



CCAS and SEAS Computing Facility



HPC Workshop 3

For new Colonial One users!!!

Login to Colonial One using your GW NetID/email address & password

```
$ ssh <netid>@login.colonialone.gwu.edu
$ cd ~
$ cd skel
$ cp .bash* ../
$ cd ~
```



HPC Workshop 3

What we're covering:

- Simple bash scripting
- Installing software in your home directory
- Virtualenv Python Demo
- Install R modules in your home directory
- Parallel module



Overview: A shell script is a computer program designed to be run by the Unix/Linux shell, a command-line interpreter.

- Shell scripts are just executable text files
- Use shell scripts to perform repetitive tasks
- Shell scripts can include any command you run on the command line
- Scripting allows you to use programming functions such as 'for' loops, if/then/else statements, etc.
- You can use command substitution in a shell script, such as using the 'date' command to name a file.



Create a simple shell script file to make a directory for your today's workshop and copy the workshop files into the directory:

- \$ cd ~
- \$ nano workshop copy.sh
- Type:

```
cd ~
mkdir ~/hpcworkshop_3
cp /groups/hpcworkshop/ws3/* ~/hpcworkshop 3
```

- Make your script executable: chmod +x workshop copy.sh
- Execute the script: ./workshop_copy.sh
- Is the new directory to see the files: ls /home/<username>/hpcworkshop_3

Alternatively, one can use git to install the materials:

```
module load git
git clone https://github.com/bindatype/hpcworkshop.git
```



Shell scripts that call modules should be loaded with the "source" command:

Text is located in ~/hpcworkshop_3/load_python.sh

- nano loadpython.sh
- Type:

```
#!/bin/bash
module load python/3.4.2
python
```

- Make your script executable: chmod +x loadpython.sh
- Execute module script with the "source" command: source loadpython.sh
- Type quit () to exit the Python command line



You can use shell scripts to load modules for specific tasks, like compiling a large program

Text is located in ~/hpcworkshop 3/load paraview.sh

To load the Paraview build environment:

```
!#/bin/bash
module load openmpi/current
module load python/2.7.6
module load cmake/3.3.1
module load openblas/openblas
module load qt
```

Save the text in a file and you can load all those modules with one command:

```
$ source load_paraview.sh
```



For Loop Example

Text is located in ~/hpcworkshop_3/for_loop.sh

```
#!/bin/bash
for i in 1 2 3 4 5
do
    echo "Welcome $i times"
done
```

Make your script executable, then run it:

```
$ chmod +x for_loop.sh
./for loop.sh
```



For Loop Example Continued ... looping over an array

```
LIST=(`ls /home/$USER`)
for NAME in ${LIST[@]}; do echo $NAME; done
for NAME in ${LIST[@]}; do echo $NAME | tr '[a-z]'; done
```

For More Complicated Loop Example

```
LIST=(`ls /home/$USER/hpcworkshop_3/*sh`)
for NAME in ${LIST[@]}; do echo $NAME | awk '{split($0,a,"\\.sh") ;print
a[1]".bash"}'; done
```



Questions & Discussion



Installing your own software

You can install software in your home directory on Colonial One

- Users do not have root on Colonial One, so software with many dependences should be installed by the C1 team
- You can compile programs directly in your home directory
- You can also download precompiled utilities and run them from your home directory



Installing your own software

Figlet Example

CTRL+C to exit

Text is located in ~/hpcworkshop_3/figlet_example

\$ cd ~
 \$ mkdir figlet; cd figlet
 \$ wget ftp://ftp.figlet.org/pub/figlet/program/unix/figlet-2.2.5.tar.gz
 \$ tar -xvf figlet-2.2.5.tar.gz
 \$ cd figlet-2.2.5
 \$ nano Makefile
 o prefix = /home/<username>/figlet
 o DEFAULTFONTDIR = /home/<username>/figlet/figlet-2.2.5/fonts
 \$ make all
 ./figlet (test out your program!)
 Type something and hit enter



