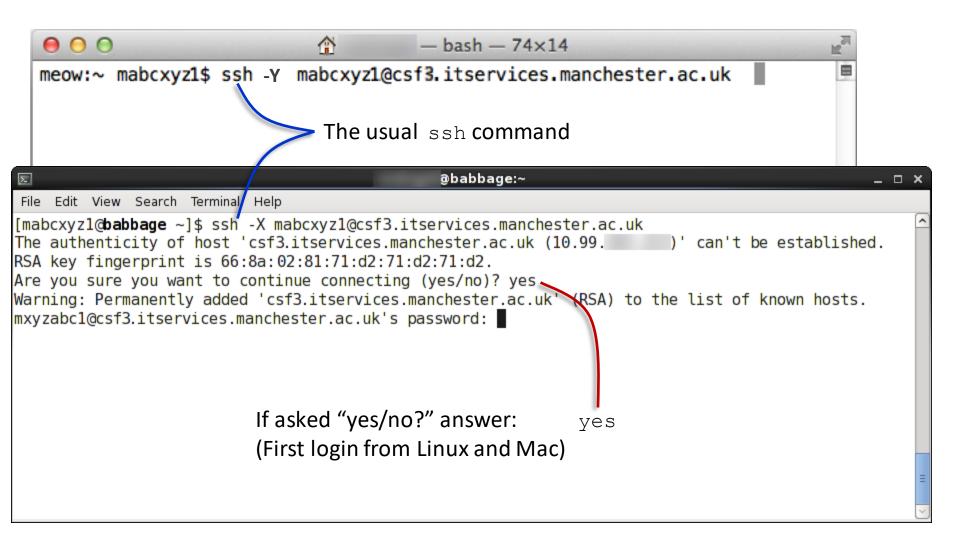


- Access the CSF from a PC / laptop using an SSH (Secure Shell) app, eg a 'terminal'.
  - There's no web-site or other fancy GUI on the CSF command-line for now.
- Mac users have a terminal application by default
  - You will need to install X-Quartz first
     <a href="https://www.xquartz.org/">https://www.xquartz.org/</a> (install then you should reboot your Mac)
  - Start a *Terminal* app and type the following command:



Finished using CSF? Log out with: logout or exit

## Linux / Mac Terminals



#### DUO 2FA (when on-campus)

 When on-campus, after you enter your password, all login methods will then ask about DUO:

Duo two-factor login for mabcxyz1

Enter a passcode or select one of the following options:

1. Duo Push to +XX XXXX XX7890

Passcode or option (1-1): 1

Success. Logging you in...

(the message of the day is now displayed)

[mabcxyz1@login1 [csf3] ~]\$

Type **1** (and press Enter) in your ssh app to generate a DUO *push* to your device.

Then accept the push on your device.

You are now logged in.

## What you see when you log in

- CSF uses CentOS Linux (c.f. Red Hat EL)
  - Command line requires the input of commands,
     can be a little scary at first to new users
  - A welcome message of the day announcements
  - The system awaits input/commands from you at a prompt (after you've logged in):

Type vour

commands

at the prompt

```
[username@login1 [csf3] ~]$
or [username@login2 [csf3] ~]$
```

- Learning Linux commands (more later):
  - <a href="https://www.chm.bris.ac.uk/unix/">https://www.chm.bris.ac.uk/unix/</a>

## Security (2)

- It is NOT permitted to share your CSF account
- CSF uses your IT password i.e. same as needed to access UoM email, ...
  - NEVER share it with ANYONE, including IT staff and your supervisor
  - Forgotten it? You can reset it via the IT Account Manager. Will affect all systems that require it.
    - https://iam.manchester.ac.uk/
- Reminder: Other general safety measures
  - —Install a virus scanner
    <a href="https://www.itservices.manchester.ac.uk/cybersecurity/advice/virusprote-ction/">https://www.itservices.manchester.ac.uk/cybersecurity/advice/virusprote-ction/</a>
  - By aware of phishing emails
     https://www.itservices.manchester.ac.uk/cybersecurity/advice/phishing/

