

Introduction to the iCSF (Incline) & CSF (Computational Shared Facility)

- WELCOME!
- Please take a seat near the front.
- Course materials can be found at

http://ri.itservices.manchester.ac.uk/course/icsf-csf/

- There is no sign-sheet. Attendance is recorded using a feedback form (see above link)
- https://ri.itservices.manchester.ac.uk/hpc-help

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Who we are - Research IT

- Central Group in IT Services offering a variety of researcher specific services
 - o HPC/Compute
 - Applications support
 - Software developers for research projects
 - Storage for Research Data
 - Virtual Machine Service
 - Advice regarding external services Training courses
 - Many other topics of relevance to researchers
- Blog and newsletter
- Events: Research IT Club, Drop-in sessions
- More Info https://research-it.manchester.ac.uk/services/



Presentation Overview

- Introduction to the iCSF (incline) & CSF
- Research Virtual Desktop Service
 - o Exercise 1
- Research Data Storage (RDS) & CIR Filesystems
 - o Exercise 2
- Using the iCSF (incline)
 - o Exercise 3
- Using the CSF
 - o Exercise 4
- Other Research IT Infrastructure Provided Services

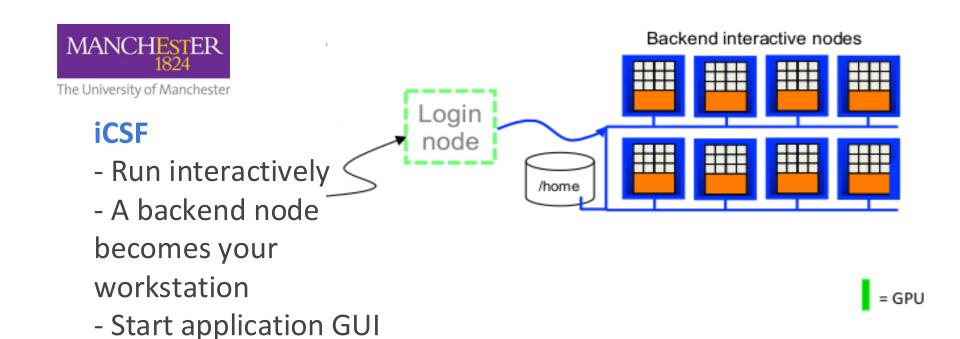


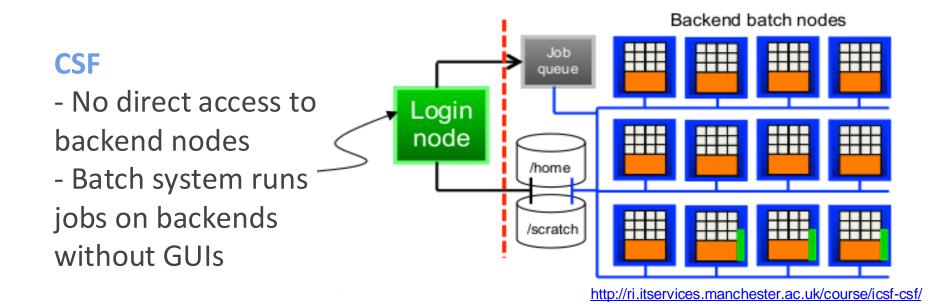
Introduction to the iCSF (Incline) & CSF



Why use the iCSF & CSF?

- Offload computational work from your PC
 - Do larger simulations/analyses that wouldn't fit on your PC
 - Work in an interactive manner using a GUI (iCSF)
- Variety of software applications
 - o MATLAB, R, Stata, SAS, many more
- 1-core (serial) jobs or many-core (parallel) jobs = speedup
 - A core is a processing unit in a CPU/processor
 - Modern CPUs have several cores: your PC/laptop may have a single CPU with 4,6,8 cores; the iCSF (32 cores per node), CSF (up to 168 cores per node), a lot more in total.
- Tackle larger problems involving larger datasets
 - Lots of memory 1.5TB, 2TB, 4TB nodes







iCSF vs CSF Differences

iCSF - run interactively

- After login a backend node becomes your 'shared' workstation
- Load and interact with applications via a GUI inc., MATLAB, Rstudio, Stata, SAS
- Easier to use then CSF
- Some command line use
- Instant access to resources

CSF - run in batch

- No direct access to backend nodes
- All jobs are submitted via a batch system jobs run on backends without GUIs
- Steeper learning curve
- Command line heavy
- Greater computational Resource
- Large Scratch Filesystem



iCSF vs CSF Similarities

- Both are a collection of servers (compute nodes) linked together to form two large systems
- Operating system on both is the same Linux (Red Hat Based)
- They share the same home directory (similar to your P Drive
 a private area for your files)
- Both platforms are built based on a contribution model, i.e. research groups cost in cost of hardware in research grant proposals.
- Limited Free at the Point of use offering (FATPOU) on both platforms.



