Introduction to IT for UCL Astrophysicists

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Where to find this presentation

url

https://github.com/Astrophysics-UCL/HPCInfo/tree/master/training/workshop_2016

Overall goals of presentation

- What software you might find useful
- Where to get more information (UCL courses, web, etc.)
- UCL-specific information (e.g. login details)
- Some hands-on work

Specific contents

13 October:

- Accessing Astrophysics group machines
- Using the Linux console
- Basics of Python

20 October:

- Commonly used programs (LaTeX, DS9, IRAF,...)
- Using High-Performance Computing (HPC) machines
- HPC best practices

Information on the Web

Astrophysics Wiki

https:

//wiki.ucl.ac.uk/display/PhysAstAstPhysGrp/Main+Page This Wiki is freely viewable and editable by all members of the department. Please use it to record information that you think will be useful to others (including your future self). Be bold!

UCL Research Computing Platforms

https://wiki.rc.ucl.ac.uk/wiki/Main_Page

Stack Overflow

http://stackoverflow.com/

Computing Environment for Astrophysics

- Large datasets requiring substantial processing followed by sophisticated statistical analysis
- Calculations often done on specialised 'high-performance computing' (HPC) machines having large filesystems and large RAM; calculations are often broken into pieces that can be run simulataneously ('in parallel') across many processors.
- Much useful software is made freely available within the community. Software quality is usually high; documentation quality is more variable. As well, many users write their own software.

Local Computing Environment

You will have your own local machine, which will have one of these operating systems:

- ► PC (Windows)
- Mac
- ► Linux

In addition there are shared Linux machines:

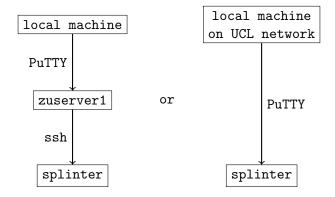
- zuserver1 (general purpose; accessable from outside UCL)
- splinter (HPC cluster)
- Others? Legion?

Work patterns

Several work patterns are possible:

- Write and test a program on your local machine; use the local machine to remotely connect to splinter; upload the program to splinter and run it there;
- Or do all your work locally (requires small data sets);
- Or use the local machine to remotely connect to splinter and do all your work there.

Accessing splinter



Accessing splinter

- ▶ You will need a *username* and *password* for splinter (and perhaps zuserver1).
- ▶ Step 1: Logon to zuserver1:
 - ► From Windows: Use PuTTY (http://www.putty.org/). On the 'Connection/SSH/X11' tab, click on 'enable X11 forwarding' and set 'X display location' to 'localhost:0'. On the Session tab, set the Host Name to zuserver1.star.ucl.ac.uk.
 - From Mac: ?? Find out what to do
 - From Linux: ?? Just do ssh?
- Step 2: From zuserver1 you can logon to splinter: ssh -YC username@splinter-login.star.ucl.ac.uk.
- ▶ If you are connecting from a machine on the UCL network then in step 1 go straight to splinter-login.star.ucl.ac.uk and omit step 2.
- Say something about X Windows forwarding.

Command shell in Linux

- ▶ In Linux you will use a 'command shell'.
- ► This is a text-based environment in which you type commands and receive text output.
- ▶ Not GUI! Reflects the hardware limitations current when Unix was created. Low-tech and reliable e.g. for remote access.
- ▶ Various command shell programs in use: bash, csh, tcsh,...

Directory structure

- ► Everything is organised around files (which may be data files or program files i.e. instructions to be executed).
- ► Files live in directories. There is a hierarchical tree structure of directories.
- ► Sample file name: /share/splinter/ucapwhi/des/foo.txt
- ▶ Note use of slash '/', not backslash '\' as in Windows.
- Case sensitivity: 'Foo' and 'foo' are different strings.

