

# Intro to Unix

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#### Winter 2025 Workshop Series

#### Winter Series

Workshop	Date	Location	Registration	
How to think in Code	Jan. 28 1PM-3PM	EDUC 133	Closed	
Intro to Git & GitHub	Jan. 30 1PM-5PM	EDUC 133	Closed	
Intro to Unix	Feb. 61PM-5PM	EDUC 133	<u>Open</u>	
Intro to Python (Part 1)	Feb. 11 1PM-5PM	EDUC 133	<u>Open</u>	
Intro to R (Part 1)	Feb. 13 1PM-5PM	EDUC 133	<u>Open</u>	
Exploring MATLAB	Feb. 18 1PM-5PM	EDUC 133	<u>Open</u>	
Statistics in R (Part 2)	Feb. 20 1PM-5PM	EDUC 133	<u>Open</u>	
Data Processing in Python	Feb. 25 1PM-5PM	EDUC 133	<u>Open</u>	
Intro to Machine Learning	Mar. 13 1PM-5PM	EDUC 133	TBA	
Intro to R (Part 1)	TBA	EDUC 133	TBA	
Intro to Python (Part 1)	TBA	EDUC 133	TBA	

https://www.mcgill.ca/micm/training/workshops-series



## Outline

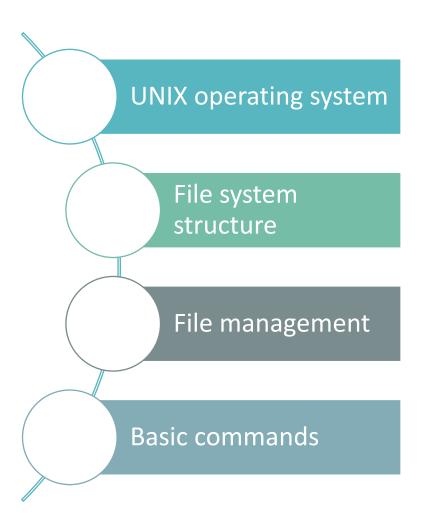
- Module 1 0
  - UNIX operating system
  - Filesystem structure
  - File management operations
  - Basic commands
- Module 2
  - Pipes and redirects
  - Variables
  - Pattern matching
  - Text processing

- Module 3
  - Shell scripting
  - Control structures
  - High performance computing (HPC)

Materials: https://github.com/McGill-MiCM/MiCM Winter2024



### Module 1







### UNIX Operating system

- General purpose and interactive
- Multiuser
- Resource-efficient
- Hierarchical file system
- Compatibility (files, devices, processes)
- Time sharing



#### Parts

#### The kernel

- The core of the operating system
- Manages processes' time, memory and resources
- Handles file storage
- Responds to system calls

#### The shell

- Let's the user communicate with the kernel
- Command line interpreter (CLI)
- Executes the instructions requested by the user (commands)

#### Files

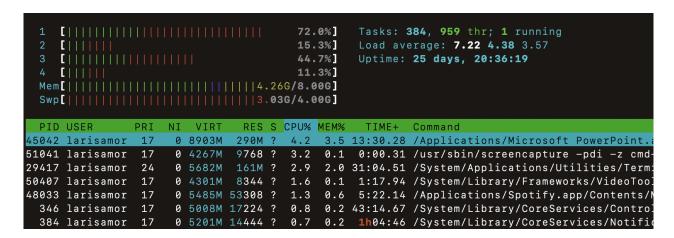
AirDrop			
(%) / III DIOP	HG002.g.vcf	<b>②</b>	20 January 2021, 11:45
Recents	HG002.g.vcf.idx	•	20 January 2021, 11:45
🙏 Applications	HG002.new.bai	•	20 January 2021, 11:30
	HG002.new.bam	<b>©</b>	20 January 2021, 11:30
OneDrive	HG002.old.bai	•	20 January 2021, 11:29
Desktop	HG002.old.bam	•	20 January 2021, 11:29
Documents	HG002.sorted.bam	<b>©</b>	20 January 2021, 11:01
	HG002.sorted.bam.bai	•	20 January 2021, 11:04
Downloads	HG002.sorted.dup.bai	0	26 January 2021, 9:36
Pictures	HG002.sorted.dup.bam	•	26 January 2021, 9:36
	HG002.sorted.dup.recal.bai	•	20 January 2021, 11:45
<b>arisamorales</b>	HG002.sorted.dup.recal.bam	•	20 January 2021, 11:45
iCloud	HG002.sorted.metric.insertSize.hist.pdf	<b>©</b>	20 January 2021, 11:29

```
total 132464
-rw-r--r--@ 1 larisamorales
                                        75629 Jan 20
                              staff
                                                       2021 HG002.g.vcf
-rw-r--r--@ 1 larisamorales
                                                       2021 HG002.g.vcf.idx
                              staff
                                        44048 Jan 20
-rw-r--r--@ 1 larisamorales
                                                       2021 HG002.new.bai
                              staff
                                        21584 Jan 20
                                        78169 Jan 20
-rw-r--r--@ 1 larisamorales
                              staff
                                                       2021 HG002.new.bam
-rw-r--r--0 1 larisamorales
                              staff
                                        21584 Jan 20
                                                       2021 HG002.old.bai
-rw-r--r--@ 1 larisamorales
                              staff
                                        76424 Jan 20
                                                       2021 HG002.old.bam
-rw-r----0 1 larisamorales
                              staff
                                     17803945 Jan 20
                                                       2021 HG002.sorted.bam
            1 larisamorales
                                      1350440 Jan 20
                              staff
                                                       2021 HG002.sorted.bam.bai
                                                       2021 HG002.sorted.dup.bai
-rw-r--r--0 1 larisamorales
                                      1350440 Jan 26
                              staff
-rw-r--r--@ 1 larisamorales
                              staff
                                     17956229 Jan 26
                                                       2021 HG002.sorted.dup.bam
-rw-r--r--0 1 larisamorales
                                      1350440 Jan 20
                              staff
                                                       2021 HG002.sorted.dup.recal.bai
-rw-r--r--@ 1 larisamorales
                              staff
                                     24793574 Jan 20
                                                       2021 HG002.sorted.dup.recal.bam
rw-r--r--@ 1 larisamorales
                                                       2021 HG002.sorted.metric.insertSize.hist.pdf
                              staff
                                        32357 Jan 20
```