Is the iCSF suitable for my work?

Yes if

- You need to work interactively
- Have low to moderate computational requirements
- Are unfamiliar with the Command Line.
- Are developing or testing code

It also

- Complements other CIR platforms
- Provides a first step into computational work

Unsure

email its-ri-team@mancheter.ac.uk we're here to help.



Is the CSF suitable for my work?

Yes, if

- You have moderate to high computational needs
- You need to run many 'Jobs' e.g. Simulations/Analyses at once.
- You need GPUs

You will need to

- Be a little more familiar with a Linux Command Line.
- use a batch system
- Don't worry, we will teach you some of the basics today
- We have 1 day training course specifically for the CSF
- Lots of docs, and friendly support team

(its-ri-team@manchester.ac.uk)

Networking & Security

- The iCSF/CSF has a private (campus only) IP address & firewall
 - Controls connections to and from the system
 - On campus
 - Connect from any PC/laptop with a wired connection or University wireless and Eduroam

No need for VPN but will need to authenticate using your 2FA device, e.g.

```
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
```

- Off campus: http://ri.itservices.manchester.ac.uk/wfh/
- University Global Protect VPN Gateway with 2FA
 - Must be running
 - Allows you to the CSF & iCSF
 - Can be some issues with some ISPs, see above WFH webpage
- Further documentation: http://ri.itservices.manchester.ac.uk/csf3/getting-started/connecting/



How do I Connect to the iCSF & CSF?

- Via the Remote Virtual Desktop Service (RVDS) focus of today's exercises.
- Directly via SSH using a Terminal or suitable Terminal Emulator such as e.g. MobaXterm, Putty - NOT RECOMMENDED for iCSF
- What operating system are you connecting from, Windows, Linux, OSX?
 - Windows users need to install a free terminal app called MobaXterm http://mobaxterm.mobatek.net/download-home-edition.html
 - Linux and Mac both have a terminal application by default
 - Mac depending on the version of OSX you may need to install X-Quartz in order to view GUI http://xquartz.macosforge.org/



RESEARCH VIRTUAL DESKTOP SERVICE



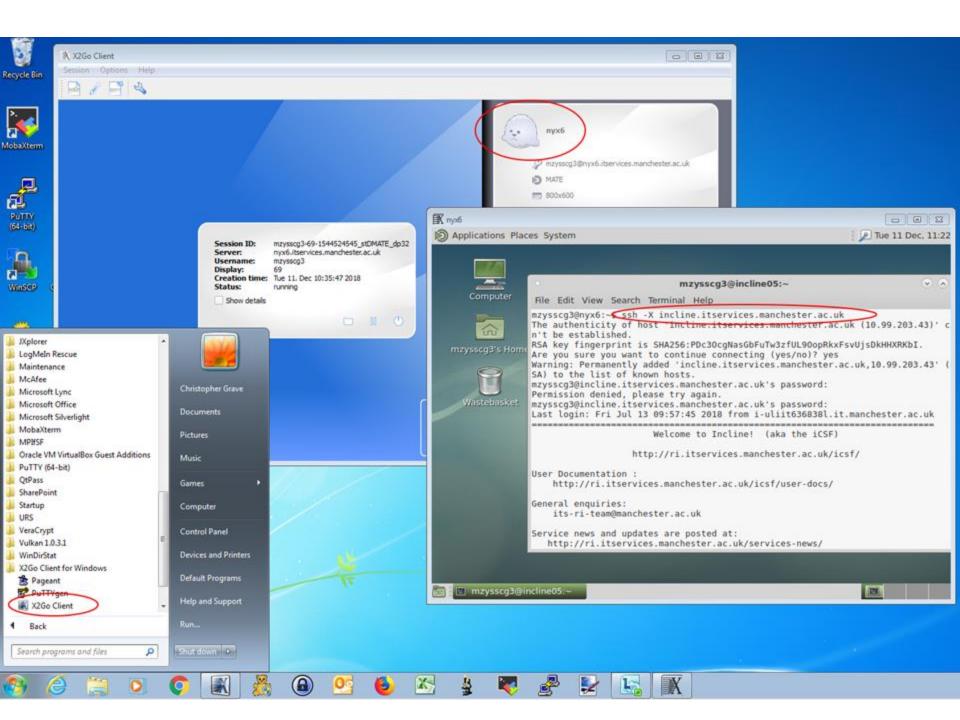
Research Virtual Desktop Service & X2Go

- Recommended method for connecting to the iCSF can be used to connect to other platforms within the CIR ecosystem
- Accessible from off/on-campus
- Requires a small piece of Software called X2Go (open-source) MAC users will also have to download Xquartz
- Setup Guide http://ri.itservices.manchester.ac.uk/virtual-desktop-service/x2go/
- PLEASE USE either nyx5, nyx6, nyx7 these provide the virtual desktop
- Login with central username & password



