Using the Computer Science Cluster

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Computer Science (CS) High-Performance Computing (HPC) Cluster

- CS cluster has over 5200 nodes.
- Designed to run large scale computing jobs in batch mode (non-interactive mode).
 - Limited graphics-based interactive computing services.
- Users should set up their code to run in batch mode.
- Users who primarily need graphics-based interactive computing should use Economics Department cluster, ISD services, or desktop computers.
- Users will benefit from learning some basix unix commands.

More information

- CS cluster webpage: http://hpc.cs.ucl.ac.uk/how_to_login/.
 - Get username and password from IT support,
- Econ GitHub page: https://github.com/UCL/ECON-CLUSTER
 - Contact Fatima or Andrew to be added to the UCL user group.
- If you need help,
 - CS support: cluster-support@cs.ucl.ac.uk
 - ECON IT support: economics.it@ucl.ac.uk

Obtain account

- Fill out and submit application form (see Fatima or Andrew).
- Wait 5-7 days and go to CS Cluster office: (4.20 Malet Place Engineering Building).
- They will set up two accounts
 - OS departmental account (for remote access).
 - ② CS Cluster account (to use cluster).
- By default, both accounts have same username and password.

Connect to cluster

There are three ways to connect to the cluster:

- Connect using ThinLinc (graphical interface).
- 2 Connect using ssh (command line interface).
- Onnect using ftp (data transfer).

Connection option 1: ThinLinc

Connecting using ThinLinc is a two-step process:

- Connect to a CSRW (Computer Science Remote Worker) using your CS Department Account username and password.
 - Instructions to download Thinling.
- 2 Then connect to one of the cluster login nodes ("vic" or "wise") using your **CS Cluster Account** username and password.
 - Open terminal inside CSRW window.
 - Then type one of the following
 - ssh vic
 - ssh uctpXXX@vic
 - Senter your cluster password.

Connection option 2: ssh

- For remote access (i.e. when not connected to UCL network), connecting is a two step process:
 - Connect to "tails", "storm" or "jet" using your CS Department Account username and password.
 - Connect to the cluster using your CS Cluster Account username and password.
- For access from within the UCL network (e.g. a desktop in Drayton House or the Econ HPC), skip step 1.

Option 2 Step 1: ssh to "tails", "jet" or "storm"

- Open an ssh client (e.g. putty) or a terminal.
- Type
 - ssh -X uctpXXX@tails.cs.ucl.ac.uk
- Enter your CS dept. password.
- Note:
 - The option '-X' allows graphics to be forwarded from the CS cluster to your computer.
 - The username "uctpXXX" is that given when you are assigned your CS department account.
 - The text after the '@' is the address of the server.

Option 2 Step 2: ssh to cluster

- Logon to the CS cluster using ssh.
- Type
 - ssh -X uctpXXX@vic
- Enter your CS cluster password.

Connection option 3: sftp (for file transfer)

- To transfer files to cluster storage use sftp instead of ssh.
 - Open a terminal or an FTP client.
 - 2 Type
 - sftp uctpXXX@tails.cs.ucl.ac.uk
 - 3 Enter your CS department password.
- From tails, you can directly access the CS data storage.
- For example, if you have previously requested CS to create a storage directory named "PROJECT_X", you can access this by typing:
 - cd /slash/economics/research/PROJECT_X
- We discuss requesting CS data storage below.

Typical workflow

- Logon to cluster.
- Transfer data and/or files to cluster (using ftp, email, or some version control software like git).
- Edit files or code.
- Write a script to submit your job to SGE (Sun Grid Engine).
 - SGE is the scheduler that manages allocation of jobs to nodes on the cluster.
- Submit your job.
- Monitor job progress if necessary.
- Download results to your local computer (using ftp or email).

SGE sessions: two types

- There are two types of SGE sessions
 - Interactive sessions
 - Start session with "qrsh".
 - Interact with compute node directly on command line.
 - Good for developing code and testing things.
 - Non-interactive sessions
 - Start session with "qsub".
 - Job runs in background.
 - Good for running large scale, long running, or multiple jobs.
 - Cluster is optimised for this type of job.
- SGE jobs are command line only jobs.
- If you need to do GUI based interactive work, it is best to use an alternative cluster.