#### **PRACTICAL SESSION 1**

Logging in

### Practical Session 1 – Logging in

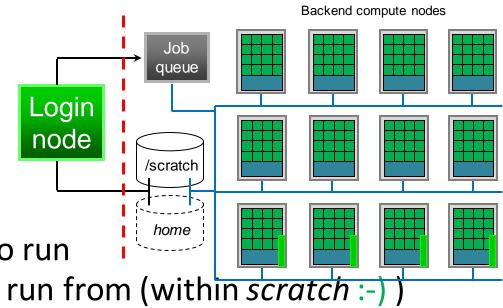
- Exercise 1 sheet (pdf) available at: https://ri.itservices.manchester.ac.uk/course/rcsf/
- Tip: During login, when you are prompted for your password, type carefully you will
  not get a cursor that moves or display any \*\*\* as you press the keys. But it IS noticing
  what you type!
- Tip: Once logged in to the CSF, Linux does not always display something after you have entered a command – this is usually a good sign – your command worked, or there was nothing for it to do. If you've got something wrong, it will usually tell you via an error message.
- By the end of this practical session, everyone should have successfully logged in to the CSF!
- PLEASE ASK FOR HELP IF YOU RUN IN TO ANY PROBLEMS WE ARE HERE TO HELP!

#### **RUNNING JOBS**

Doing real work on the CSF

#### Jobs, Jobscripts and the Batch System

We want to do computational work - "jobs"



- You decide:
  - What program(s) to run
  - Which directory to run from (within scratch:-)
  - What resources it needs (#cores, CPU type, memory)
- Write these requirements in a jobscript
- Submit your jobscript to the batch system (SGE)
- SGE decides exactly when and where the job runs

## Warning: The login nodes

- Do not run computational work here:
  - Not enough cores
  - Not enough memory
  - 100+ users connected, so running work causes serious problems
- You can do the following:
  - Transfer files on and off the CSF
  - Set up and submit your jobs (covered in next few slides)
  - Basic data processing/viewing
- Computational work running on the login node will be killed without warning!

#### Reminder: Creating Jobscript files (1)

- You need to be able to create a small text file to describe your job
- Run gedit on the CSF login node a simple text editor
  - Creates and saves the file on the CSF
  - gedit is similar to notepad (other Linux editors: emacs, vi)
- Once logged in to the CSF type:
  - gedit & '&' allow you to carry on using the command-line. Try it without to see.
  - Navigate to a file or start typing and then save

#### Can I Write Jobscripts on Windows?

- A warning about Windows text files (EG: in notepad)
  - There's an inconsistency over the (hidden) end-of-line characters in text files:
    - Windows: CR (carriage return) + LF (line feed)
    - Linux/Unix: LF (line feed)
  - The extra CR from Windows is a problem in jobscripts.
     Causes your job to fail immediately.
- Solutions
  - Use gedit on the CSF login node (writes Linux text files)
  - Or use notepad, upload then run dos2unix myfile.txt
    - Use only on jobscripts. Do not come to rely on this it is too easy to forget to do it – use gedit!

# A simple Jobscript – Serial (1 core)

myjob.txt



#! on first line only (a special line)

#\$ indicates a batch **system parameter** to specify our job requirements. We'll use various combinations of these.

# lines are just comments - anything on the line after it will be ignored.

```
#!/bin/bash --login
#$ -cwd
#$ -N myjob •
#$ -1 resource
# Let's do work
date
```

Actual Linux commands we run in our job. They will execute on a compute node.

hostname

date

sleep 120

First line indicates we use the *bash* script language to write our jobscript.

- -cwd indicates we'll run from our current (working) directory. Input / output files will usually be found here.
- -N (optional). Set the jobname. Otherwise will use name of your jobscript as the name.
- -1 (optional) used to add extra resource requirements e.g. memory, time limits

#\$ -I course only works on the day of a course.