Your own Python Environment

Create your own Python environment with Virtualenv

- Regular Colonial One users do not have permission to alter the system-wide Python installations
- Virtualenv creates your own custom python build inside a directory you choose
- You can install additional python libraries without going through the C1 team
- You can build multiple Python environments for different projects, including different versions of the Python executable and different libraries for each project



Your own Python Environment

Virtualenv Example

Text is located in ~/hpcworkshop_3/virtualenv_example

- 1. cd ~
- 2. mkdir my virtual env
- 3. cd my virtual env
- 4. module load python/3.4.2 < --- The environment will build based off the version of Python you've loaded
- 5. virtualenv test env
- 6. source ~/my virtual env/test env/bin/activate
- 7. pip install numpy, scipy, times, etc.

You can unload the virtual environment by deactivating it:

deactivate



Your own Python Environment

Questions & Discussion



Build your own modules

You can build your own modulefiles if you want to customize environment variables.

Create a folder called "modulefiles" in your home directory:

```
$ cd ~
$ mkdir modulefiles
$ cp ~/hpcworkshop_3/mypython ~/modulefiles/
$ cd ~/modulefiles/
```

Look at the text file for your new module:

```
$ nano mypython
```



Build your own modules

The modulefile:

```
"Adds python 2.7.5 to your environment"
module-whatis
                          /c1/apps/python/2.7.5
set
             root
                          $root/include
prepend-path
             INCLUDE
             LD LIBRARY PATH $root/lib
prepend-path
                          $root/lib
prepend-path
             LIBRARY PATH
                          $root/bin
prepend-path
             PATH
```

CTRL+o to save CTRL+x to exit



Build your own modules

Configure the module system to use your new module directory:

\$ module use /home/<username>/modulefiles (tab complete does not work)

List available modules:

\$ module avail

Load your new module:

\$ module load mypython

Confirm that your module loaded:

\$ module list



Your own R libraries

Installing R libraries in your home directory is very simple - you create a directory, set an environment variable, and that's it!

```
$ cd ~
$ mkdir R_libs
$ export R_LIBS="/home/<username>/R_libs"
$ module load R
$ R
   install.packages('sqldf', repos="http://cran.r-project.org")
   quit()
```

Check your R_libs directory: ls ~/R libs

To make the change permanent, add the export command to your .bashrc file:

```
$ echo export R_LIBS="/home/<username>/R_libs" >> ~/.bashrc
```



Your own R libraries

Questions & Discussion



Parallel is a module that lets you run a single process across multiple cores or CPUs.

Basic usage:

```
parallel [options] [command [arguments]] < list_of_arguments
parallel [options] [command [arguments]] (::: arguments|::: argfile(s))...</pre>
```

The command can be one command or a script you've written. Parallel will launch concurrent instances of the command.

Parallel is a module:

```
$ module load parallel
```



 \mathbf{E}

The parallel module

Parallel will automatically try to use as many threads as necessary based on the amount of input arguments:

```
$ parallel echo ::: A B C D E
A
B
C
D
```



You can specify the total number of processes to use at the same with with the -j option. the {%} string represents the Job slot.

```
$ parallel -j 2 echo {%} {} ::: A B C D E
```

- 1 A
- 2 B
- 1 (
- 2 D
- 1 E



Parallel will index the total number of processes run. The {#} string represents the total process number.

```
$ parallel -j 2 echo {#} {} ::: A B C D E
```

- 1 A
- 2 B
- 3 C
- 4 D
- 5 E



Serial Fibonacci number example. Use the fibonacci_serial program to find specific numbers in a Fibonacci sequence:

```
$ parallel ~/hpcworkshop_3/fibonacci_serial ::: 42 43 44 45
```

Result:

42 267914296

43 433494437

44 701408733

45 1134903170



You can use a text file with the same arguments instead of typing them out on the command line:

```
$ nano input.txt
Type:
42
43
44
45
$ parallel ~/hpcworkshop 3/fibonacci serial < input.txt</pre>
Result:
42 267914296
43 433494437
44 701408733
45 1134903170
```



Questions?

Please fill out the survey for the workshop. Thank you!

https://goo.gl/HiGqZC