HPC Workshop for Beginners

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You can find the slides in www.hannokase.com/HPC/

How do I get the HPC account?

To access the HPC cluster, you need to request an account via **EUI Helpdesk**.

For this workshop, we have created you **test accounts**:

User name: cmatlab01-cmatlab16

Password:

How to connect to the HPC cluster?

You can access the HPC cluster through Secure Shell Protocol (SSH), using any SSH client.

On Linux, Windows 10, and Mac a SSH client is preinstalled and you can use it through command line (Terminal, Command Prompt, PowerShell).

Command for using SSH: ssh username@host

ssh hkase@hpceui.iue.private

hkase@hpceui:~		×
C:\Users\kaseh>ssh hkase@hpceui.iue.private hkase@hpceui.iue.private's password: Last login: Mon Oct 28 18:16:17 2019 from 192.168.3.104		
######################################		
The HPC Cluster is newly available for simulations since the upgrade of the Workload Management Systems has been implemented.		
Moreover, the HPC Cluster now features two new nodes termed as matlabcl8 and matlabcl9 accessible via qmw36 queue.		

[hkase@hpceui ~]\$ _		

Are you tired of typing your password?

To avoid typing your password every time you enter the HPC cluster you can add your **public** SSH key to ~/.ssh/authorized keys file on the HPC cluster.

On Windows 10 you can find your public SSH key in:

```
C:\Users\<username>\.ssh\id rsa.pub
```

This is the default location, it might be somewhere else. If you don't find it, you can generate one by ssh-keygen command. Choose the location (default location is usually fine), choose a password (or leave it empty).

Some essential Linux commands

Command	What does it do?
man [COMMAND]	Manual page for [COMMAND]
passwd	Change your password
pwd	Current working directory
cd [FOLDER]	Change directory to [FOLDER]
cd	Move to parent directory
cd ~	Change directory to home directory
ls	List files in the folder
mv [SOURCE] [DESTINATION]	Move or rename files
cp [SOURCE] [DESTINATION]	Copy files
rm [FILE]	Delete file
mkdir [FOLDER]	Make a directory named [FOLDER]
rm -r [FOLDER]	Recursively delete the folder and all files in it
nano [FILE]	Open text editor (nano)

How to transfer files to the HPC cluster

- Command Line
 - SFTP
 - SCP
- Graphical User Interface
 - Using a SFTP client (<u>FileZilla</u>, <u>WinSCP</u>)
 - Map a network drive*

*Easy on Linux, but a bit tricky on Windows 10 and Mac

- Windows 10: install WinFsp and SSHFS-Win
- Mac: this might be helpful

Transfering files - Command Line: SFTP

Interactive command, that opens a connection for submiting several commands for transfering files.

Command for opening SFTP connection: sftp username@hpceui.iue.private

sftp hkase@hpceui.iue.private

SFTP interactive commands

Command	What does it do?
cd	Change the active directory on REMOTE machine
lcd	Change the active directory on LOCAL machine
ls	List files in the active folder on REMOTE machine
lls	List files in the active folder on LOCAL machine
rm	Remove files on REMOTE machine (there is no lrm)
mv	Move files on REMOTE machine (there is no lmv)
ср	Copy files on REMOTE machine (there is no lcp)
put [FILE]	Put a file from the LOCAL machine to the REMOTE machine
get [FILE]	Get a file from the REMOTE machine to the LOCAL machine

Close the connection by typing <code>exit</code> or <code>bye</code> or pressing CTRL+C.

SFTP Example

```
sftp hkase@hpceui.iue.private
```

Let's try to transfer file1.txt file from local machine to the remote machine.

```
put file1.txt
```

The file file1.txt from the active directory on our local machine got transferred to the active directory on the remote machine.

Transfering files - Command Line: SCP

Non-interactive command. Takes only one command and then exits.

Command for using SCP: SCP [SOURCE] username@hpceui.iue.private:[TARGET]

```
scp file1.txt file2.txt hkase@hpceui.iue.private:~/demo
```

For faster file transfer, we can use compression -C.