Benefits of Research Virtual Desktop Service

- Accessible from off/on-campus
- Provides a remote desktop enabling users to connect to the platforms
- Start a session at work power off your work PC at the end of the day, go home and reconnect to the same session continue from where you left off.
 - Connecting directly to CIR platforms without going via the RVDS does NOT provide the reconnect functionality e.g. Closing Mobaxterm which is directly connected to the iCSF will also log you out of iCSF and kill any running jobs. :(
- X2go by design uses compression to make GUI's run smoothly even over slow connections





RVDS Exercise

Connect to incline using the RVDS



Research Data Storage (RDS) & CIR Filesystems



Research Data Storage Service

- Also commonly known as isilon (the technology on which it relies)
- Data stored on RDS can be considered secure e.g. files can be recovered if accidently deleted
 - Automatically backed up once an hour (changes kept for 24 hours
 - Automatically backed up daily and copy kept for 35 days
- Does <u>not</u> provide Encryption at Rest therefore if handling restrictive/highly restrictive data please speak to us first.
- Can be accessed from your PC both on and off campus
- Groups can share files easily



Can I have some RDS

- Each funded Research group entitled to 8TB free of charge
 - Over 8TB costs money
- Must be requested by academic responsible for the group /project e.g. Pl
- All users' home directories reside on RDS
 - Project data should reside in dedicated Project folders not home
 - Projects should request further storage if required
- If requesting RDS
 - Research Data Management Research Council requirement that you have a Research Data Management plan – Outside scope of this presentation for more info
 - http://www.library.manchester.ac.uk/services-andsupport/staff/research/services/research-data-management/



Filesystems cont....

 After logging onto the iCSF/CSF you will be automatically placed into your home directory.

```
[username@incline31 ~]$ pwd (tells you where you are)
/mnt/hum01-home01/username
```

- Home space is shared by everyone in the group and space is limited. It is not designed for long term storage of data - use project space
- Only you can access your home directory. However sharing is possible using file permissions. Please contact Research IT regarding this.

Filesystems cont....

You can see how much space you are using with the command

```
$ du -sh
```

How big is that file? -

```
$ ls -lh filename
```

How much space is used/free overall?

```
$ df -h . (don't forget the.)
```



Uploading and Downloading Files

- Various ways this can be achieved. MobaXterm, WinSCP, RDS-SSH,
 SSHFS & command line utilities e.g. scp, rsync.
- How large are the files you need to transfer?
- What operating system are you using?
- Documentation can be found at the following location

http://ri.itservices.manchester.ac.uk/userdocs/file-transfer/



Documents - mzysscg3@incline.itservices.manchester.ac.uk - WinSCP

Queue -

· 🐸 🔽 | 4 · 4 · | 🔁 🔯 🎧 🗗 🐉

Parent directory

File folder

File folder

File folder

File folder

File

Text Document

Microsoft Word D...

Local Mark Files Commands Session Options Remote Help

mzysscg3@incline.itservices.manchester.ac.uk | New Session

Upload + 📝 Edit - 🗶 🚮 🕞 Properties ᢨ New • 📳 🖃 🔻

0 KB

Size Type

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My documents

Name

£ ..

Corel

test

MobaXterm

My Palettes

Outlook Files

email addresses.txt

0 B of 19.7 KB in 0 of 7

RDS Support Matrix.d...

