# 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 -1 )
- 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 <code>[eyfv]</code> . Using <code>^</code> makes this an inverse set <code>(not these letters/numbers)</code>

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 -1 and use head

```
cat new_fruits.txt | sort | uniq -c | head -n 2
```

```
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 (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
- 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

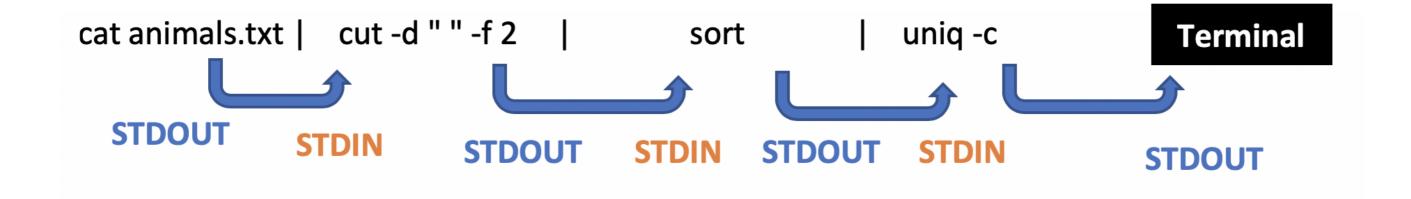
By default, these streams will come from and write out to the terminal.

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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