Bash Shell Basics

This is a concise refresher on the basic commands available with the Bash shell. It is designed as a prerequisite module to the Supercomputing User Training offered by **Pawsey Supercomputing Research Centre**, Perth (Australia).

For a more detailed tutorial on the Bash shell, see The Unix Shell, by the Software Carpentries.

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Definitions

- Command Line Interface (CLI): a type of interface to use a computer, that
 requires the user to type textual commands and read their textual outputs by
 means of a prompt. Some commands may require optional or mandatory
 arguments.
- Shell: a program with a prompt providing a command line interface, and an
 interpreter to execute commands. There are different implementations of a shell,
 using different shell languages for the commands.
- **Bash**: one of the most popular shells in the Linux world.
- Shell script: a text file containing a sequence of shell commands. It can be
 executed via the shell, and is useful to increase reproducibility of a workflow.
- Main advantages of using a shell:
 - Productivity (lots of actions with few keystrokes)
 - Task automation
 - Reproducibility (using shell scripts)
- Text editor: a program used to open and edit textual files; some of them are
 designed for use within a command line interface, such as nano, vi and emacs.

NOTE: in all the examples below, by convention commands are identified by lines starting with \$, which represents the prompt (not to be typed). Other lines represent command outputs. Also note how # denotes a comment line in Bash.

Hands-on

Handling directories

First, navigate to the home directory using cd; here, ~ refers to the home, and can be omitted if used alone:

```
$ cd ~
$ # or also
$ cd
```

Check the path of the current working directory with pwd:

```
$ pwd
/home/username
```

Create a new directory with mkdir:

```
$ mkdir training
```

Enter it, and check where you are:

```
$ cd training
$ pwd
/home/username/training
```

The current directory can be referred to using a single dot, .:

```
$ cd .
$ pwd
/home/username/training
```

The **parent directory** can be referred to using a double dot, ...:

```
$ cd ..
$ pwd
/home/username
$ cd training
```

Create two more directories inside training, create one more into the first one of these, and then access the first one:

```
$ mkdir dir1 dir2
$ mkdir dir1/one-more-dir
```

```
$ cd dir1
```

A **full path** does not depend on the current working directory, and starts with a slash, /, e.g.: /home/username/training.

A **relative path** depends on the current working directory, and starts with a directory name, a . or a . . , e.g.: one-more-dir, . . / one-more-dir, . . / dir2.

Go back to the training parent directory:

```
$ cd ..
$ pwd
/home/username/training
```

Handling files

Now let us play with files. Make new empty files using touch:

```
$ touch file1
$ touch file2 file3
$ touch dir1/file4
```

The command is actually designed for altering timestamps of existing files; however it will create an empty file, if the provided filename does not match an existing one.

Check the content of a directory with 1s:

```
$ ls
dir1 dir2 file1 file2 file3
$ ls dir1
file4 one-more-dir
```

Get more details about files and directories with 1s -1:

```
$ ls -l total 8 drwxr-x--- 3 username username 4096 Aug 26 14:10 dir1 drwxr-x--- 2 username username 4096 Aug 26 14:10 dir2 -rw-r---- 1 username username 0 Aug 26 14:10 file1 -rw-r---- 1 username username 0 Aug 26 14:10 file2 -rw-r---- 1 username username 0 Aug 26 14:10 file3
```

To make a copy of a file with another name use cp:

```
$ cp file1 copy1
```

To copy a directory use cp - r:

```
$ cp -r dir1 copydir1
```

To rename a file use mv:

```
$ mv copy1 another-file
```

To move a file to another directory use mv again, but now with the second argument as an existing directory:

```
$ mv another-file dir2/
```

Use rm to remove a file:

```
$ rm file3
```

Use rm -r to remove a directory:

```
$ rm -r copydir1
```

Multiple files and/or directories can be selected using the **wildcard** characters? and *. The question mark, ?, means any value of a single character:

```
$ ls file?
file1 file2
```

The asterisk, *, means any value of any number of characters:

```
$ ls f*
file1 file2
```

When the * also includes directories, 1s will output their content, too:

```
$ ls *
file1 file2

dir1:
file4 one-more-dir

dir2:
another-file
```

You can assign a **value**, *i.e.* a string of characters, to an environment **variable**. We call **shell environment** the whole set of variables defined in the current shell session. Assign a variable using the equal operator, =:

```
$ HELLO="world"
```

The value can be reused later in the same shell session by referring to the corresponding variable. For instance, use echo to just display the variable value:

```
$ echo $HELLO
world
```

To make the variable accessible inside sub-processes, you need to export it:

```
$ export BYE="moon"
```

A process could be a program, a shell script, or just another Bash shell. Let us verify the latter case. Start a new Bash shell with bash, and then exit it when done:

```
$ bash
$ echo $HELLO

$ echo $BYE
moon
$ exit
exit
```

You can use a variable to store a directory path:

```
$ pwd
/home/username/training
$ MYDIR="/home/username/training"
```

And then use it:

```
$ echo $MYDIR
/home/username/training
$ ls $MYDIR
dir1 dir2 file1 file2
```

There is a method to store the output of a command inside the variable: ncapsulating the command within the syntax \$(). For instance, use such syntax to capture and store the output of pwd:

```
$ MYDIR_AGAIN="$(pwd)"
$ echo $MYDIR_AGAIN
/home/username/training
```

Every shell session automatically defines a number of environment variables, some of which can be quite useful. Among others, \$HOME contains the path to your home directory, \$USER contains your username, and \$PATH has the list of paths where the shell looks for executable programs and scripts:

```
$ echo $HOME
/home/username
$ echo $USER
username
$ echo $PATH
/usr/local/bin:/usr/bin
```

Editing and visualising text files

Let us now use the text editor nano to create, edit and save a text file. First, start the editor with a new empty text filw called text:

```
$ nano text
```

Now, type the following in the editor:

```
Line 1
Fancy line 2
Super line 3
Extra line 4
Last line 5
```

To save this text, press Ctrl-0 (Control + 0), then confirm with Enter. To exit the editor, press Ctrl-X.