C:\Users\mzysscg3\Documents\

WinSCP

Transfer Settings Default

Name

N R

scratch

out.txt

Changed

28/08/2018 15:38:3

07/06/2017 10:28:1

04/09/2017 12:11:0

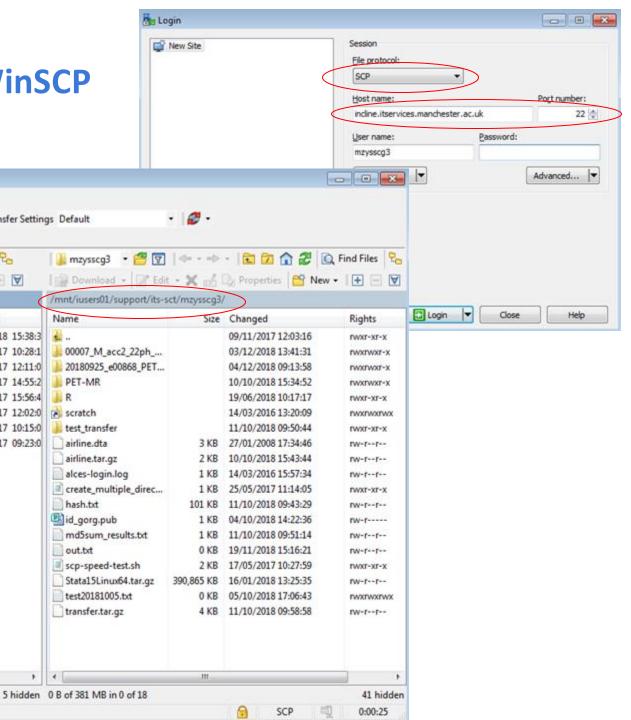
28/09/2017 14:55:2

07/07/2017 15:56:4

06/04/2017 12:02:0

20/09/2017 10:15:0

17/08/2017 09:23:0





Uploading & Downloading Files via the Command Line e.g. Linux or OSX

Upload a file from your computer to your iCSF home dir

scp file1.txt username@incline2000.itservices.manchester.ac.uk:
Secure Copy Protocol

No
destination
after the:
means "use
your home
directory

Download a file from your iCSF home dir to your computer

scp username@icsf.itservices.manchester.ac.uk:results.outy.

The . translates to "the current directory" on your computer

Upload a file from your computer to an CSF folder in home

scp file2.txt username@csf.itservices.manchester.ac.uk:mydata/

Download a file from your CSF folder in home to your computer

scp username@csf.itservices.manchester.ac.uk:mydata/results.out results.copy

Will rename the copied file to "results.copy"



File Transfer Exercise

- Upload a file to the iCSF using WinSCP
- Manipulate a file using Linux commands



USING THE ICSF

(logging in, setting up and running applications)



iCSF Resource

- The iCSF is a shared resource you will likely be sharing the compute nodes with other people
- The login node assesses the load on them before deciding which one to place you on.
- 7x Super-High-Memory nodes 32core 2TB (2000GB) of memory

ssh -X
centralusername@incline2000.itservices.manchester.ac.uk

Connecting to the iCSF

- Open X2go application and connect to to Virtual Desktop Service i.e. nyx 5,6,7.
 - From Menu Applications > System Tools > MATE Terminal
- In Terminal log into the iCSF
 - o ssh -X username@incline2000.itservices.manchester.ac.uk
 - Perform 2FA when prompted
- You will end up on a backend node e.g. incline31
- The node is indicated by the 2 digit number in the above example
 - O 31-37 = Super High Memory 2TB nodes



Starting Apps

- All software is controlled and loaded using module files
- List available software

```
module avail
```

Not all software will be available due to license restrictions

All software is loaded via modulefiles e.g.

```
module load apps/binapps/rstudio/0.98.1103-any-r module load apps/binapps/matlab/R2017a
```

List currently loaded modules i.e.

```
module list
```

Once module file is loaded open the software (the & lets you keep using the terminal)

```
rstudio & matlab &
```

- Voila! GUI/window should open
- More information on available software, and module files please follow this link