#### **Processes**

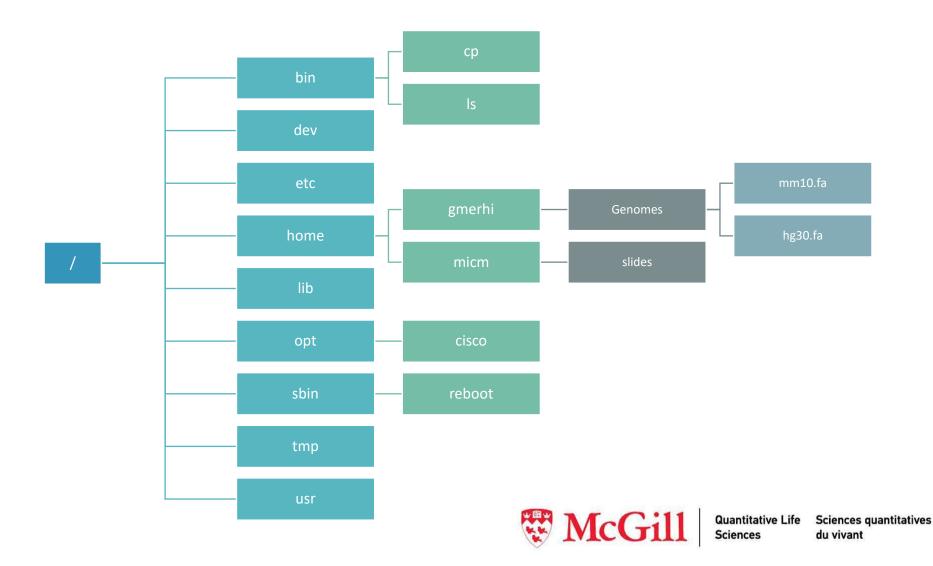
Р	Process Name v	% CPU	CPU Time	Threads	Idle Wake Ups	% GPU	GPU Time	PID	User
ZO	om.us	1.7	52:25.01	26	262	0.0	0.00	47243	larisamorales
Хр	protectService	0.0	0.08	2	0	0.0	0.00	50628	root
Хр	protectService	0.0	0.06	2	0	0.0	0.00	50596	larisamorales
xar	rtstorageremoted	0.0	2.62	2	0	0.0	0.00	235	root
Wi	irelessRadioManagerd	0.0	0.13	2		0.0	0.00	50460	root
Wi	indowServer	24.6	12:53:27.49	14	46	2.1	3:43:30.81	127	_windowserve
wif	fianalyticsd	0.0	2.60	2	0	0.0	0.00	293	root
Wi	i-Fi	0.0	5:28.75	4	0	0.0	0.00	369	larisamorales
System:		11.10%	6	CPU LOAD		Threads:			1,882
User:		14.32%	6			Proce	esses:		384
		74.58%	6						