## Submit Jobscript to Job Queue

Submit the jobscript from the login node with:

```
qsub jobscript # EG: qsub myjob.txt
```

 You will be given a unique JobID (6/7-digit number)

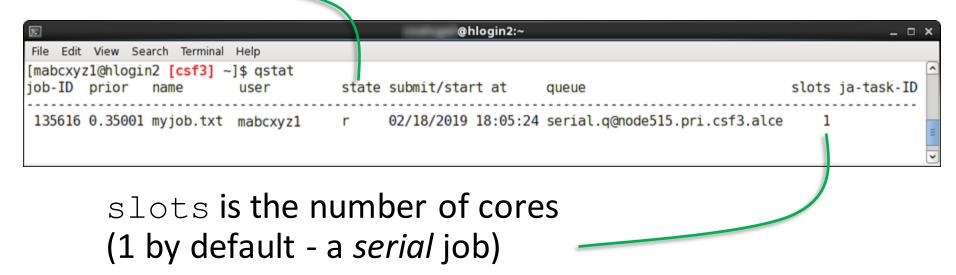
```
Your job 598052 ("myjob.txt") has been submitted
```

- You can then:
  - carry on with other work, submit other jobs that run other applications without disturbing previous jobs
  - log out of the CSF and your jobs will still run

## Check status of your jobs

To see your job(s) in the batch system, run:
 qstat

state is either queued (qw), running (r) or error (Eqw)



## Serial (1-core) Job Properties

- Our simple example:
  - Serial (unless specified, only 1 core is used see later)
  - Standard memory (no #\$ line asking for more)
    - We get: 4GB RAM (per core we're just using one core!)
  - Standard 7 day runtime (no #\$ line asking for shorter)
- Default: standard serial jobs will be placed on:
  - Intel nodes (Ivybridge or Haswell)
    - The system looks for a free core on a list of compute node
- We didn't use any: #\$ −1 option
  - EG: #\$ −1 mem256 to land on higher memory node

## Optional Serial Job Resources

https://ri.itservices.manchester.ac.uk/csf3/batch/serial-jobs

Jobscript flag	Description	
-l haswell	5GB/core haswell	NB: This flag is not so useful. The other flags below <i>are</i> .
-l short	1 hour runtime limit, test or pre/post proc	
-1 mem256	16GB/core haswell (default)	
<pre>-1 mem512 -1 mem512 -1 ivybridge -1 mem512 -1 haswell</pre>	32GB/core ivybridge or haswell (system will choose) 32GB/core ivybridge 32GB/core haswell	

- 7-day runtime limit on jobs unless otherwise indicated in table.
- Our simple jobscript did not use any of the above. Not needed in most cases.
- If you limit a job by node-type or memory it may wait longer in the queue.
- You will see that the example jobscripts in the exercises have #\$ -1 course
  - Only for use today (reserved nodes on a teaching day).
  - Remove if you are practising outside of today.
  - Never use on your regular jobs (they'll wait forever!)

## So where did my results go?

- If qstat returns no output means job has finished!
- Three possibilities:
  - 1. If app prints to screen: A text file called *jobname.oJobID* 
    - Jobname is the name of your jobscript script (or -N name setting)
    - JobID is the number of your job
    - Previous example: myjob.txt.o598052
    - (Technically: "the std output stream is redirected to the file")
  - 2. An output file specific to your application
  - 3. Your job had a problem or failed: it may be reported in one of the above files or in *jobname.eJobID* 
    - (Technically: "the std error stream is redirected to the file")
- Various options to view the file (they are plain text):

```
cat filename
less filename (allows you to page through with spacebar)
gedit filename (not recommended if it is large)
```

## Why is my job still waiting?

- Your job will wait until there are cores available (meeting your jobscript's requirements).
- All jobs can run for up to 7 days (wallclock)
  - (#\$ -1 short jobs up to 1 hour)
  - Jobs are starting and finishing all the time
- Initially frustrating (perhaps) but advantages:
  - You can log off, switch off your PC and your job will stay on the CSF. Log in later to check on job / collect the results.
  - You can submit many jobs.
    - They might not all run at once but SGE will decide...
  - More than one may run at the same time (make sure you have different files/folders for each job).