iCSF Exercise

- Load and Run Rstudio
- Logout of the iCSF



Using the CSF

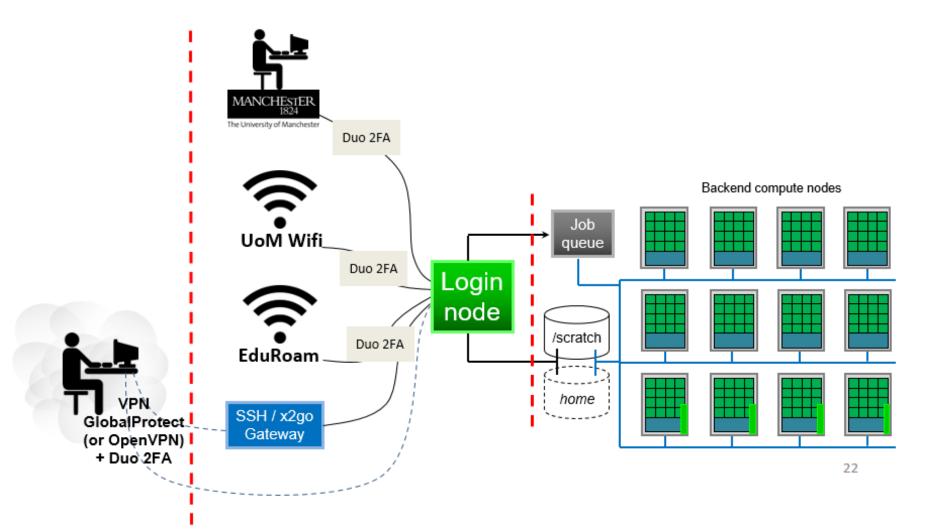
What we'll be using today - the CSF

- CSF3 current config:
 - A large Linux cluster system
 - 14,016 CPU cores (Intel "Xeon" or AMD "EPYC" CPUs)
 - 152 Nvidia GPUs (68 x v100, 72 x A100, 12 x L40s)
 - Got big datasets to process? Can run large-memory jobs
 - (we'll cover all of these details throughout the course)

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

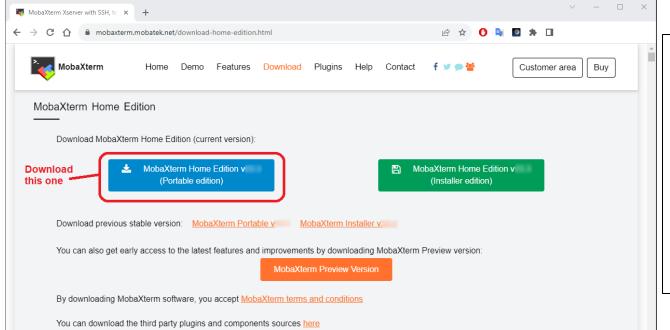


How can I access the CSF



Connect to CSF from Windows

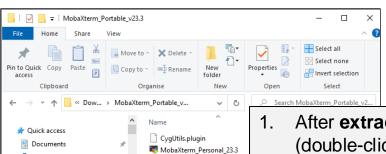
- Job queue Pacratch
- Access the CSF from a PC / laptop using an SSH (Secure Shell) app
 - Sometimes called a "terminal".
 - There's no web-site or other fancy GUI on the CSF use the "command-line".
- Windows users need to install a free terminal app called MobaXterm
- https://mobaxterm.mobatek.net/download-home-edition.html
 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.