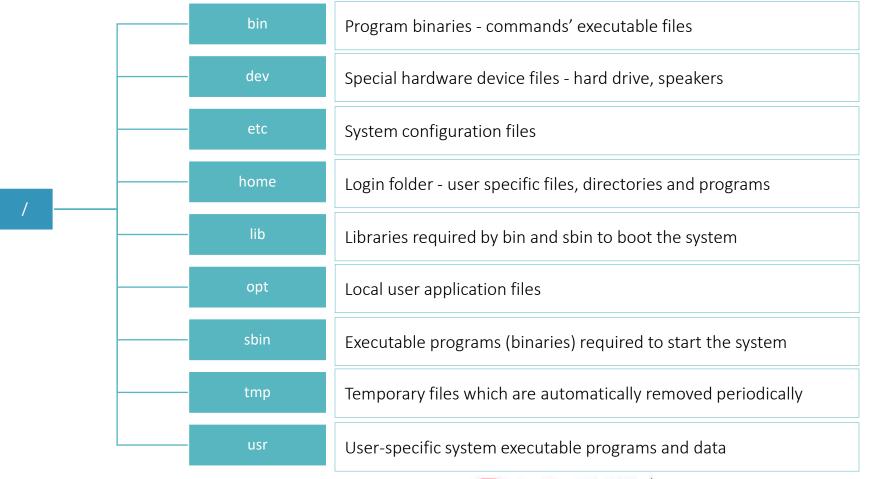


## Everything is a **file** or a **process**

# File system structure

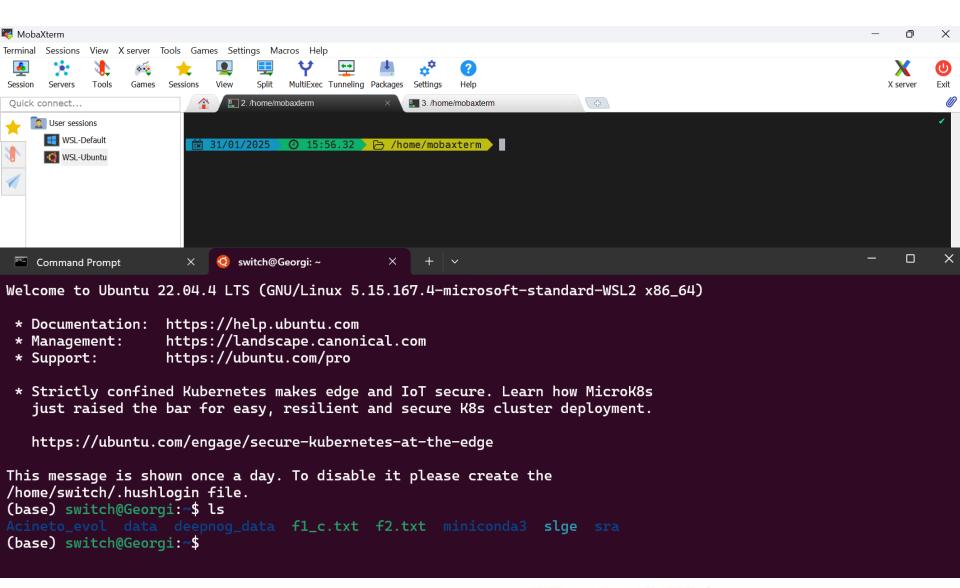


## File system structure





#### The terminal



Quantitative Life

Sciences

Sciences quantitatives

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## Keyboard shortcuts

Ctrl+A

Ctrl+E

Ctrl+C

q

Go to the start of the line

Go to the end of the line

Stop the current process (less, more, etc)

#### Unix commands

Programs built in the shell that perform specific actions



#### Basic commands

top

man

history

clear

See active processes and the resources they're using

Shows the manual page of a command

List your previous commands

Clear your terminal window

% htop

% man ls % man cd % man htop

% history

% clear

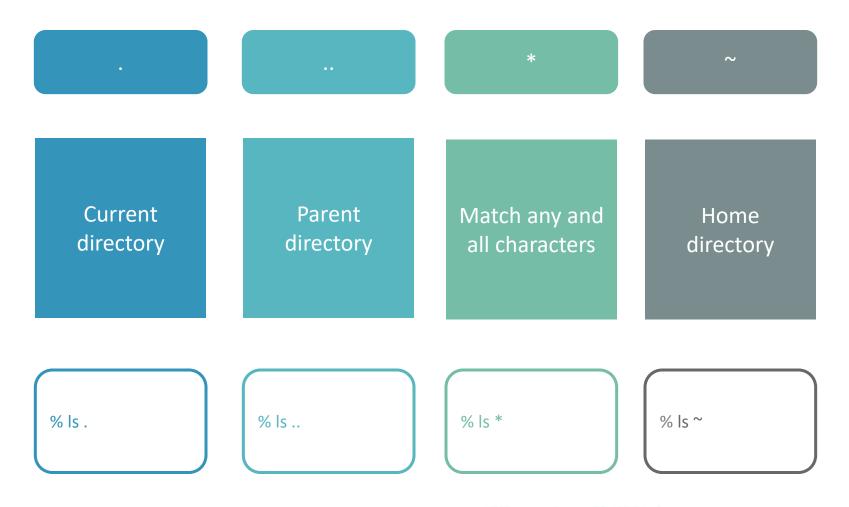


#### Basic commands

whoami Is pwd cd Shows the Prints the Displays Access a current current current user id directory directory directory % whoami % pwd % Is folder1 % cd folder1

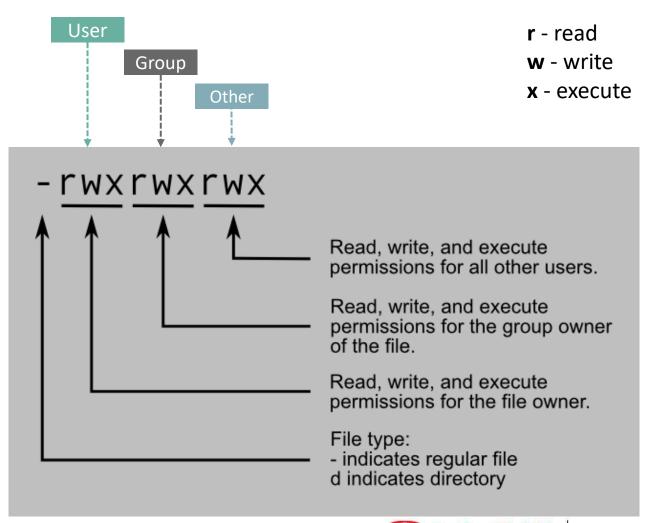


## Special characters





## File permissions



## File management commands

touch

mkdir

chmod

echo

Creates a new file

Creates a new directory

Change file permissions

chmod [ugo][+-][rwx]

u: user +: grant w: write g: group -: revoke r: read o: other x: execute Prints something to the terminal