## Many Users Sharing the CSF

- 100s of users running 1000s of jobs
- SGE gives each job a priority: depends on
  - Size of research group's / school's CSF contribution
  - Amount of work already put through by that group and by you as an individual (this month)
- The time for your job to start depends on
  - Priority
  - Availability of requested resource (is CSF busy?)
- Jobs submitted after yours may start before yours!
- A few Jobs may never start
  - SGE tries to spot errors in jobscripts when you run qsub
  - Some may still get through then never run
- We try to ensure that if you submit some jobs, some of them will start within 24 hours.
  - We make a check every morning of the waiting jobs

## How busy is it?

- The system is usually very busy.
- However, jobs frequently finish, allowing waiting ones to start
- To see all the jobs for everyone

```
qstat -u "*"
```

- Note: all jobs shown as one list by qstat
  - It displays running and waiting jobs
  - Your job is not necessarily stuck behind all others above yours in the qstat output.
  - CSF is split into a few chunks the very big jobs do not compete with the smaller jobs for cores
- Do not try to guess when is a good time to submit your jobs.
  - If you have work ready to run, submit it
  - If your jobs are not in the queue the scheduler cannot consider them
  - You will waste time, not gain it, by not submitting

# What does Eqw mean? How do I delete a job?

- qstat reports your job as 'Eqw'
  - System tried to start it, but something went wrong
  - Usually no output file (job hasn't run) to indicate what happened, use:

```
qstat -j JobID | grep error
```

- Error can be cryptic. Most common causes:
  - Missing directory (cannot chdir to ...)
  - You created your jobscript on windows (exec error)
  - Unusual characters or spaces in file and directory names
  - No disk space on the filesystem did you forget to use scratch?
- Detailed advice:

https://ri.itservices.manchester.ac.uk/csf3/batch/job-monitoring

• To delete an Eqw job (it'll never run), or one you simply no longer want:

```
qdel jobID
```

#### **PRACTICAL SESSION 2**

Serial job

#### Practical Session 2 - Submitting jobs

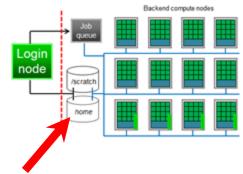
- Follow the handout 'Practical Session 2'
  - Use qsub to submit a simple serial job on the CSF
  - Use qstat to look at the queues
  - Use qdel to kill jobs
  - Use qacct to look at finished jobs
- Exercise sheet (pdf) available at:

https://ri.itservices.manchester.ac.uk/course/rcsf/

## **CSF STORAGE (FILESYSTEMS)**

Where to store your files...

## Storage – Home filesystem



 Upon login, automatically placed in your home directory (folder)

/mnt/iusers01/group01/username

- Limited space, quota shared by everyone in the group
- Uses the Research Data Service (networked storage)
  - Large files can be slow-ish to read/write (implications for jobs)
- Which directory (folder) am I currently in?

pwd

How much space am I using? (Linux commands!)

```
du -sh dirname # Can take a while
```

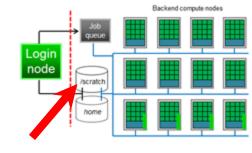
How big is that file?

```
ls -lh filename # Letter el not number 1
```

How much space is used/free overall?

```
df -h . # The . is important!
```

## Filesystems - Scratch (1)



- Filesystem local to CSF for:
  - Temporary files can be huge
  - Running jobs from (it is faster!). Recommended!
- Shared by all CSF users, but we have 1.2PB
- Tidy up after each job finishes
- Clean-up policy applies: files that have not been accessed for the past 3 months may be deleted automatically
- Not backed up!
  - Move/copy important results to home area
  - Not considered safe for long term storage hardware failure could cause data loss

