

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 simultaneously ('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:

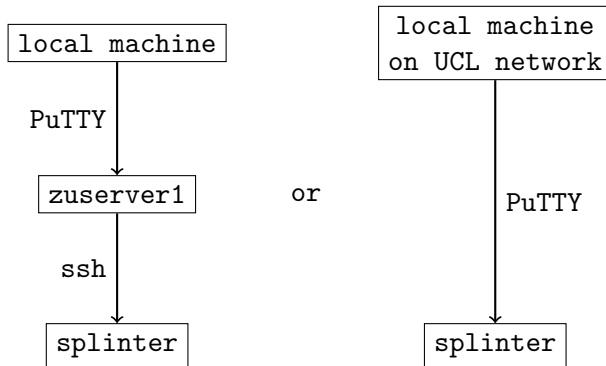
- ▶ zuserver1 (general purpose; accessible from outside UCL)
- ▶ splinter (HPC cluster)
- ▶ Others? Legion?

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*, *csch*, *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

& ; | ; i

& (background)

; (combine)

\ (next line)

| (combine)

* (wildcard)

> (output)

< (input)

Text editors

emacs

vi

gedit

process management

kill

top

nohup

compressed files

gunzip

tar

images

gthumb

ds9

publishing

latex

bibtex

Exercises I

1. Go to your home directory and create a directory called `linux_hpc_workshop`.
2. Change directory to `linux_hpc_workshop`.
3. Find the name of the present working directory.
4. 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).
7. 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`.
5. 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 `ls` that shows the output in Kilobyte, Megabyte.

Exercises III

1. 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.
5. 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`.
6. Delete the directory `to_be_compressed` and extract the files from `to_be_compressed.tar.gz`.
7. 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.
10. 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`.

Information on the Web

Documentation

<http://scipy.org/>

<http://matplotlib.org/>

<http://www.astropy.org/>

SciPy Tutorials (Also NumPy and Matplotlib)

<https://conference.scipy.org/scipy2013/tutorials.php>

SciPy Lectures (Also NumPy and Matplotlib)

<http://www.scipy-lectures.org/>

Stanford's Introduction to Scientific Python

<http://web.stanford.edu/~arbenson/cme193.html>

Dictionaries

Functions

Classes

NumPy Arrays

Basic Statistics

Slicing

Sorting

I/O

matrices?, masked arrays?

Constants

More Stats

Fitting

PyFits

World Coordinate Systems

Symbolic Calculus

Different Plotting Styles

Image Plotting

Multiple Subplots

Contour?

Exercises

Common and Useful Programs

LaTeX

Talk about Tikz in here

DS9

IRAF

Features

- ▶ Aligning with WCS (World Coordinate System)
- ▶ Scaling image contrast
- ▶ Funky colour sets (and inverse or negative)
- ▶ Regions and annotating the image
- ▶ Multiple frames; blinking and matching
- ▶ Contour plots

DS9 Website

<http://ds9.si.edu/site/Home.html>

Image Reduction and Analysis Facility (IRAF)

Features

- ▶ Uses basic Linux commands to navigate around directories, copy/move files, run tasks in the background
- ▶ A series of packages each containing various tasks
- ▶ ? to view all tasks in current package, ?? to view all tasks
- ▶ Each tasks contains various editable parameter files
- ▶ `lpar [parameter_file_name]` to view and
`epar [parameter_file_name]` to edit a parameter file
- ▶ `unlearn [parameter_file_name]` to set reset task parameters

IRAF Websites

<http://iraf.noao.edu/>

<http://iraf.net/irafdocs/>

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](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

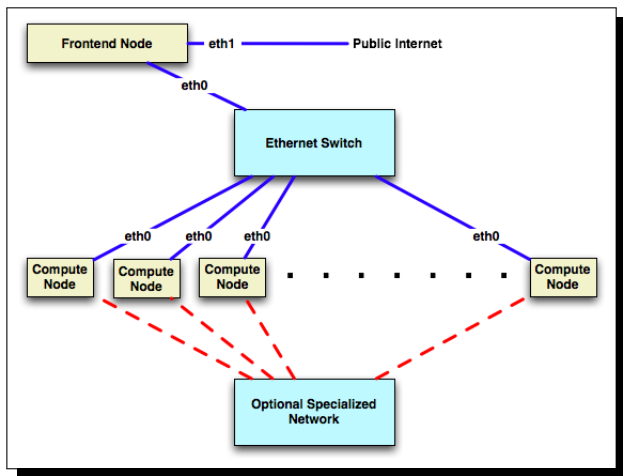
`https://www.mailinglists.ucl.ac.uk/mailman/listinfo/splinter-users`

- ▶ please subscribe
- ▶ post any issues regarding splinter

Splinter specs

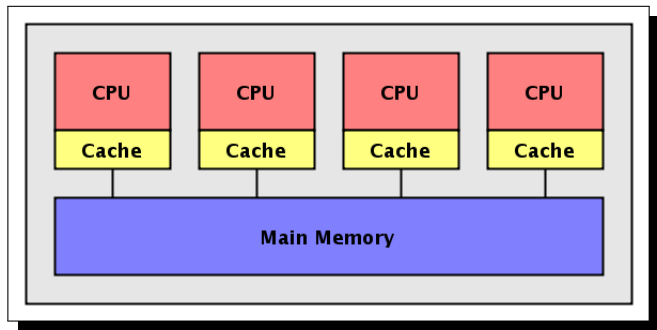
- ▶ As of October 9, 2016, *Splinter* has 528, 4TB memory
- ▶ 8 nodes, dual 6-core 2.8GHz, 48GB memory
- ▶ 20 nodes, dual 8-core 2.0GHz, 128GB memory
- ▶ SMP node, 40 2.4GHz cores, 1TB memory
- ▶ login node, dual 10-core, 2.4GHz 98GB memory
- ▶ head-node, dual 8-core, 2.4GHz, 164GB memory

SPLINTER distributed



1

SPLINTER shared



2

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 -l nodes=1:ppn=6
# PBS -l mem=32gb
# PBS -l walltime=120:00:00
```

Set some environment variable

```
setenv 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/