% touch f1.txt

% mkdir ~/intro\_unix

% chmod o-w f1.txt

% echo "hello world"



## File management commands

ср

mv

cat

zcat

Copy a file

Move a file or rename it

Print the contents of a file(s) to the terminal

Print the contents of a zipped file(s) to the terminal

% cp f1.txt f1\_copy.txt

% mv f1\_copy.txt f1.txt

% cat f1.txt

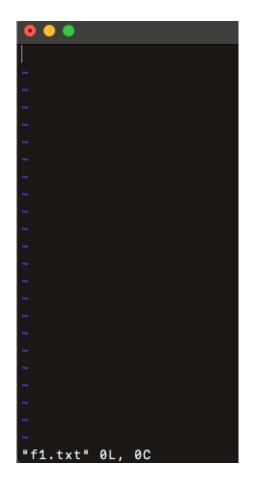
% zcat f1.txt.gz



### How to edit files in the terminal

Type :xq then press Enter

vi nano Recommended Recommended for code for plain text files/scripts files To open % nano f1.txt % vi f1.txt the file Press Ctrl+X Press Esc then Type Y hold To close **Press Enter** Shift+zz the file





#### How to download a file and use it in the terminal

- % cd ~/intro\_unix
- % mv ~/Downloads/cars.tsv data/ho1
- scp
   [username]@[IP\_address\_of\_the\_server]:[path/of/files/on/server] [path/on/your/machine]

## File management commands

gzip gunzip tar rm **Bundle files** Decompress a with Compress a file Removes file(s) file compression (optional) % tar -cvzf % gzip cars.tsv % gunzip \* % rm cars.tsv cars.tgz f\*



## File management commands

Print the first N lines of a file

Print the last N lines of a file

Print the last N of a file

Count words, characters lines or bytes

% head -3 cars.tsv

% tail -3 cars.tsv

% more cars.tsv

% wc cars.tsv



## Text processing commands

cut

sort

uniq

paste

Extract columns of file

Order elements

Get set of uniq elements

Paste two files column-wise

% cut -f1 cars.csv > col1.tsv

% sort cars.tsv > cars.sort.tsv

% uniq cars.sort.tsv

% paste col1.txt cars.tsv





## Bonus skill



## Create your own commands





Hands on 1



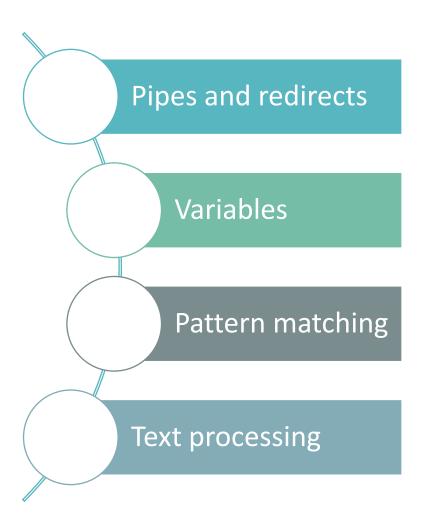
- 1. Print the current directory and user
- 2. Create a directory named intro\_unix in the home directory
- 3. Create a directory called data within intro\_unix
- 4. Create a sub-directories ho1 ho2 and ho3
- 5. Create a directory called folder1 under data/ho1
- 6. Go into folder1 and create two files: f1.txt and .f2.txt
- 7. Write the numbers from 1 to 10 in f1.txt (one number per line)
- 8. Write the following sequence in .f2.txt (one letter per line)
  - aaabbbbcccccddddddd

- 1. List all the contents of the directory (including .f2.txt) hint: look at the manual of the **ls** command
- 2. Change the name of .f2.txt to f2.txt
- 3. Change the permissions of f1.txt so that only the user can read and write the file
- 4. Write only the first 10 lines of f1.txt and all the lines in f2.txt to a new file called f3.txt

#### **Bonus skill**

- 1. Using the file cars.tsv:
  - 1. List the unique set of countries included in the dataset
  - 2. Find the number of unique car models

### Module 2



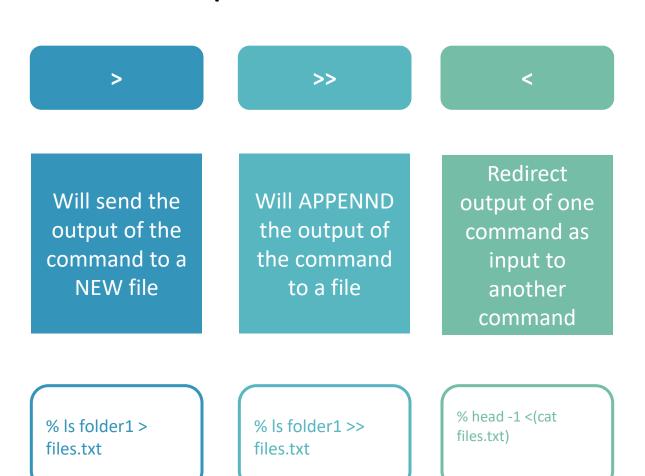


## Pipes

- A way to connect the end of something with the start of something else
- They are specified with the control operator " | "