## Storage - Home filesystem

- Home is backed up and mirrored to another datacentre
  - Keep important files here (results, jobscripts, source code, ...)
  - Deleted a file by mistake? <u>its-ri-team@manchester.ac.uk</u> can tell you how to retrieve it
- Only you can access your home directory
  - File permissions can be used to give others access
  - Contact <u>its-ri-team@manchester.ac.uk</u> if you want advice on this as they can be complex
- Do not run jobs from your home area (see later)
  - Can generate a lot of files, some of them large
  - Using up all of the shared space will make your colleagues unhappy!
  - Consider compressing large (text) files with gzip

## Filesystems - Scratch (2)

Using scratch is easy: after log in, change to it:

```
cd ~/scratch
```

- Uses a 'symlink' (short cut) in your home dir to /scratch/username
- Create a directory (now we're in scratch):

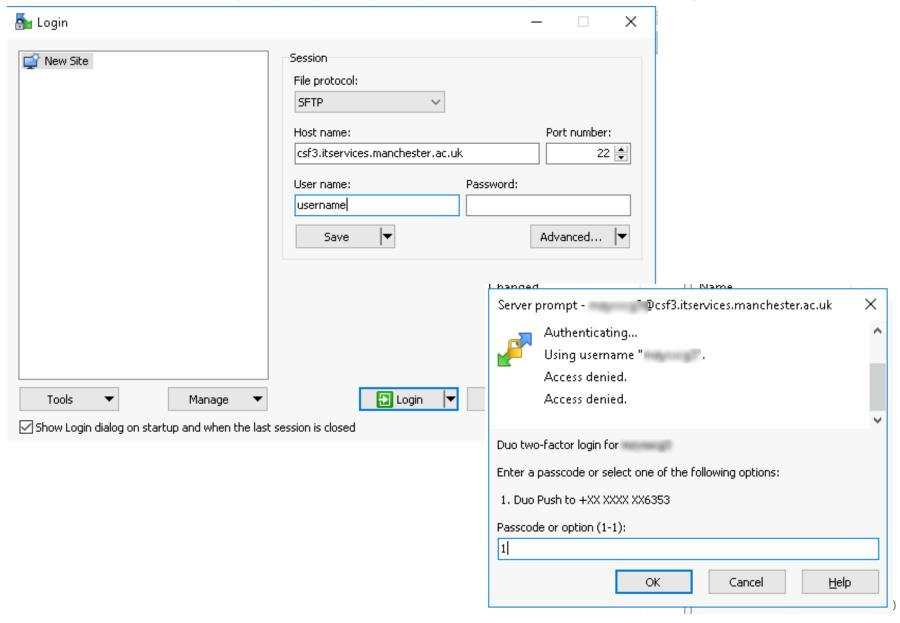
```
mkdir myjobdir
```

- Put all files relevant to your job in that directory and run your jobs from there - we'll try this out soon...
- All compute nodes see the same scratch area