Interactive SGE sessions

- Use the "qrsh" command and specify running time and memory.
- For example, at the command line, type:
 - qrsh -I h_vmem=1.9G,tmem=1.9G,h_rt=8:0:0
- This command starts the session and requests 1.9 GB memory and 8 hours running time.
 - 'qrsh' is the login command for an interactive session.
 - '-I' is a flag for resource requests for the interactive session.
- Resource options listed after the '-I' flag are:
 - 'h_vmem=XG,tmem=XG' requests X Gb of memory
 - 'h_rt= H:M:S' requests that the session run for 'H' hours, 'M' minutes,
 'S' seconds
- For further command line options for 'qrsh' type: 'man qrsh'

User tips

- After starting SGE session, you need to load and open the software you require. See below for details.
- It may take a short while (1 5 minutes) to be allocated a node.
- To increase chances for quick allocation, request as little memory as necessary,
- For a small job, 2G is likely to be sufficient. For Matlab, request at least 4G.
- If you need a lot of memory (i.e. X > 2G), omit the 'h_rt' option from your 'qrsh' command.
- For example, to request a 14G session type:
 - qrsh -l h_vmem=14G,tmem=14G

Submitting batch jobs

To run a batch job:

- First, write a script (e.g. a text file names 'job1.sh') detailing what resources to request from SGE and what commands/programs to run.
- Then, submit the script using the command 'qsub'.
- For example, first create the file 'job1.sh' and then type the command
 qsub job1.sh
- Instructions for writing a script: job script instructions.

Some example command options to include in script

Request memory and running time:

```
#$ -1 h_rt=1:10:35
#$ -1 tmem=1.9G,h_vmem=1.9G
```

Each line containing SGE flags starts with '#\$'.

Requesting a parallel environment

- To run a job using more than one worker, you need either to specify a parallel environment or submit multiple jobs using "tasks".
 - We will discuss using "tasks" later.
- To specify the parallel environment, add the following lines to your shell script:

```
#$ pe [pe_option] [NumWorkers]
#$ -R y #
```

- replace [pe_option] option from next slide
- Replace [NumWorkers] with an integer.
- Second line reserves resources (needed when using multiple nodes).
- In most cases the best pe_option will be one of 'orte', 'smp', or 'matlabpe2014b'

Parallel environment options

- Options for parallel environments are
 - 'smp': single node with multiple workers
 - 'matlabpe2014b': parallel environment specific to Matlab. CS only support b release each year. Matlab 2015b is yet to be supported.
 - 'mpi': Old MPI interface
 - 'mpich': New MPI interface
 - 'orte': Distributed computing across nodes, but tries to cluster processes on nodes
 - 'para': Distributed computing across nodes, no clustering of processes
- Default option is 'orte'.

Examples

- Example 1: Use a single node with 4 workers
 - #\$ pe smp 4
- Example 2: Use a single node with 16 workers
 - #\$ pe smp 16
 - This example may "hang" in the queue for a long time; the scheduler will have to wait until a node with 16 cores is free.
- Example 3: 20 workers clustered on a few nodes

```
#$ pe orte 20
#$ -R y
```

Checking job status or deleting jobs

- For more details see Econ wiki.
- You can use the command 'qstat' or 'qstat-rn'.
- Note: 'qstat-rn' is only available if you have loaded the module 'econutils' by typing
 - module load econutils
- To delete a job with id 123456, type qdel 123456

Software available

- To find full details of software available on cluster go to cluster webpage or contact cluster support.
- The main software packages for economics are most easily used by using a module environment. See module environments.
- To see list of software currently available type

module avail

Software available

Current list is

dot
econutils
gcc/5.2.0
gcc/6.2.1
git/2.8.3
intel/composer/2013.1.117
intel/composer/2015.1.133
julia/0.4.5
knitro/10.0.1-z
knitro/10.1.2-z

```
matlab/r2012b
matlab/r2013b
matlab/r2014a
matlab/r2014b
matlab/r2015b
module-info
modules
nag/fll6i25dc
nag/mbl6a23dml
nag/mbl6a24dnl
```

openmpi/gcc/1.10.0 openmpit/gcc/1.8.1 openmpi/intel/1.10.0 openmpi/intel/1.8.1 R/3.4.2 stata/14 totalview/8.15.7-6 use.own

Loading modules

- Each software package depends on a large number of files.
- Loading the module file sets "environment" variables on the system so that the software will run correctly.
- To use modules, they must be loaded on every compute node on which you are running.
 - Interactive sessions: first start the interactive session using 'qrsh', then load the modules you need.
 - Non-interactive sessions: include the 'module load' commands in your batch job script or create a '.bash_rc' file to automatically load them at startup.