Special symbols for directories

Symbol	Meaning
/	Top of the directory tree (the root directory)
	Current directory
	Parent of the current directory
~	User's 'home' directory

Structure of commands

```
Structure
[command] -[option(s)] [argument]
Examples
ls -la
mkdir hello_world
cp hello.cpp new_hello.cpp
```

Linux command reference

There is a very useful summary of Linux commands at: http://www.computerhope.com/unix.htm

Basic Linux commands I

navigation and help

ls -la
cd dir_name
man command_name
pwd
exit

copy or move

cp src dest
mv src dest
scp usr@host:file dest

create or delete

touch file.txt
mkdir dir_name
rm -i file.txt

find and system info

whereis file which echo \$VAR_NAME

file contents

cat file
more file
head file

Basic Linux commands II

```
process management
                                 kill
&; | ¿ i
                                 top
& (background)
                                 nohup
 (combine)
  (next line)
                                 compressed files
  (combine)
                                 gunzip
* (wildcard)
                                 tar
> (output)
< (input)
                                 images
                                 gthumb
Text editors
                                 ds9
emacs
vi
                                 publishing
gedit
                                 latex
                                 bibtex
```

Exercises I

- 1. Go to your home directory and create a directory called linux_hpc_workshop.
- Change directory to linux_hpc_workshop.
- 3. Find the name of the present working directory.
- Make a directory level_1/level_2, and move to level_1/level_2 in one command.
- 5. Move back to the previous directory.
- 6. Remove the directory level_1 (and its contents).
- In the current directory make a symbolic link to usr/lib called my_sybolic_link.
- 8. Create a file called foo.txt with contents "This file contains the word foo".
- 9. Add another line in foo.txt with contents "This is the second line".
- 10. Check to see if it worked.
- 11. Search for the phrase foo in foo.txt.

Exercises II

- 1. Find the location of your python installation.
- 2. Find the installation location(s) of liblapack.a.
- 3. Find whether an object daxpy is in liblapack.a.
- 4. Find the value the environment variable PATH and LD_LIBRARY_PATH.
- Set the environment variable MY_LINUX_HPC_VAR to equal the absolute path to linux_hpc_workshop.
- 6. Add (i.e append) to the PATH the absolute path to linux_hpc_workshop.
- 7. Use the source command do the last two steps from source file.
- 8. Use the *man* command to find the option of 1s that shows the output in Kilobyte, Megabyte.

Exercises III

- Find hostname, processor type and operating system version and write this info into a text file called info.txt.
- 2. List the people who are currently logged into the system.
- 3. Find the process that is taking most of the CPU at the moment.
- 4. Find the IDs of the processes that you are running.
- Make a directory called to_be_compressed. Add the files hello.cpp and hello.py in this dir. Then compress this directory using tar and zip.
- Delete the directory to_be_compressed and extract the files from to_be_compressed.tar.gz.
- Use wget to download files from ftp://heasarc.gsfc.nasa.gov/software/fitsio/c/cfitsio3370.tar.gz.
- 8. Find the size of the item you just downloaded in MB.
- 9. Extract all files from this downloaded archive file.
- In the extracted files, find all occurrences of ffopentest in all the files with extension .h.
- 11. Remove all the files with extension .h.
- 12. Copy the files with extension .c into a new directory c_files.

Python

Vanilla Python

Numpy

Talk about slicing somewhere...

Scipy/Astropy

Matplotlib

Common and Useful Programs

LaTeX

Talk about Tikz in here

DS9

IRAF

Information on the Web

This presentation

https://github.com/Astrophysics-UCL/HPCInfo/

Splinter on the UCL Astrophysics Wiki

https://wiki.ucl.ac.uk/display/PhysAstAstPhysGrp/

Splinter+User+Guide

UCL Research Computing Platforms

https://wiki.rc.ucl.ac.uk/wiki/Main_Page

DiRAC

http://www.dirac.ac.uk/

Mailing list

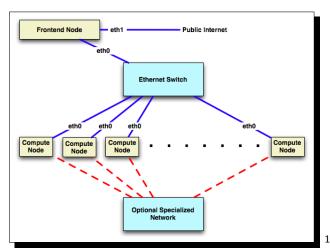
https://www.mailinglists.ucl.ac.uk/mailman/listinfo/splinter-users