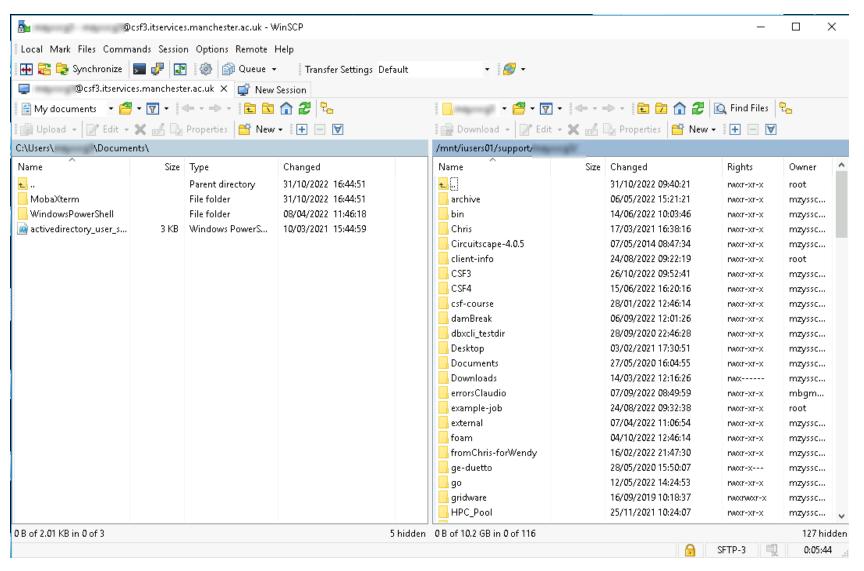
#### File transfer with MobaXterm

2. First select files in the MohaXterm browser. **3**. The *Upload* button opens a file-browser Then the **Download** button opens a file-browser to select files on your computer to upload to to select a destination folder on your PC. **CSF** (current directory). - - X @hlogin2:~ Terminal Sessions View X server Tools Games Settings Macros ... Quick connect... @hlogin2:~ [2019-02-18 16:55.51] ~ mabcxyz1.babbage] > ssh mabcxyz1@csf3.itservices.manchester.ac.uk mabcxyz1@csf3.itservices.manchester.ac.uk's password: /mnt/iusers01/support/ Last login: Mon Feb 18 10:40:46 2019 from nyx5.itservices.manchester.ac.uk Name This is the CSF3 abagus plugins adtemp altera \*\*\* Please note - we are still working on some aspects of the configuration. archive Docs: http://ri.itservices.manchester.ac.uk/csf3/getting-started babbage Help: its-ri-team@manchester.ac.uk Fri 25th Jan - Batch limits have been adjusted to accomodate the move of CATSettings users from CSF2 to CSF3. components 1. Drag-n-drop cronlogs csf2vpn files from 'SCRATCH' STORAGE MUST NOT BE USED FOR LONG TERM STORAGE - PLEASE READ http://ri.itservices.manchester.ac.uk/csf3/filesystems/scratch-cleanup/ Windows Explorer or Follow terminal folder [mabcxyz1@hlogin2 [csf3] ~]\$ Desktop UNREGISTERED VERSION - Please support MobaXterm by subscribing to the professional edition here: https://mobaxterm.mobatek.net to/from here.

#### File transfer with WinSCP



#### File transfer with WinSCP



#### home file transfer with Linux / Mac

(can also type in a local MobaXterm window)

Transfer a file from your computer to your CSF home dir

scp myfile.txt username@csf3.itservices.manchester.ac.uk:~/training/

Exercise: Create a file on your PC named myfile.txt containing some text then transfer it to the CSF.

Destination directory on the CSF

important!

Destination directory on the CSF

is shorthand meaning "your home directory"

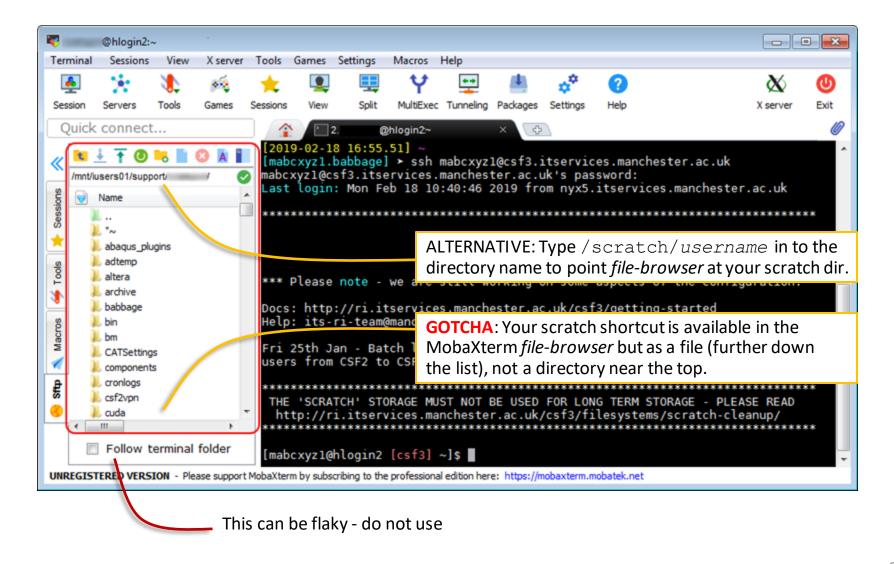
If no destination after the: then uses "your home directory"

