Introduction and refresher

INTRODUCTION TO BASH SCRIPTING



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Introduction to the course

This course will cover:

- Moving from command-line to a Bash script
- Variables and data types in Bash scripting
- Control statements
- Functions and script automation

Why Bash scripting? (Bash)

Firstly, let's consider why Bash?

- Bash stands for 'Bourne Again Shell' (a pun)
- Developed in the 80's but a very popular shell today. Default in many Unix systems, Macs
- Unix is the internet! (Running ML Models, Data Pipelines)
 - AWS, Google, Microsoft all have CLI's to their products

Why Bash scripting? (scripting!)

So why Bash scripting?

- Ease of execution of shell commands (no need to copy-paste every time!)
- Powerful programming constructs

Expected knowledge

You are expected to have some basic knowledge for this course.

- Understand what the command-line (terminal, shell) is
- Have used basic commands such as cat, grep, sed etc.

If you are rusty, don't worry - we will revise this now!

Shell commands refresher

Some important shell commands:

- (e)grep filters input based on regex pattern matching
- cat concatenates file contents line-by-line
- tail \ head give only the last -n (a flag) lines
- wc does a word or line count (with flags -w -l)
- sed does pattern-matched string replacement

A reminder of REGEX

'Regex' or regular expressions are a vital skill for Bash scripting.

You will often need to filter files, data within files, match arguments and a variety of other uses. It is worth revisiting this.

To test your regex you can use helpful sites like regex101.com



Some shell practice

Let's revise some shell commands in an example.

Consider a text file fruits.txt with 3 lines of data:

banana apple carrot

If we ran grep 'a' fruits.txt we would return:

banana
apple
carrot



Some shell practice

But if we ran grep 'p' fruits.txt we would return:

apple

Recall that square parentheses are a matching set such as [eyfv]. Using ^ makes this an inverse set (**not** these letters/numbers)

So we could run grep '[pc]' fruits.txt we would return:

apple carrot



Some shell practice

You have likely used 'pipes' before in terminal. If we had many many fruits in our file we could use sort | uniq -c

- The first will sort alphabetically, the second will do a count
- If we wanted the top n fruits we could then pipe to wc l and use head

```
cat new_fruits.txt | sort | uniq -c | head -n 3
```

```
14 apple
```

13 bannana

12 carrot

Let's practice!

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Your first Bash script

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Bash script anatomy

A Bash script has a few key defining features:

- It usually begins with #!/usr/bash (on its own line)
 - So your interpreter knows it is a Bash script and to use Bash located in /usr/bash
 - This could be a different path if you installed Bash somewhere else such as /bin/bash (type which bash to check)
- Middle lines contain code
 - This may be line-by-line commands or programming constructs

Bash script anatomy

To save and run:

- It has a file extension .sh
 - Technically not needed if first line has the she-bang and path to Bash (#!/usr/bash), but a convention
- Can be run in the terminal using bash script_name.sh
 - Or if you have mentioned first line (#!/usr/bash) you can simply run using
 ./script_name.sh

Bash script example

An example of a full script (called eg.sh) is:

```
#!/usr/bash
echo "Hello world"
echo "Goodbye world"
```

Could be run with the command ./eg.sh and would output:

Hello world Goodbye world



Bash and shell commands

Each line of your Bash script can be a shell command.

Therefore, you can also include pipes in your Bash scripts.

Consider a text file (animals.txt)

```
magpie, bird
emu, bird
kangaroo, marsupial
wallaby, marsupial
shark, fish
```

We want to count animals in each group.

Bash and shell commands

In shell you could write a chained command in the terminal. Let's instead put that into a script (group.sh):

```
#!/usr/bash
cat animals.txt | cut -d " " -f 2 | sort | uniq -c
```

Now (after saving the script) running bash group.sh causes:

```
2 bird
1 fish
2 marsupial
```

Let's practice!

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Standard streams & arguments

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STDIN-STDOUT-STDERR

In Bash scripting, there are three 'streams' for your program:

- STDIN (standard input). A stream of data into the program
- STDOUT (standard output). A stream of data **out** of the program
- STDERR (standard error). Errors in your program

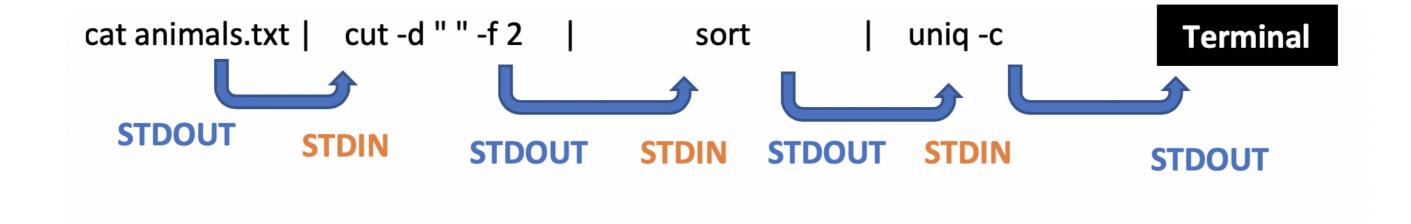
By default, these streams will come from and write out to the terminal.

Though you may see 2> /dev/null in script calls; redirecting STDERR to be deleted. (
1> /dev/null would be STDOUT)



STDIN-STDOUT graphically

Here is a graphical representation of the standard streams, using the pipeline created previously:



STDIN example

Consider a text file (sports.txt) with 3 lines of data.

football
basketball
swimming

The cat sports.txt 1> new_sports.txt command is an example of taking data from the file and writing STDOUT to a new file. See what happens if you cat new_sports.txt

football basketball swimming



STDIN vs ARGV

A key concept in Bash scripting is arguments

Bash scripts can take **arguments** to be used inside by adding a space after the script execution call.

- ARGV is the array of all the arguments given to the program.
- Each argument can be accessed via the \$ notation. The first as \$1, the second as \$2 etc.
- \$@ and \$* give all the arguments in ARGV
- \$# gives the length (number) of arguments

ARGV example

Consider an example script (args.sh):

```
#!/usr/bash
echo $1
echo $2
echo $@
echo "There are " $# "arguments"
```

Running the ARGV example

Now running

bash args.sh one two three four five

```
one
two
one two three four five
There are 5 arguments
```

```
#!/usr/bash
echo $1
echo $2
echo $@
echo "There are " $# "arguments"
```

Let's practice!

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