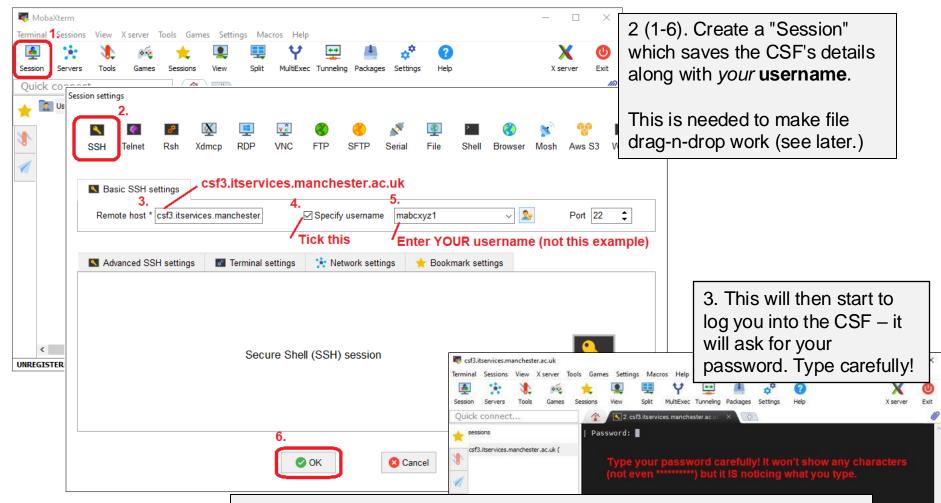


Downloads

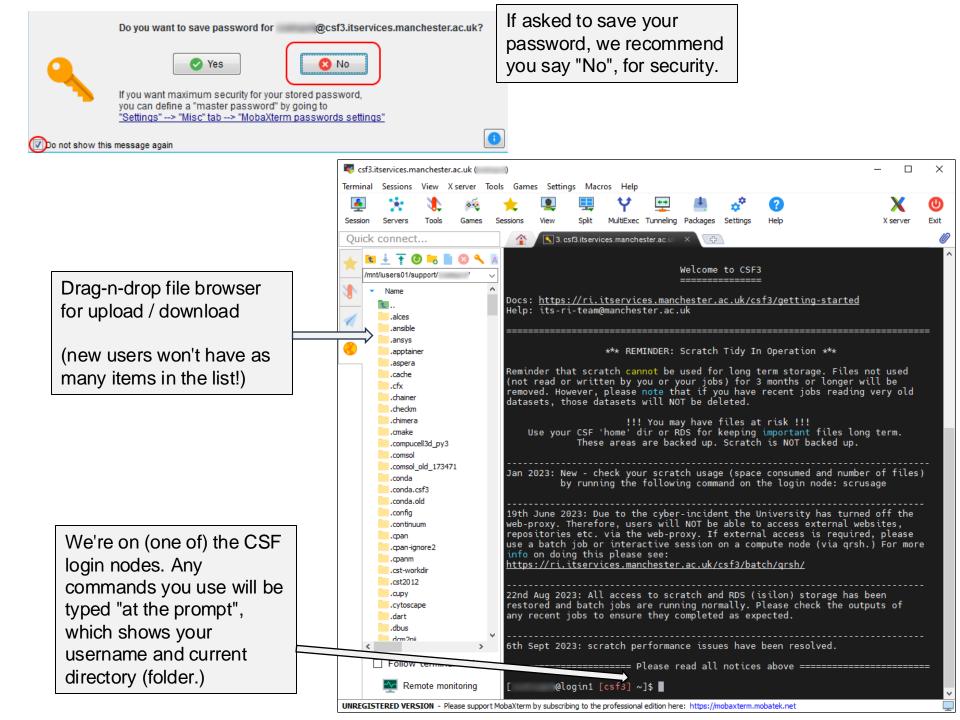
MobaXterm "Session"

(username saved in the session setup)

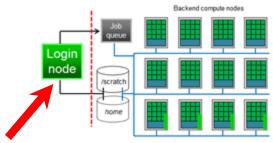
 After extracting the .zip file, start MobaXterm_Personal_xy.z (double-click on the icon)



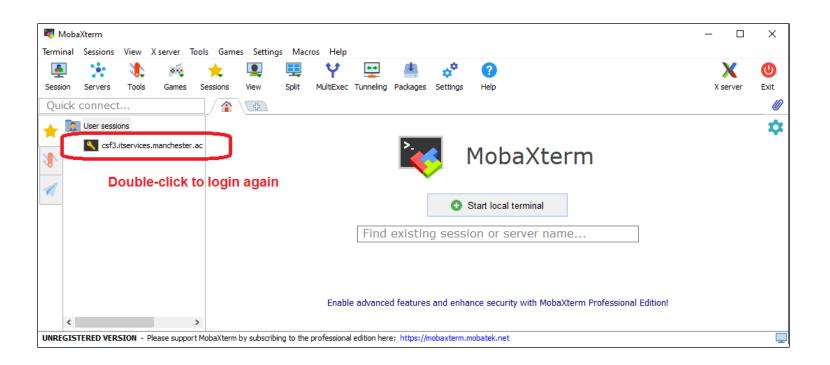
4. See slide about 2FA - you may be asked for DUO after your password



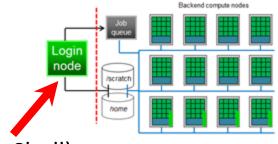
Next time you want to login to CSF from Windows



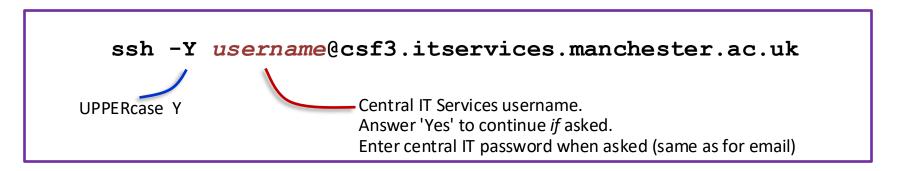
- Just double-click the csf3 "session" in the list of "User sessions"
- The CSF details are saved in the "session"
- (this also makes the file browser work, for drag-n-drop file transfers.)



Connect to CSF from a Mac

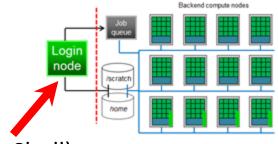


- Access the CSF from a PC / laptop using an SSH (Secure Shell) app, eg a 'terminal'.
 - There's no web-site or other fancy GUI on the CSF command-line for now.
- Mac users have a terminal application by default
 - You will need to install X-Quartz first
 http://xquartz.macosforge.org/ (install then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