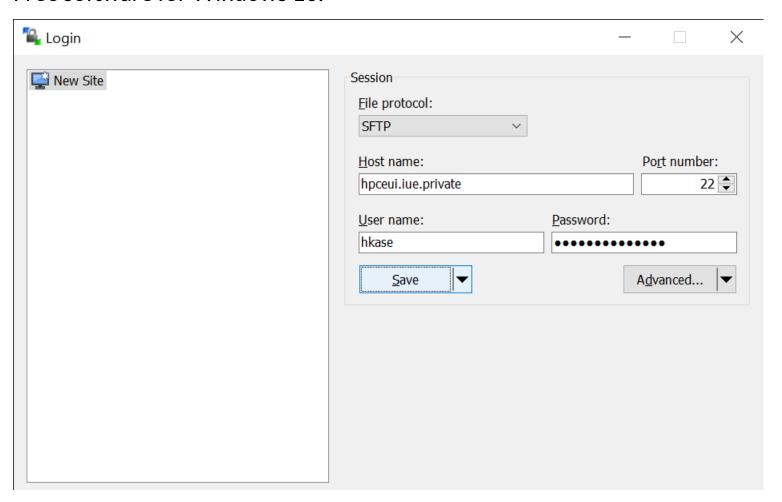
```
scp -C file1.txt hkase@hpceui.iue.private:~/demo/file_rename.txt
```

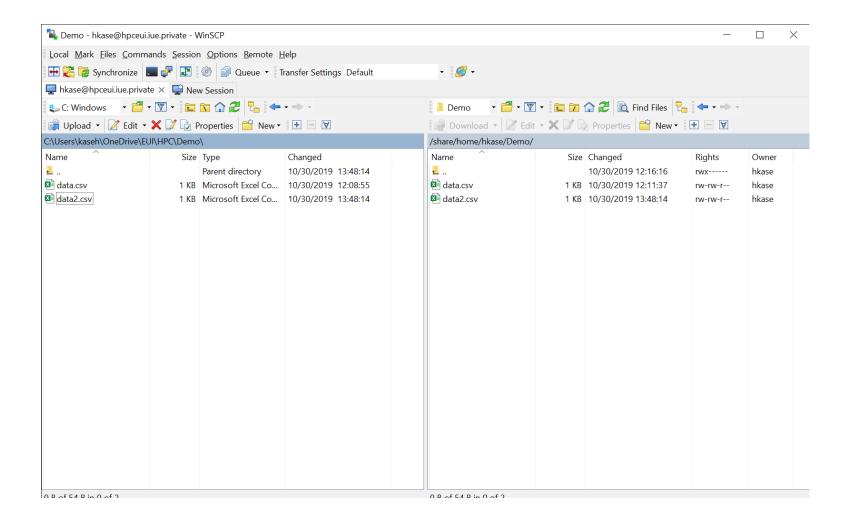
Add -r flag to copy the whole folder.

```
scp -r folder hkase@hpceui.iue.private:~/new_folder
```

Transfering files - GUI: WinSCP

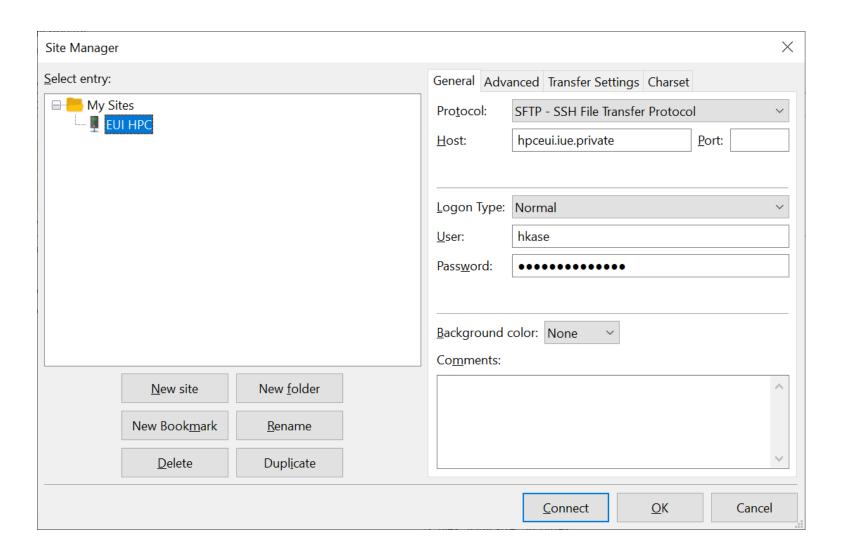
Free software for Windows 10.