Visualise the full contents of the file from the shell using cat:

```
$ cat text
Line 1
Fancy line 2
Super line 3
Extra line 4
Last line 5
```

For long files, the command less allows showing the contents one screen at a time. The next screen can be accessed using Space or Ctrl-F (forward), the previous screen

using Ctrl-B (backward). Quit the less screen with q. You can trial this with the file you just created, although the output will most likely fit in a single screen:

```
$ less text
Line 1
Fancy line 2
Super line 3
Extra line 4
Last line 5
text lines 1-5/5 (END)
$ # quit by typing `q`
```

Use head/tail to visualise the first/last lines of a file. By default 10 lines are displayed, but a custom number can be specified using a specific command option:

```
$ head -1 text
Line 1
$ tail -1 text
Last line 5
```

Manipulating outputs and text files

Use wc to get the line, word and character count of a text:

```
$ wc text
5 14 58 text
```

You can see that the text file has 5 lines, 14 words and 58 characters (including newlines).

Use grep to filter text lines matching a specific string pattern. Search for the string Super in the text file created above:

```
$ grep Super text
Super line 3
```

Commands can be connected together to feed the output of a command as the input for the following command. To connect commands use the **pipe operator**, I:

```
$ echo "Hello world from Pawsey." | wc
1 4 25
```

Redirection operators can be used to connect a file with the output or input of a command.

Use > to redirect a command output into a file (file will be overwritten):

```
$ echo "Hello world from Pawsey." >hello
$ cat hello
Hello world from Pawsey.
$ echo "Hello again." >hello
$ cat hello
Hello again.
```

Use >> to append a command output into a file (file will not be overwritten):

```
$ echo "Bye bye from Pawsey." >>bye
$ echo "Bye bye again." >>bye
$ cat bye
Bye bye from Pawsey.
Bye bye again.
```

Use < to feed the contents of a file as command input:

```
$ wc < bye
2 7 36
```

Other useful commands for file and output manipulations include cut, tr, sort, sed and awk.

More commands

Use hostname to output the name of the computer/server where the shell session is open:

```
$ hostname
setonix-01
```

Use which to get the full path of a program:

```
$ which grep
/usr/bin/grep
```

Some commands are not programs (see next command to know more):

```
$ which cd
which: no cd in (/usr/local/bin:/usr/bin:/bin)
```

Use type to get information about the typology of command:

```
$ type grep
grep is /usr/bin/grep
```

```
$ type cd
cd is a shell builtin
```

Sometimes it is useful to group together a set of files and/or directories within a single archive file, which is typically called tarball and assigned the extension .tar. The tar program is used to this end.

Create a tarball containing all files and/or directories (with their contents) whose name starts with file or dir:

```
$ tar cf archive1.tar file* dir*
```

Visualise its contents:

```
$ tar tf archive.tar
file1
file2
dir1/
dir1/one-more-dir/
dir1/file4
dir2/
dir2/another-file
```

Now create a directory expand, enter it, unpack the tarball there, and check the directory contents:

```
$ mkdir expand
$ cd expand
$ tar xf ../archive1.tar
$ ls
dir1 dir2 file1 file2
cd ..
```

If you also want to compress the tarball to reduce its size, add the option z to the commands above, and use the conventional extension .tar.gz:

```
$ tar czf archive2.tar.gz file* dir*
$ ls -l archive*
-rw-r---- 1 username username 20480 Aug 26 15:59 archive1.tar
-rw-r---- 1 username username 581 Aug 26 16:03 archive2.tar.gz
```

The latter tarball takes 581 bytes instead of 20480. Note that compression rates may vary significantly depending on the file contents.

Other useful Bash syntax constructs include for/while loops and if/elif/else conditionals.

Keyboard tricks

Use Up-arrow and Down-arrow to browse the shell history of commands:

```
$ # hit Up-arrow once
$ ls -l archive*
$ # hit Up-arrow again
$ tar czf archive2.tar.gz file* dir*
$ # hit Down-arrow once
$ ls -l archive*
```

Note you will need to hit Enter to actually execute the selected command, as usual.

Tab completion is a handy feature that can reduce typing and make shell usage more convenient. Start typing a command, then hit Tab to see available commands whose name start with the typed characters:

```
$ # type without hitting Enter
$ host
$ # hit Tab twice
host hostid hostname hostnamectl
$ hostn
$ # hit Tab again
$ hostname
```

Again, to execute the selected command hit Enter, after eventually having edited it or added more arguments.

If you need to stop what you are typing, or a running program, and get back to an empty shell within the same session, use Ctrl-C:

```
$ # type without hitting Enter
$ hostname
$ # hit Ctrl-C
$ hostname^C
$
```

This is all for this Bash shell refresher. You can close the shell session with exit!