% cat cars.tsv | head -10

## Redirect output



#### Variables

- Used to store information
  - number, text, file name, etc
- The name can only contain:
  - Letters (a to z or A to Z)
  - Numbers (0 to 9) \*not at the beginning
  - Underscore ("\_")
- We can have several types:
  - Environment variables
  - Shell variables
  - Global and local variables

#### Global variable

- Are variables that are accessible from any part of a script, within functions and sourced scripts.
- They do not disappear when the process is done but when the shell session is closed

```
% MY_NAME_LOCAL=I_am_groot
```

% echo \$MY\_NAME\_LOCAL

## Shell variables

 Environment variables required by the shell to function

\$USER

\$HOME

SPATH

\$PWD

Stores the name of the current user

Has the path to the user's home Contains all the directories where executable files are stored

Contains the path of the current directory



## Environment variables

Are available to any process in the shell

% export MY\_NAME=Larisa

% echo \$MY NAME

# Variable expansions

- Substituting the variable with its value
- It occurs before calling a command
- If the result contains spaces, the expansion should be quoted
- We can use operators to modify the content of the variable before passing it on to a command

# Variable expansions

Allow us to modify the content of the variable before passing it to a command

Extracting a part of the string

\${VAR:offset}

Extracts the content starting from the offset

\${VAR%pattern}

Extracts the content until it finds the pattern

# Variable expansions

Replacing substrings

\${VAR/pattern/replace}

Changes **pattern** for **replace** the first time it appears

\${VAR//pattern/replace}

Changes **pattern** for **replace** all the times it appears

\${VAR%pattern}replace

Extracts content until **pattern** and writes **replace** after

# Groupings

(...)

{...}

((...))

#### Subshell

Contains a list of commands. If preceded by \$(...) then its expanded to the output of the commands

Group commands and affect their parsing. When written as \${VAR} it is a parameter expansion Arithmetic instructions
They operate at the level of numbers.
The expansion is also \$((...))

files=\$(ls \*xt)

x=2 {x=4} echo \$x x=2 (x=4) echo \$x

x=2 y=6 echo \$((\$x+\$y))



# Example

#### Extract

```
% FILE_NAME=$(Is ../data/ho1 | tail -1)
% echo ${FILE_NAME:4}
% SUFFIX=${FILE_NAME:4}
```

### Replace

```
% echo ${FILE_NAME%${SUFFIX}}.csv
% echo ${FILE_NAME%_*}.csv
```

# Pattern matching

grep

Search a pattern in a file

% grep Male happiness.csv Country Gender Mean N=

AT **Male** 7.3 471

AT Female 7.3 570

AT Both 7.3 1041

BE **Male** 7.8 468

BE Female 7.8 542

BE Both 7.8 1010

BG **Male** 5.8 416

**BG Female 5.8 555** 

BG Both 5.8 971

# Regular expressions

#### **Groups and ranges**

	Any character except new line (\n)
(a b)	a or b
()	Group
(?:)	Passive (non-capturing) group
[abc]	Range (a or b or c)
[^abc]	Not (a or b or c)
[a-q]	Lower case letter from a to q
[A-Q]	Upper case letter from A to Q
[0-7]	Digit from 0 to 7
\x	Group/subpattern number "x"

#### **Character classes**

\s	White space
\S	Not white space
\d	Digit
\D	Not digit
\w	Word
\W	Not word



# Regular expressions

#### **Anchors**

٨	Start of string, or start of line in multi-line pattern
\A	Start of string
\$	End of string, or end of line in multi- line pattern
١Z	End of string
\b	Word boundary
\B	Not word boundary
\<	Start of word
\>	End of word

#### Quantifiers

*	0 or more	{3}	Exactly 3
+	1 or more	{3,}	3 or more
?	0 or 1	{3,5}	3, 4 or 5

#### **Special characters**

٨	[		\$	
{	*	(	\	
+	)	1	?	
<	>			
The escape character is usually \				



# Example

All lines that start with a letter

```
% grep -E "^[A-Z]+" happiness.csv
```

Lines that do not start with a letter

```
% grep -v -E "^[A-Z]+" happiness.csv
% grep -E "^\W" happiness.csv
```

Lines with a country from the list

```
% grep -f countries.txt happiness.complete.tsv
```



## Bonus skills



## awk

Scripting language that provides much more flexibility for text processing

#### **Built-in functions**

- length(string)
- tolower(string)
- toupper(string)
- match(string,pattern)

#### **Built-in variables**

- Entire line \$0
- Fields (specified by delimiter) \$1,\$2,...