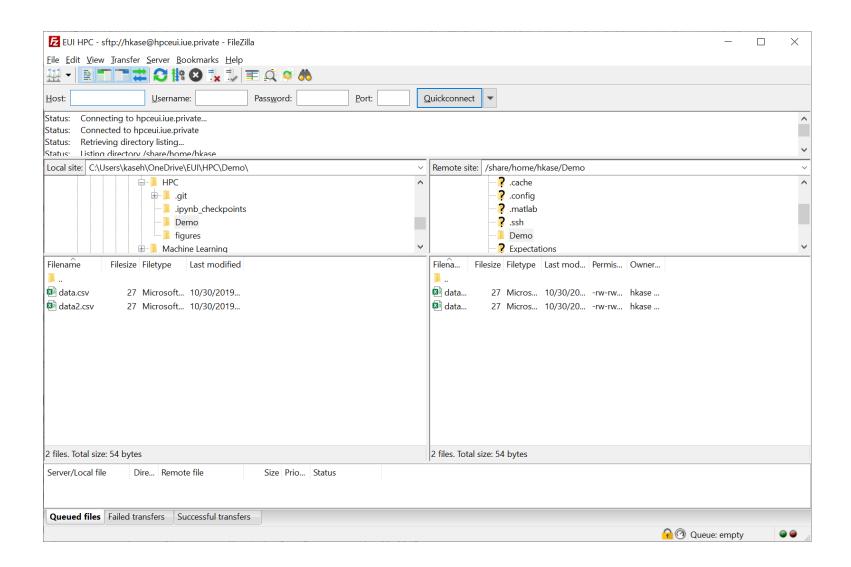




Transfering files - GUI: FileZilla

Free software available for Linux, Mac, Windows 10. Careful when installing it. You probably don't want the included Avast Antivirus.





Transfering files - Map a network drive

Windows 10

- Install WinFsp and SSHFS-Win. Look for latest release tag and .msi files.
- Map a network drive sshfs//<username>@hpceui.iue.private

Linux - Ubuntu 18.04

• Add remote location sftp://<username>@hpceui.iue.private/

Mac

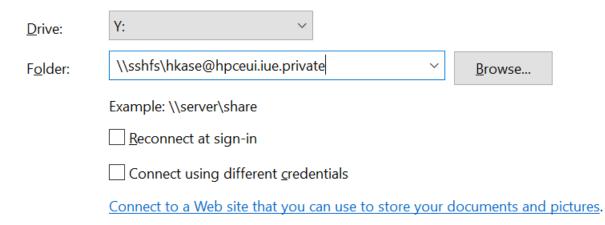
• This might be helpful

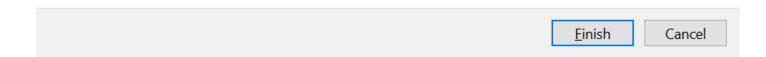




What network folder would you like to map?

Specify the drive letter for the connection and the folder that you want to connect to:





Queues in the HPC cluster

At the heart of the cluster is a workload management system (PBS Professional); a program that allocates computational resources.

You use the cluster by first asking for resources. Depending on your computational needs, you will choose from which **queue** to ask the resources from.

The EUI HPC cluster is designed such that there is a direct mapping from queues to computing nodes (servers). All the resources you request are allocated within one node.

Queue	Slots	Nodes	CPU cores	Memory	GPU	Walltime
qwe36	4	matlabcl6-9	36	128GB	No	12h
qwe24	2	matlabcl4-5	24	128GB	No	12h
qwe12	1	matlabcl3	12	32GB	Yes	72h
qwe36long	?	matlabcl6-9	36	128GB	No	168h
qwe24long	?	matlabcl4-5	24	128GB	No	168h

Which queue should I choose?

qstat -Q or qstat -q to find out what kind of queues are there.

qstat or qstat -n1 to monitor the queues.

pbsnodes -av to get a detailed overview of all the nodes.

How to submit a batch job to the HPC cluster?

The most common use of the cluster is to submit a **batch job**. For this we will need to specify for the workload manager:

- 1. what kind of resources we need and ...
- 2. what program we would like to run.