Transfer a file from your CSF home dir to your computer

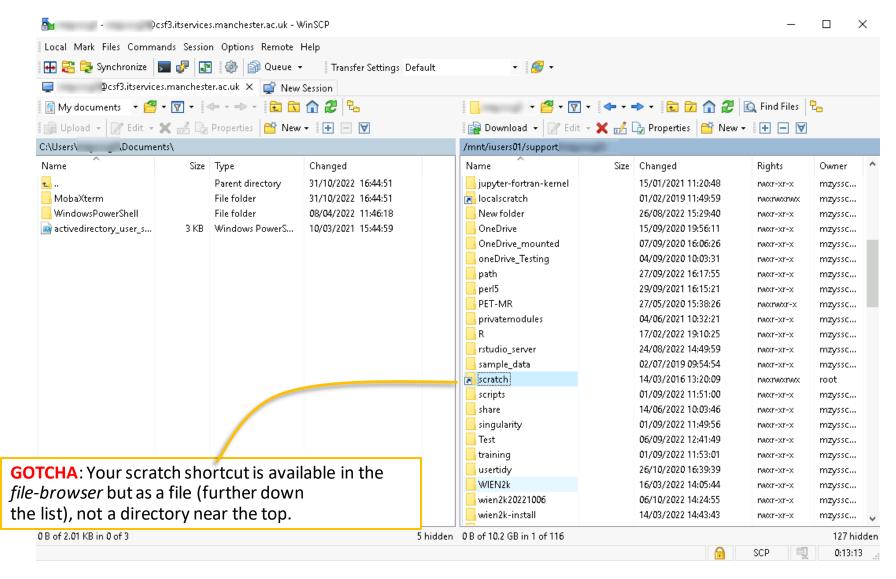
```
The. is shorthand meaning "the current directory" on your computer
```

Change directory & filenames...

## Accessing scratch with MobaXterm



## Accessing scratch with WinSCP



#### scratch file transfer with Linux / Mac

(can also type in a local MobaXterm window)

- Very similar to commands used earlier for our home directory
- Transfer a file from your computer to your CSF scratch dir

```
scp file2.txt username@csf3.itservices.manchester.ac.uk:~/scratch/

We give a destination after the: meaning "use my scratch shortcut"
```

Omit the destination to transfer to *home* dir.

Transfer a file from your CSF scratch to your computer

```
Now you get a copy with a different name on your computer.

Use . to keep the same name (results2.out)
```

Change directory & filenames as required...

## Extra Storage Space (Optional)

• Some research groups have extra space, example path:

```
/mnt/eps01-rds/group/username
```

No shortcut from your home? To access it use:

```
cd /mnt/eps01-rds/group/username
```

To create a shortcut (named data) in your home area:

```
cd ~
ln -s /mnt/eps01-rds/group/username data
```

- Also backed up
- Often many TB, but again shared by everyone else from your group
  - Be fair in your usage

#### Additional filesystem/file transfer info

- We have additional info about how to manage your files and your disk usage:
  - https://ri.itservices.manchester.ac.uk/userdocs/file-management/
- Docs about file transfer:
   https://ri.itservices.manchester.ac.uk/userdocs/file-transfer/
- If you need to transfer a lot of files or big files to and from the CSF please do not do it on the login node
  - Ask for an account on the RDS-SSH service
- Many file management tasks can be included in your batch jobs – see the FAQ.