# Example

Print the 3rd column and then the 1st column

```
% awk '{print $3 "\t" $1}' happiness.complete.txt
```

• Print columns 1,3 and 2 if they contain the word "Female"

```
% awk '/Female/ {print $1 "\t" $3 "\t" $2}' happiness.complete.txt
```

Count the number of characters in "Female" entries only

```
% awk -F "," '/Female/ { print length($0) "\t" $1 "\t" $2}' happiness.complete.csv
```



## sed

Command or streamline editor with multiple text processing functionalities

#### **Built-in functions**

- s/search/replace/ for pattern substitutions
  - /g replace all occurrences
  - /1,/2,... specifying which occurrence to replace
  - /I Ignore case
  - /w write to a file with /w filename
- -e to run multiple commands
  - sed -e 's/a/A/' -e 's/b/B/'

# Examples

Delete the first line

```
% Is -I | sed 1d
```

Replace capital A for lowercase a

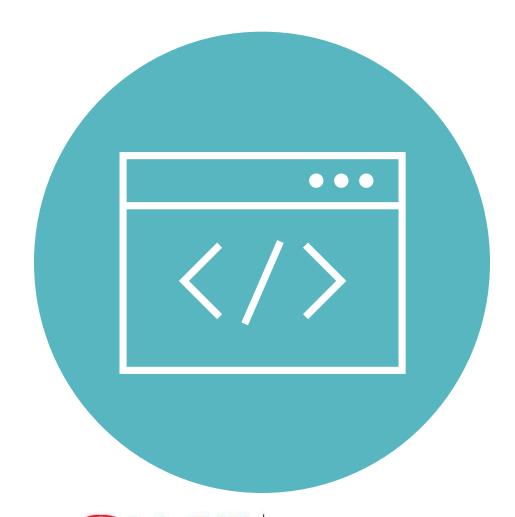
```
% sed 's/A/a/g' happiness.complete.txt | head
```

Print the first line every 3 lines

```
% awk 'NR % 3 == 0' happiness.csv
```



Hands on 2.1

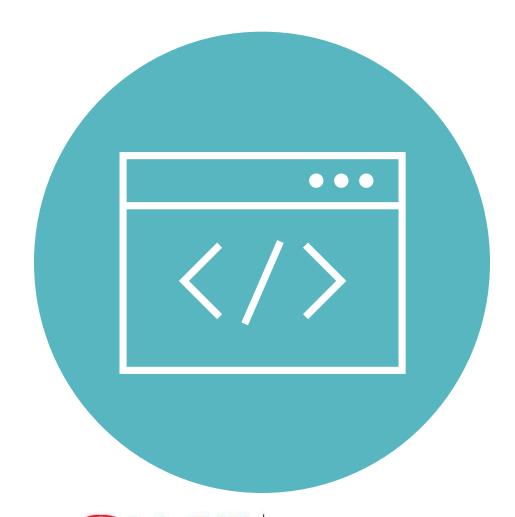




- 1. Go to the directory data/ho2 and list the files in it (move files there if missing)
- Count many "Female" and "Male" entries are listed in the file happinhowess.complete.tsv
- 3. Count how many unique countries are listed in the file happiness.complete.tsv



# Hands on 2.2





- 1. Go to the directory data/ho2 and list the files in it
- 2. Using the file happiness.complete.tsv
  - Write a new file in which all the countries are written in lowercase
  - b. Replace "Female" for "F", "Male for "M" and "Both" for B and look at the first 10 lines
  - c. Write a new file with all the "**Both**" entries only for the countries included the file countries.txt

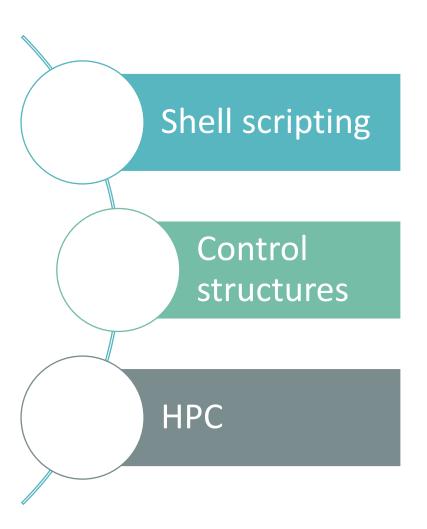
# What will happen?

```
% export MY_NAME=LARISA
% MY_NAME_LOCAL=larisa
% vi my_name.sh
```

```
#!/bin/sh
echo "My name is ${MY_NAME}"
echo "My name is ${MY_NAME_LOCAL}"
```

```
% chmod u+x my_name.sh
% ./my_name.sh
```

## Module 3





# For loop

for element in [list]
do
[COMMANDS]
done

- This loop iterates over all the elements in the list
- Executes some command until the list is over

#### Numbers smaller than 10

```
for i in {1..10}
do
echo number $i
done
```

#### **Change file extension**

```
for file in *tsv
do
  cp "$file" "${file%.tsv}.txt"
done
```



## Conditional statements

```
if [TEST]
then
[COMMANDS]
fi
```

If the output of the test is *True* then the commands are executed. If it's false, nothing happens.

```
if [TEST]
then
  [COMMANDS1]
else
  [COMMANDS2]
fi
```

If the test is *True* then executed commands1 If it's *False* then executes commands2.

```
if [TEST1]
then
[COMMANDS1]
elif [TEST2]
then
[COMMANDS2]
else
[COMMANDS2]
fi
```

If test1 is *True* it executes commands1. If it is *False* then evaluates test2. If test2 is *True*, executes commands2. If it is false, executes commands3

## Operators for variables

-n VAR

True if the length of VAR is greater than zero

-z VAR

True if the length of VAR is zero

-v VAR

True if the variable VAR has been assigned a value

S1 == S2

True if the strings S1 and S2 are equal

S1 != S2

True if the strings S1 and S2 are not equal

S1 < S2

*True* if string **S1** sorts before **S2** 

S1 > S2

True if string S1 sorts after S2



## Operators for arithmetic operations

INT1 –eq INT2

*True* if INT1 is equal to INT2

INT1 –gt INT2

*True* if INT1 is greater than INT2

INT1 –lt INT2

True if INT1 is less than INT2

INT1 –ge INT2

*True* if INT1 is equal or greater than INT2

INT1 –le INT2

True if INT1 is equal or less than INT2

<sup>\*</sup>Note: For arithmetic operations the statement need to be inside double square braces:



## Operators for files

-f FILE

True if FILE exists and is a regular file

-d FILE

*True* if FILE exists and is a directory

-e FILE

True if FILE exists

-r FILE

True if FILE exists and is readable

-w FILE

True if FILE exists and is writeable

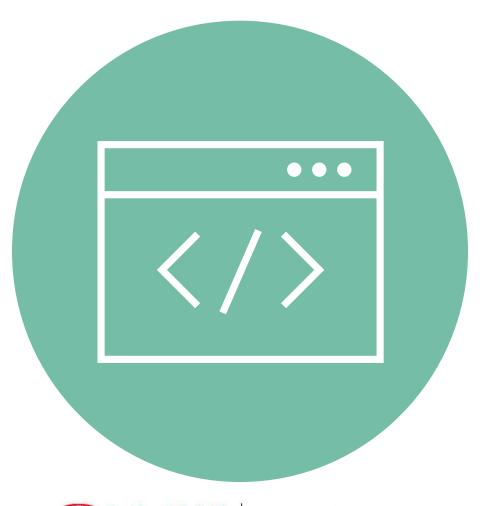
-s FILE

True if FILE exists and has a size greater than zero





Hands on 3.1 and solution







### **Counting and Analyzing Car Manufacturers**

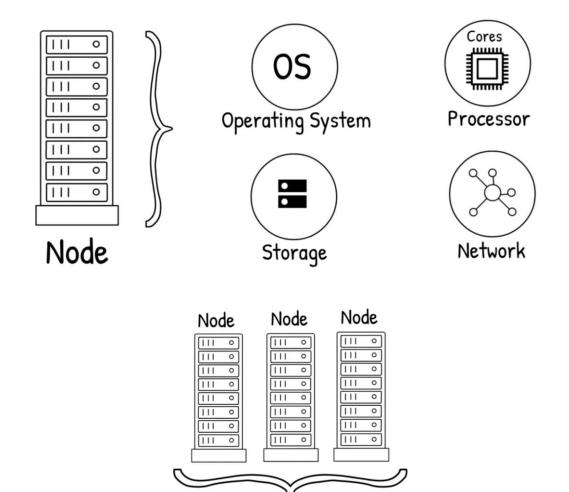
- You have a TSV file named cars.tsv in your ho1 current directory.
- The file contains data about cars, with at least the following columns: ID, Manufacturer, Model, Year.

### **Requirements:**

- Write a Bash script to process the cars.tsv file and:
  - Count the total number of cars (excluding the header).
  - Count the number of cars for each manufacturer (e.g., Toyota, Honda, Ford).
  - Calculate the percentage of cars for each manufacturer.
  - Print the results in a readable format.



# High performance computing



Cluster

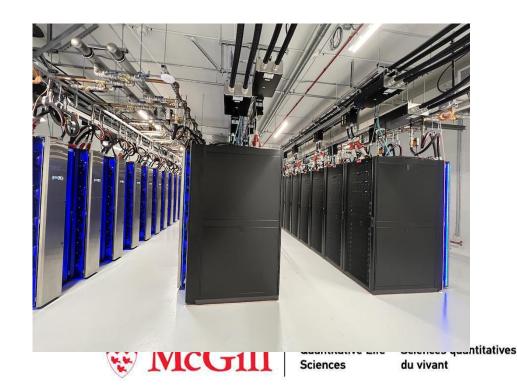
# Digital Research Alliance of Canada (Compute Canada)



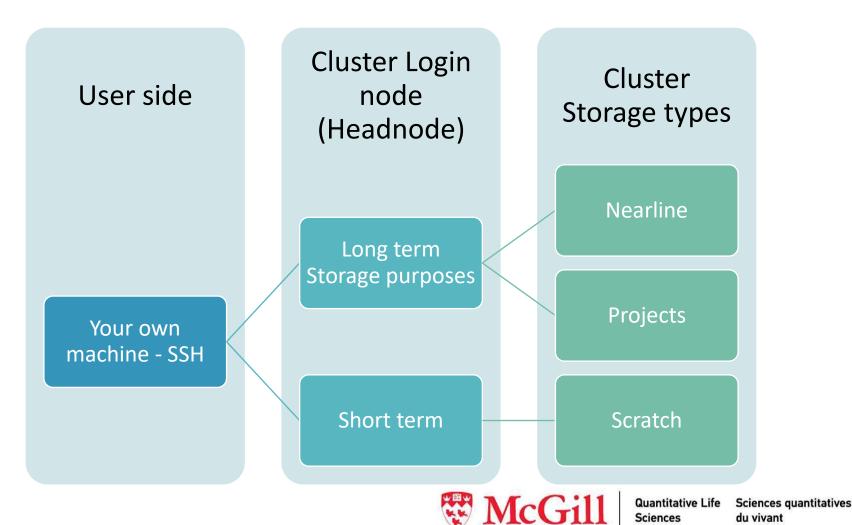
**Digital Research Alliance** of Canada

Alliance de recherche numérique du Canada





# Digital Research Alliance of Canada (Compute Canada)



du vivant

# Digital Research Alliance of Canada (Compute Canada)

Feature	Scratch Storage	Nearline Storage	Projects Storage
Purpose	Temporary, high-performance storage	Long-term, infrequently accessed data	Persistent, project-related storage
Performance	High	Moderate to Low	Moderate to High
Capacity	Large	Large	Moderate to Large
Backup	No	Yes	Yes
Purge Policy	Regular purges	Data retained long-term	Retained throughout project duration
Accessibility	Individual user	Individual user	Shared among project members