We do it all by creating a **PBS script** (like a bash script), that contains some special instructions for the workload manager.

Then we submit the job into the queue using <code>qsub [pbs_script]</code> command. When there are available resources, it will get executed.

Script for running MatLab code (CPU)

```
#!/bin/bash
# 1. Job name
#PBS -N MATLAB
# 2. Request resources
# set the number of nodes and processes per node
#PBS - q qmw12
# use submission environment
\#PBS - V
# 3. Run the program
# start the job from the directory it was submitted
cd $PBS O WORKDIR
# set some environment variables for MatLab
source /apps/Matlab var R2019a.sh
# Run your the matlab code
matlab -batch "R2019a_mod_paralleldemo_parfor_bench" >& matlab.log
```

Script for running MatLab code (GPU)

```
#!/bin/bash
# 1. Job name
#PBS -N MATLAB
# 2. Request resources
# set the number of nodes and processes per node
#PBS - q qmw12
# use submission environment
\#PBS - V
# 3. Run the program
# start the job from the directory it was submitted
cd $PBS O WORKDIR
# set some environment variables for CUDA and MatLab
source /apps/cuda-10.1.sh
source /apps/Matlab var R2019a.sh
# Run your the matlab code
matlab -batch "paralleldemo_gpu_backslash" >& matlab_gpu.log
```

Script for running Julia code

```
#!/bin/bash
# 1. Job name
#PBS -N Julia
# 2. Resource request
#PBS -q qmw12 -1 walltime=12:00:00, select=1:ncpus=12
# use submission environment
\#PBS - V
# 3. Program to run
# start the job from the directory it was submitted
cd $PBS O WORKDIR
# set some environment variables for Julia
source /apps/Gnu var 4.9.4.sh
export PATH=/opt/pkgs/julia-1.1.0/bin:$PATH
# Run your Julia code
export JULIA NUM THREADS=4
julia -03 -- spectralnorm.julia 5500 >& spectralnorm.julia.log
```

Script for running FORTRAN code (OpenMP)

```
#!/bin/bash
# set the number of nodes and processes per node
#PBS -q qmw12 -1 select=1:ncpus=12
# set name of job
#PBS -N Fortran
# OMP Environment Variable #
export OMP NUM THREADS=12
export OMP THREAD LIMIT=12
# use submission environment.
\#PBS - V
# start job from the directory it was submitted
cd $PBS O WORKDIR
# compile the Fortran Source....
echo "Running gfortran on `hostname`: gfortran -fopenmp matrix mult.f -o matri
x mult.exe"
gfortran -fopenmp matrix mult.f -o matrix mult.exe
# run my application over a compute node within the PBS ring (pbsnodes -av)
echo "Running line on `hostname`: ./matrix mult.exe > 12 cores output file "
./matrix mult.exe > 12 cores output file
```

I would like to run R/STATA/Python...

You can find many other PBS script samples on the cluster in /share/apps

Monitoring your job

Command	What does it do?
qstat	Shows the status of all the jobs
qstat -n1	Shows the status of all the jobs and their nodes
qstat -u [username]	Shows the status of the jobs belonging to [username]
qstat -f [jobid]	Shows the detailed information about [jobid]
qdel [jobid]	Delete the [jobid]
tail -f [output_file.log]	Prints the output file in the terminal as it is being written

How is my code doing?

If you are not quite sure that your code is running nicely in parallel, you can log into the compute node and check the task manager htop.

- 1. Find the node where your job is running using qstat -u [username] -n1.
- 2. Log into the node ssh [node name], for example ssh matlabcl3.
- 3. Run htop to open the task manager
- 4. Close the task manager by pressing F11
- 5. Close the SSH connection by typing exit

How to use the cluster interactively?

Using cluster interactively can be useful for developing your code or debugging.

Add -I to the reqular qsub command. The following command will start an interactive job with the parameters specified in the pbs script.

```
qsub -I pbs_script
```

You can start an interactive session without a PBS script.

```
qsub -I -N MyInteractiveJob -q qmw12 -l select=1:ncpus=12:host=matlabcl3
```

Thank you!

E-mail us if you have questions: hpc.support@eui.eu
Request an account via EUI Helpdesk.