Linux HPC Workshop

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How to access training materials?

url

https://github.com/Astrophysics-UCL/HPCInfo/tree/master/training/linux_hpc_workshop_oct_2014



What will you learn?

- Accessing Astrophysics group machines
- Using linux console for your research
- Running your programs in HPC machines

Accessing machines from outside

You will need a username and password

```
# step 1 login to zuserver
ssh -YC username@zuserver.star.ucl.ac.uk

# step 2 login to other machines from
# zuserver

ssh -YC username@splinter.star.ucl.ac.uk
```

command structure

structure

```
# [command] -[option[s]] -[argument]
```

Example

```
ls -la
makdir hello_wrold
cp hello.cpp new_hello.cpp
```

Linux console cheat sheat I

navigation and help

```
ls -lah dir_name
cd dir_name
cd ..
cd -
man command_name
pwd
exit
```

copy or move

```
cp src dest
cp -r src dest
mv src dest
ln -s src targ
```

create or delete

```
touch file.txt
mkdir dir_name
mkdir -p prt/dir
rm -i file.txt
rm -rf dir_name
```

find or search

```
locate file
whereis file
grep "bla" file
awk 'pattern' file
```

Linux console cheat sheat II

cat file more file less file head file tail file nm object_file readelf shared_obj_file ldd executable

process management

file contents

```
ps -e
kill
killall
top
```

ssh

```
ssh usr@host
ssh -YC user@host
scp usr@host:file dest
```

system info

```
uname -a
who
whoami
whois
which
finger
ping
echo $VAR_NAME
```

Linux console cheat sheat III

```
&:|i|
& # backgraound
; # combine
\ # next line
| # combine
* # wildcard
> # output
< # input</pre>
```

```
Text editors

emacs
vi
gedit
```

firefox google-chrome wget curl

```
publishing

latex
pdflatex
bibtex
```

Linux console cheat sheat IV

gzip gunzip tar xvzf tar cvzf tar xvjf tar xvJf

```
eog
xfig
gimp
gthumb
convert
```

```
make
cmake
python
gcc
g++
gfortran
```

```
gnuplot
R
matlab
IDL
```

Exercises I

- In your home directory create a directory called linux_hpc_workshop
- Change directory to linux_hpc_workshop
- What is the present working directory
- Make a directory level_1/level_2, and move to level_1/level_2 in one command
- Move back to previous directory
- 6 Remove the directory (and its contents) level_1
- Make a symbolic link to usr/lib in the current directory called my_sybolic_link
- 6 Create a file called bla.txt contents "this file has a word called bla"
- Add another line in bla.txt called "this is the second line"
- Check if it worked
- Search for the phrase bla in bla.txt



Exercises II

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



Exercises III

- Find hostname, processor type, operating system version and write these info inot a text file called info.txt
- 2 Find the list of people who are loged into the system
- 3 Find the process that is taking most of the CPU at the moment
- Find 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 Now 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 What is the size of the item you just downloaded in MB
- Find the number of occurrences of the phrase table is easy in all the files with extension .h
- 10 Remove all the fiels with extension .h
- Copy the files with extension .c into a new directory c_files

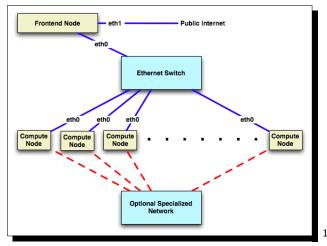


HPC Facilities

machine	type	cores	memory
SPLINTER-1	distributed	96	48GB
SPLINTER-2	shared	40	1TB
PHALANX	shared	32	512GB



SPLINTER distributed

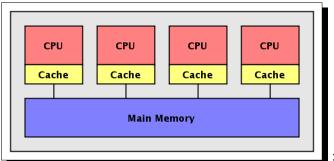


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¹http://www.rocksclusters.org/

SPLINTER shared



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²http://www.cs.rit.edu/

Best practices I

- Choose the machines that are suited for your problem
- Read the User Guide
- Do not run your programs in the login node
- Do not install common software locally
- Request optimum resouces
- Minimise data transfer between nodes,
- Backup! Backup! Backup!

Submitting jobs

commands

```
qsub jon_script
qsub -I
checkjob job_id
qstat
showq
qdel
```

Example

```
#!/bin/bash
#PBS -N hello_world_program
\#PBS - l \quad nodes = 1: ppn = 4
\#PBS - l mem = 2qb
#PBS -j oe
#PBS -V
# source the required scripts
# this sets the PATH
source /home/sbalan/binpaths.sh
# this sets the LD_LIBRARY_PATHS
source /home/sbalan/libpaths.sh
# run my program
/home/sbalan/hello.exe
```

Exercises III

- Login to your HCP machine and find the path to your HOME directory and your quota
- Find the processor type and the version of your operating system
- Request an interative queue and run the hello_world.exe
- Sumbit hello_world.exe using a job script, find its jobid, check the output log.
- Ompile big_mem_example, submit it using a job-script and see how much menory it uses
- Compile time_pause_example, submit it using a job-script and kill this job using its jobid.
- In the previous example see what happens when you play with the time requested.



More information

ap-wiki

http://www.ucl.ac.uk/star/GroupAWiki

UCL Research Computing Platforms

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

DiRAC

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