# Digital Research Alliance of Canada (Compute Canada)

- Overview of some important terms related to the cluster:
  - Compute Nodes → one node = one computer
  - Job Scheduling System → SLURM
  - SLURM job priority → Fairshare

## Connect to a cluster

ssh user@cluster.computecanada.ca

```
[(base) larisamorales@MBP-de-Larisa ~ % ssh lmoral7@graham.computecanada.ca
Last login: Wed Sep 15 11:32:04 2021 from modemcable076.215-203-24.mc.videotron.ca
***********************************
Welcome to the ComputeCanada/SHARCNET cluster Graham.
Documentation: https://docs.computecanada.ca/wiki/Graham
Current issues: https://status.computecanada.ca/
      Support: support@computecanada.ca
Graham has several types of GPUs, some of which are available with less wait:
320 p100 2/node, 12GB, original
 70 v100 8/node, 16GB, newer, about 50% faster than P100 and with tensor cores
144 t4 4/node, 16GB, newer, about half a V100 for compute & AI, except much slower FP64
More details: https://docs.computecanada.ca/wiki/Graham#GPUs on Graham
```

## Jobs

[georgi26@narvall scratch]\$ cat tarring.sh

Requesting a computing allocation to execute your code

```
#!/bin/bash
#SBATCH --job-name=tarball_creation_GM
#SBATCH --account=ctb-shapiro
#SBATCH --nodes=1
#SBATCH --repus-per-task=16
#SBATCH --mail-type=ALL
#SBATCH --mail-user=georgi.merhi@mail.mcgill.ca
#SBATCH --output=tarball_creation_%j.out
#SBATCH --error=tarball_creation_%j.err
#SBATCH --time=23:00:00
#SBATCH --tme=86

# Navigate to the directory you want to archive
cd /home/georgi26/scratch/
# Create a tar.gz archive of everything in the directory
tar -czvf Georgi_all_13MAI2024_v2.tar.gz *
```

# Job arrays

A job script that internally behaves like a for loop and submits individual jobs

```
[georgi26@narval1 phylo_RAW]$ cat phylo_snp_2.sh
#!/bin/sh
#SBATCH --account=ctb-shapiro
#SBATCH -o /home/georgi26/scratch/phylo_RAW/output_e_o/gatk_call_variants_%j.o
#SBATCH -e /home/georgi26/scratch/phylo_RAW/output_e_o/gatk_call_variants_%j.e
#SBATCH -t 08:00:00
#SBATCH --mem=64G
#SBATCH -c 8
#SBATCH --job-name=GM_gatk_variant_call
#SBATCH --array=1000-1273 #Indicate number of isolates to call variants from.
## Calling SNPs using GATK HaplotypeCaller for phylogeny construction
# Load packages
module load StdEnv/2020 gcc/9.3.0
module load bwa
module load samtools
module load java picard
module load gatk
# ID of Reads for analysis
RUNID=$(sed -n "$SLURM_ARRAY_TASK_ID"p $RUNDIR/genomes_all_list.txt)
```

## Jobs - Bonus

You can use the sbatch command with flags without adding parameter directives in a job script.

#!/bin/sh

module load iq-tree iqtree2 -s ./core\_gene\_alignment\_Acinetobacter\_GM.aln -T 32 -m MFP -B 1000 echo done>run.done

Then run the command in the terminal as follows

sbatch -A ctb-shapiro --time 168:0:0 --tasks 32 --mem 800g run.sh



# Job submission/monitoring & Useful commands

sbatch job script.slurm

Submit a job using a job script

squeue -u \${USER}

List all the processes of \$USER

sq

Same as above but shorter

scancel [PID]

Cancel a process using its process ID (PID)

scancel -u \${USER}

Cancel all processes that belong to \$USER

quota -s

Check your home directory usage and quota

quota --per user

Check all users' usage and quota

## To summarize

- ✓ UNIX is an operating system with two parts: the kernel and the shell
- ✓ The shell (terminal) contains many built-in programs called commands
- ✓ Commands allow the user to perform operations on files
- ✓ HPC can help solve problems that go beyond the capacity of personal computers

#### Now you are ready to:

- Automate file management operations
- Extract relevant information from files without opening them
- Create pipelines that connect multiple tools
- Determine when a problem could be better solved with HPC

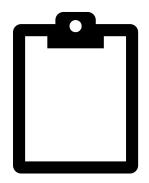
# Thank you for attending!





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### Datasets references

Hands on 1 – Cars dataset

https://perso.telecom-paristech.fr/eagan/class/igr204/datasets

Hands on 2 – Happiness surveys dataset

https://perso.telecom-paristech.fr/eagan/class/igr204/datasets

#### Resources

**UNIX** cheatsheet