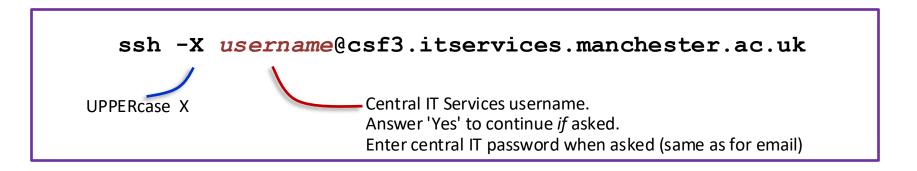


Finished using CSF? Log out with: logout or exit

Connect to CSF from Linux

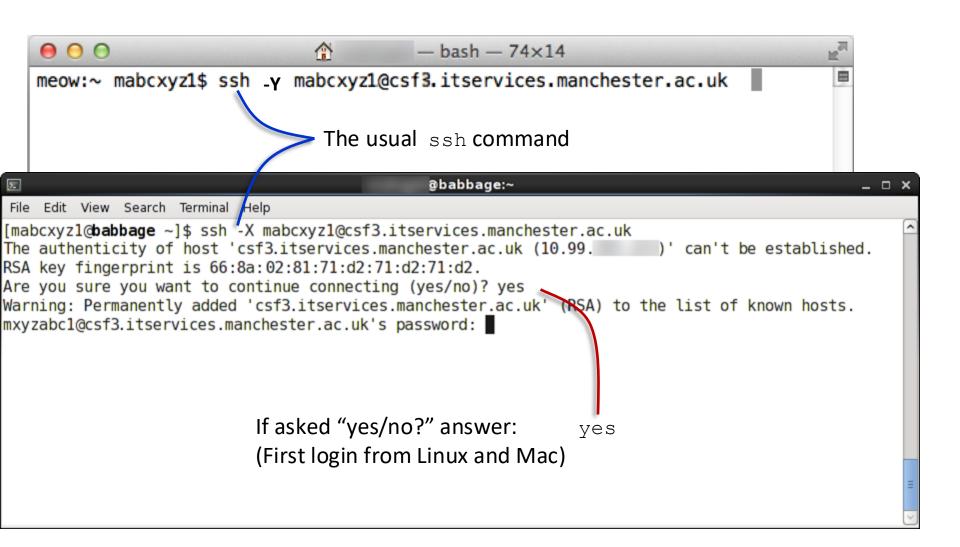


- Access the CSF from a PC / laptop using an SSH (Secure Shell) app, eg a 'terminal'.
 - There's no web-site or other fancy GUI on the CSF command-line for now.
- **Linux** users have a *terminal* application by default
 - Start a Terminal and type the following command:



Finished using CSF? Log out with: logout or exit

Linux / Mac Terminals



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):
 - http://www.ee.surrey.ac.uk/Teaching/Unix

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!



CSF Scratch filesystem

- An extra and very large filesystem available on the CSF
- Shared by all CSF users
- Run jobs here it is fast storage, sometimes home can fill up causing jobs to fail
- You should have a 'symlink' to it in your home directory (if not ask us to create one)

cd ~/scratch

- Not backed up
 - When job finishes move important files to home or other RDS area
 - Not considered safe for long term storage hardware failure could cause loss
 - Files that have not been accessed in the past 3 months will be deleted



A simple batch job

- We recommend you have a directory for every job
 All files relevant to the job need to be in the folder
 Be tidy, it makes managing results easier
- We need a jobscript Create it on the CSF using this command gedit &

this avoids a problem with linux/batch not being able to read text files created on windows.

Like iCSF software is accessed by modulefiles

eg: module load apps/R/3.4.2

Unlike iCSF, most of the time we only use module load in a jobscript

A simple Jobscript – Serial (1 core)



#! 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.

myjob.txt

#!/bin/bash --login

#\$ -cwd

#\$ -N myjob •

#\$ -1 resource

Let's do work

date

hostname

sleep 120

date

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

To give the job to the system:

```
qsub jobscript
```

Replacing 'jobscript' with the name of the text file you have created.

To see the job in the batch system

```
qstat
```

- You should see the job either queued (qw) or (r)
- Your job will have a unique numerical JobID
- If you can't see the job it has finished. :)



Where did my results go?

- Two possibilities
 - A file called jobname.oJobID
 Where jobname is the name of your submission script
 JobID is the number of your job
 - An output file specific to your job
- If your job had a problem or failed it may be indicated in one of the above or jobname.eJobID
- Various options to view the file:

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

 Depending on the application and type of files produced you might want to use the iCSF to view them, postprocess, visulaise, etc.