Examples

MATLAB

```
module load gcc/5.2.0 nag/mbl6a24dnl matlab/r2015b module load gcc/5.2.0 nag/mbl6a24dnl matlab/r2014b module load gcc/5.2.0 nag/mbl6a24dnl matlab/r2014a module load gcc/5.2.0 nag/mbl6a24dnl matlab/r2013b module load nag/mbl6a23dml matlab/r2012b
```

Stata

module load stata

R

module load gcc/5.2.0 R/3.4.2

Examples

Fortran

```
module load gcc/5.2.0 nag/fll6i25dc
module load intel/composer/2015.1.133
module load openmpi/intel/1.10.0 totalview
```

Julia

module load git/2.8.3 julia

- Note: Julia users should speak to cluster support about getting setup on the CS cluster. Amongst other things, there are file limit, version control, library paths, and non-trivial batch-mode issues to address before job processing can begin.
- KNITRO

module load knitro/10.1.2-z



Running software

- Interactive sessions (command line versions)
 - Type the relevant command: 'matlab -nodisplay', 'stata-mp', 'R', 'ifort'.
- Non-interactive sessions
 - Add the relevant command to your job script: 'matlab -nodisplay', 'stata-mp', 'R', 'ifort'.
 - See examples below.
- Interactive GUI sessions
 - First logon to 'jake' or 'elwood' using ssh with the '-X' option for x-forwarding of graphics
 ssh -X uctpXXX@jake
 - Load the relevant modules.
 - Start your program using 'matlab', 'xstata-mp', 'R', 'ifort'
 - Note: There are only two servers with x-forwarding enabled. You do not need to use SGE to access these servers. They can only be used for small scale temporary jobs.

Disk storage

- All accounts have a limited amount of 'local space' for storing working files. This space is limited (a few gigabytes), is not backed up, and should not be used to store important outputs or large datasets.
- Large scale storage (project store) with back ups is available on the CS SAN (storage area network).
- Storage can be made accessible to individuals or to groups.
- To request storage for a project named 'PROJECT_X', fill in the online storage request form.
- Space will then be allocated to you in the location '/SAN/economics/PROJECT_X' where 'PROJECT_X' is the name of your project.

Transferring Files to and from the CS HPC

- Use sftp.
- Use email.
- The SAN can be accessed directly from a CSRW or from 'tails' at the location '/slash/economics/research/PROJECT_X'

Cluster rules/best practice

- Never run jobs on head nodes (wise, vic)
- Store important data and/or large data on /SAN/economics/...
- Avoid high frequency reading/writing of large datafiles to/from disk
- If you need support, send email to cluster support, go visit them in person.
- Update wiki page with tips for colleagues about how to solve common problems.
- See cluster etiquette.

Memory management

- Most nodes have between 2 and 8GB per core. The physical machines vary from 4GB to 1TB.
- If you want to run a job requiring more than 47.2GB please contact request@cs.ucl.ac.uk.
- For the best job throughput please request less than 1.9G. That is:
- There are only a few machines with more than 64GB of memory. If you request more than 8G you will probably need to add a resource reservation to your job, unless the cluster is very empty.
- Although machines have 8, 16, 24, 48, 96, 128, 256 and 1TB of memory, in reality this translates into requestable memory of 7.9, 15.7, 23.5, 62.9, 47.2, 94.4GB. i.e. if you request 16G you will not be able to run on any of the 16G machines and your job will queue for longer.
- See memory management for more details.

Use SGE 'task arrays' to run parallel jobs

Add a line like the following to your script

- Creates an array of 100 identical tasks each with an id: SGE_TASK_ID between 1 and 100.
- For example, use this to run 100 bootstrap replications or 100 Monte Carlo replications.
 - Highly efficient, highly robust way to run multiple jobs on cluster.
- Use SGE_TASK_ID in your program to control inputs/outputs
 - e.g. set random number seed to SGE_TASK_ID and save results with name based on task id.
- Task arrays can also be used to solve much more complex parallelization tasks.