- please subscribe
- post any issues regarding splinter

Splinter specs

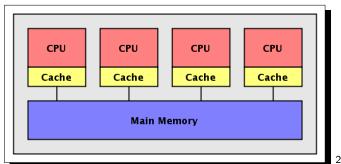
- ▶ As of October 5, 2016, *Splinter* has 528, 4TB memory
- 8 nodes, dual 6-core 2.8GHz, 48GB memory
- ▶ 20 nodes, dual 8-core 2.0GHz, 128GB memory
- SMP node, 40 2.4GHz cores, 1TB memory
- ▶ login node, dual 10-core, 2.4GHz 98GB memory
- head-node, dual 8-core, 2.4GHz, 164GB memory

SPLINTER distributed



¹http://www.rocksclusters.org/

SPLINTER shared



Workspaces I

/home/user_name

- this is your home directory
- login scripts can be put here
- 1GB quota
- private

/share/splinter/user_name

- create the directory if not already there
- can be used as a workspace
- no quota
- public unless made private

Workspaces II

/share/data1

- ▶ for storing large data
- you can create a directory for your, .e.g, /share/data1/SKA

/share/apps

- for installing software
- module-files

Login script

- everytime you login this file will be executed
- ▶ this file is in your \$HOME
- ▶ it is called .login
- you can load modules, envvars, etc.

Examples

Load my aliases

source ~/aliases.csh

Load python

module load dev_tools/nov2014/python-anaconda

Modules

- easy and flexible way use software
- available to everyone in splinter

Examples

Print the available modules module avail

Load a module module load module_name

List the loaded modules module list

Unload a module module unload module name

Unload all modules module purge

Help
module --help

Submitting jobs

- computing jobs should be submitted to the scheduler
- you will have to write a job script
- interactive job

Examples

Submit a job

qsub job_script

Submit an interactive job

qsub -I

Check the status of a job

checkjob job_id

List the status of all jobs

qstat

Show the queue

showq

Delete a job qdel job_id

Queues

- ▶ compute
- ▶ cores16
- ▶ cores12
- ▶ smp

Structure of a job script

```
#!/bin/tcsh
# PBS -q cores12
# PBS -N a_name_for_your_job
# PBS -1 nodes=1:ppn=6
# PBS -1 mem=32gb
# PBS -1 walltime=120:00:00
Set some environment variable
seteny OMP NUM THREADS 6
Source paths if needed
source /home/username/libpaths.csh
Run my program
/home/username/hello_world.exe
```

Jobscripts: things to remember

- ► Submit the job to the right queue
- ▶ Request the correct number of nodes and ppn
- Specify the memory required
- Always specify the walltime
- ▶ If your program is not parallel, please use nodes=1,ppn=1
- ▶ Use -q compute for single processor jobs
- Use qsub -I for interactive job
- If using most of the resources, please send an email to the mailing list.

More PBS commands

```
Specify output
PBS -o path/to/file.out
Specify error output
PBS -e path/to/file.err
Mail alert at (b)eginning, (e)nd, and (a)bortion of execution
PBS -m bea
Send mail to the following address
PBS -M your_email_id@ucl.ac.uk
```

Using Ganglia

http://splinter.star.ucl.ac.uk/ganglia/

- is tool for analysing splinter
- can only be loaded from splinter (using firefox)
- will give you load/memory information
- can look into nodes

Collaborative projects

- collaboration between two splinter users
- can share common data in /share/data1/my_collaboration
- give read/write permission to other users using chmod

Best practices

- ▶ Choose the machines that are suited for your problem
- Read the User Guide
- ▶ Do not run your programs in the login node
- Install common software locally if and only if absolutely necessary
- Request optimum resources
- Minimise data transfer between nodes,
- Backup! Backup! Backup!

Exercises

https://github.com/Astrophysics-UCL/HPCInfo/tree/master/training/workshop_2016/