#### **Basic Linux File Commands**

A good Linux tutorial is available at: <a href="https://www.chm.bris.ac.uk/unix/">https://www.chm.bris.ac.uk/unix/</a>

Command	Description
less file1 zless file2.gz	Display the content of file1 (text file) a page at a time on screen. If you've compressed file2 with gzip, no need to uncompress first. Press space to page down through a long file Press return to scroll down a line at a time Press b to scroll back up a page Press G to go to end of file Press q to quit/exit
<pre>cat file1 zcat file2.gz</pre>	Dump entire file to screen (a quick way to look at text files). If you've compress file2 with gzip, no need to uncompress first.
gedit file1	Edit file1 using a simple graphical text editor (similar to notepad on Windows). See later for more on opening graphical programs on the CSF so that they display a window on your computer.
file filenameA	Try to tell us what type of data is in filenameA. Useful to determine the output of some program where you are not sure what type of output it has generated. For example:  file output.dat  Might be ASCII text (so we can look at it with less or gedit) or might be data (you'll need some other program to read it)
du -sh .	How much disk space is current directory (all files and subdirs) using?
df -h .	How much free space is there in the current area? 68

#### **Basic Linux File Commands**

A good Linux tutorial is available at: <a href="https://www.chm.bris.ac.uk/unix/">https://www.chm.bris.ac.uk/unix/</a>

Command	Description	
<pre>cd dir1 cd ~/dir1/dir2 cd cd</pre>	Change directory (go in to dir1 which is located in the current dir)  Go in to dir2 in dir1 in home (~ is shorthand for home)  Go up to parent directory (e.g., from ~/dir1/dir2 to ~/dir1)  Go back to home (useful if you become lost)	
pwd	Lost? Print Working Directory (display current location)	
<pre>ls ls -lh ls -lh file1 dirA ls -lh dirA/*.dat</pre>	List content (names of files and directories) of current directory List in long form (dates, file sizes, names) current directory List in long form (dates, file sizes, names) specified files, directories List in long form all files ending in .dat in directory dirA	
mkdir dirA	Make directory named dirA (in the current directory)	
cp fileA fileB	Copy (duplicate) a file (copy fileA to a new file fileB)	
<pre>mv fileC fileD mv fileE dirA mv fileF dirA/fileG</pre>	Rename a file (from fileC to fileD). Works for directories too.  Move fileE in to sub-directory dirA (dirA must exist)  Move fileF AND rename it all in one go (dirA must exist)	
rm fileH	Delete (remove) a file (caution!!)	
rm -rf dir1	Delete directory and all files (and other sub-dirs) in there (caution!!!!!)	
gzip bigfile gunzip bigfile.gz	Compress a file (becomes bigfile.gz) to make better use of disk-space. Text files usually compress well.  Uncompress previously compressed file (becomes bigfile). 69	

#### **PRACTICAL SESSION 3**

File Transfer

#### Practical Session 3 – File Transfer

- Follow the hand-out 'Practical Session 3'
  - Transfer a file from the CSF to your PC
  - Transfer a file from your PC to the CSF
  - Windows: use MobaXterm, Mac/Linux: use "scp"
    - Or, if time permits, Windows users can try option 2 https://winscp.net/eng/download.php
- This is not a 'real' world example, but:
  - You may need to generate files on your PC for processing on the CSF (e.g. an "abaqus" input file)
  - Your supervisor may give you files that you then need to transfer to CSF
- Exercise sheet (pdf) available at: https://ri.itservices.manchester.ac.uk/course/rcsf/

## Need more help with the CSF?

 Extensive documentation about all aspects of the service:

https://ri.itservices.manchester.ac.uk/csf3/

- Contact the Research Infrastructure Team <u>Its-ri-team@manchester.ac.uk</u>
- See you:
  - After lunch (in-person courses)