Jobs do not always start quickly

- Your job will wait until the resources requested become available.
- You can log off, switch off your PC and your job will stay on the CSF. Log in again to collect the results.
- You can submit more than one job at once, more than one may run at the sametime (good to have different folder for each)



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

- qstat reports your job as Eqw
 - CSF tried to start it, but something went wrong
 - O Usually no output file so to indicate what happened use qstat -j JobID | grep error
- Error can be cryptic
- Most common causes
 - Missing directory
 - You created the jobscript on windows
 - Unusual characters or spaces in file and directory names
 - No disk space on the filesystem highly recommend you use scratch to avoid this
- To delete a problem job or one you simply no longer want:
 - o qdel jobID



Parallel Jobs on the CSF

- CSF excels at doing this
- Each compute node has multiple processors, each of which has multiple cores, each of which can run a single task
- A parallel job uses more than one of these cores
 - Specify how many cores are required
 - Software splits your job into chunks and runs them all at the sametime
 - Usually means jobs run faster
 - Allows access to more memory
 - At the end pulls everything together into your results
- Requires a parallel environment (pe) -

https://ri.itservices.manchester.ac.uk/csf3/batch/parallel-jobs/#Intel_parallel_environments

There's another way to run in parallel! High Throughput Computing (HTC)

- Lots of independent computations
 - Processing lots of data files (e.g., image files)
 - Running the same simulation many times over with different parameters ("parameter sweeps")
- Run many copies of your program
 - Programs may be serial (single core) but running lots of them at once. They don't communicate.

Parallel Jobscript – Multi-core (single-node)

```
smp.pe is the parallel environment
 #! and #$ see
                           myparajob.txt
                                                 name. This one means: app will use a
 serial jobscript
                                                 single compute node (up to 32 cores.)
 earlier.
                 #!/bin/bash --logi
                                                             4 is the number of cores
                 #$ -cwd
-pe indicates.
                                                             we want to reserve in the
                 #$ -pe smp.pe 4
we'll run a
                                                             system. Each PE has a
parallel job in a
                                                             maximum allowed.
                 # Set up to use a chemistry app
particular parallel
                 module load apps/binapps/anaconda2/2021.11
environment.
                 # Inform app how many cores to use
                 export OMP_NUM_THREADS=$NSLOTS
# indicates line is
a comment so
                 # This job runs a python script
does nothing.
                 python3 name_of_python_script.py
                                                                    Key concept!
```

Any commands we run in our job.
They will execute on a backend
node that has required number of
cores free. mdrun d is gromacs.

Must *somehow* inform the app how many cores we *reserved*. **\$NSLOTS** is automatically set to number **(4)** given on **-pe** line. **Our app** wants **OMP_NUM_THREADS** environment variable setting. **Your app** might use a different method!



What else can the CSF do?

- Access more cores across multiple-nodes. Up to 1024 via the HPC
 Pool
- Job Arrays Multiple Similar Jobs https://ri.itservices.manchester.ac.uk/csf3/batch/job-arrays/
- High Memory jobs, access up to 4TB ideal for processing large
 DataSets -https://ri.itservices.manchester.ac.uk/csf3/batch/high-memory-jobs/
- Over 150 GPUs, ideal for machine learning type workloads
- One day CSF course covers the above



CSF Exercise 1:

- Connect to CSF via RVDS
- Run a serial & parallel python jobs
- View the results on the CSF



Other Research IT Infrastructure Provided Services



HPC Pool

Dedicated pool for "true" HPC jobs

- 4096 cores
- Minimum 128-core job size, maximum 1024
- Frontend shared with CSF3
- Lightweight application process must be made by PI
- Currently free
- http://ri.itservices.manchester.ac.uk/csf3/hpc-pool



Research Virtual Machine Service (RVMS)

- Another component of the CIR ecosystem
- Centrally-hosted virtual machines (VMs) are provided for University research staff and PG students
- RVMS is suitable for VMs which are expected to generate large amounts of network traffic, may be moderately CPU-intensive or require tight integration with the computational platforms (CSF, iCSF, etc.) or the Research Data Service (RDS).
- Administrative control of VMs provided maybe handed over to research groups.
- More info http://ri.itservices.manchester.ac.uk/rvms/



Condor

- Another component of CIR ecosystem
- Utilises 100s of Desktop PCs which are idle on campus
 - Hence typically only available overnight, weekends, holidays.
- Suitable for jobs which require the same computation i.e.
 repeated many times over different data sets/ranges. Also known as High Throughput Computing.
- Burst into the Cloud using Amazon Web Services
- More Information can be found by following this link

http://ri.itservices.manchester.ac.uk/htccondor/



Research IT

- Thank you for attending!
- Course Materials -http://ri.itservices.manchester.ac.uk/course/icsf-csf/
- Feedback PLEASE COMPLETE https://goo.gl/forms/zfZyTLw4DDaySnCF3
- Support/Queries <u>-https://ri.itservices.manchester.ac.uk/hpc-help</u>

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