

## FIT1043 Introduction to Data Science

Week 9

Ian K T Tan

School of Information Technology Monash University Malaysia

With materials from Wray Buntine, Mahsa Salehi



# UNIX Shell Scripts for Data Science





## Week 9 Outline

- Characterising data and "big data"
  - the V's
  - Metadata
  - Dimensions of data
  - Growth laws
- Introduction to Unix Shell for data science
  - Why Unix shell
  - Useful commands to read/manipulate large data files



# **Learning Outcomes**

Week 9

## By the end of this week you should be able to:

- Characterize data sets used to assess a data science project
- Explain what Big data is
- Understand the V's in Big data
- Understand and analyse the growth laws: Moore's Law, Koomey's Law, Bell's Law and Zimmerman's Law
- Analyze and use shell commands to read and manipulate big data



## What is a UNIX Shell

↑ mcarman — bash — 80×24

Last login: Mon Aug 29 12:55:11 on ttys002

dyn-49-127-70-237:~ mcarman\$ ■

Command line interface to a Unix computer

Different shells have been around since the 70s.

Why are shells interesting for Data Scientists?

- Provide powerful & easy way to manipulate large data files
- And move data around a network

Available on most Unix based operating systems

- Linux
- MacOS (BSD based)



# Why Shell Scripting

Super-computers are typically UNIX/Linux based

- Explore data before you use it in Python or R
- Easier to manipulate (aka wrangle) Big Data
  - Simple and easy to learn.
  - Ideal for textual data, e.g. unstructured data for social networks, life sciences, system logs and so on.
  - Quick to sort, search, match, replace and clean your data.





# **BASH Basics I**

```
#!/bin/bash
echo "Hello world!"
```



# **Navigating the Filesystem**

#### Change directory:

```
cd [destination]
cd /my/favourite/place
```

#### Special cases:

```
cd .. <- Takes you up one directory</li>cd <- Without argument, takes you to home directory</li>
```

#### List files in the current directory:

ls

#### Copy files from one location to another:

```
cp [source] [destination]
cp /Downloads/myfile .
```



# Reading a Text File

#### Open a text file for reading using less:

```
less myfile.txt
```

#### Navigate within the text file using



## **Some Useful Commands**

Count the number of words/lines in a file

```
wc myfile.txt
```

Find lines in a file containing a keyword

```
grep "elephant" myfile.txt
```

Print the first/last few lines of a file

```
head myfile.txt tail myfile.txt
```

Print the contents of a file to the screen

```
cat myfile.txt
```



# Flags and Arguments

Many programs take flags and command line arguments that modify their behaviour, for example:

Sort the contents of a file lexicographically(alphabetically)

```
sort myfile.txt
```

Sort the lines of a file by numeric value

```
sort -n myfile.txt
```

Sort the data by column one, then column two and finally column three:

```
sort -k1,3 myfile.txt
```



# **BASH Basics II**

```
#!/bin/bash
echo "Hello world!"
```



# **Pipes**

Sometimes we'd like the output of one program to be used as the input to another.

Doing this is super easy in the shell. We just use the pipe operator "|"

```
program1 | program2
```

We can chain as many programs together as we want, for example:

```
cat hourly_44201_2014-06.csv.gz | gunzip | less
```



# Pipes (Notes)

The pipe is buffered

Each program in the list only generates data as it is needed by the next stage in the pipeline.

Thus **memory requirement** for processing the data is limited

Crucial for **scaling** up processing to **enormous data files** 



## Redirects

If we want to save the results in a file rather than pipe them to a new program:

Just change the pipe operator "|" to be a greater than symbol ">" and provide a filename:

```
cat hourly_44201_2014-06.csv.gz | gunzip > newFile.txt
```



## Wildcards

Some unix commands can take multiple files as input, for example:

```
cat myfile1.txt myfile2.txt
```

In order to avoid listing large number of files, we can use the wildcard syntax to specify all files in a directory with a certain pattern, e.g.:

```
cat myfile*.txt
```



## awk

In the tutorial, we'll have a look at a powerful command for processing text files one line at a time called awk

#### awk **syntax**:

```
awk '[select line?] {do something}'
```

#### Example

```
awk 'rand()<1/100 {print $6,$7,$14}'</pre>
```

Since awk processes data one line a time, it can scale up to massive datasets!



# **Scripts & Parallel Execution**

You can also use scripts written in Python as programs in the shell (you are encouraged to explore this).

And find out how to run programs in parallel using the ampersand notation:

myprogram &



## **End of Introduction**

We'll be experimenting with the Unix shell in this week's tutorial (or you may already have ©)

There are MANY excellent shell tutorials online if you'd like to learn more!



# Re-cap: Learning Outcomes

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