



Informatics on High-throughput Sequencing Data

(Summer Course 2020)

Day 6



Agenda

- ▶ Unix-based systems.
- ▶ Why Linux!
- ▶ Let's start!
- ▶ Linux Commands for:
 - Files & Directories.
 - System.
 - Process Management.
 - Networking.
 - Compression.
 - Searching.
- ▶ Piping output.
- ▶ Wildcard character.
- ▶ Redirecting output.
- ▶ Stream Editor (**Sed**).
- ▶ Linux tools for text files processing.
- ▶ Shell Scripting

Piping output

- ▶ Pipes are represented by the **|** character.
- ▶ It is possible to send the output of one program to another program as input.

history | less List all remembered commands page by page.

history | grep Bio
List all remembered commands containing string “Bio”.

Redirecting output

```
ls -l > myfile.txt
```

List items in current directory and show in long format to see permissions, size, and modification date. The output is directed to display in `myfile.txt`.

```
ls -l >> myfile.txt
```



More Unix commands

```
wc -l
```

```
cut -f1,2,3,4 athal.genes.gtf > sara.gtf
```

```
Cut -d ',' -f1 GDC.CSV
```

```
sort sara.gtf
```

```
sort -r sara.gtf
```

```
sort -k4 sara.gtf
```

```
sort -k4 numbers.txt
```

```
sort -k 4n numbers.txt
```

More Unix commands

```
sort -k 4nr numbers.txt
```

```
sort -u numbers3.txt
```

```
sort -k2,3 numbers1.txt
```

```
sort -t ',' -k2r GDC.CSV
```

```
sort -n -k2,2 -k6,6 numbers1.txt
```

```
uniq numbers3.txt
```

```
uniq -c numbers3.txt
```



More Unix commands

```
diff kmers1.txt kmers2.txt
```

```
sort kmers1.txt > S1.txt
```

```
sort kmers2.txt > S2.txt
```

```
comm S1.txt S2.txt
```

```
comm -1 -2 S1.txt S2.txt | wc -l
```

Display text in terminal

```
echo 'Hello World'
```

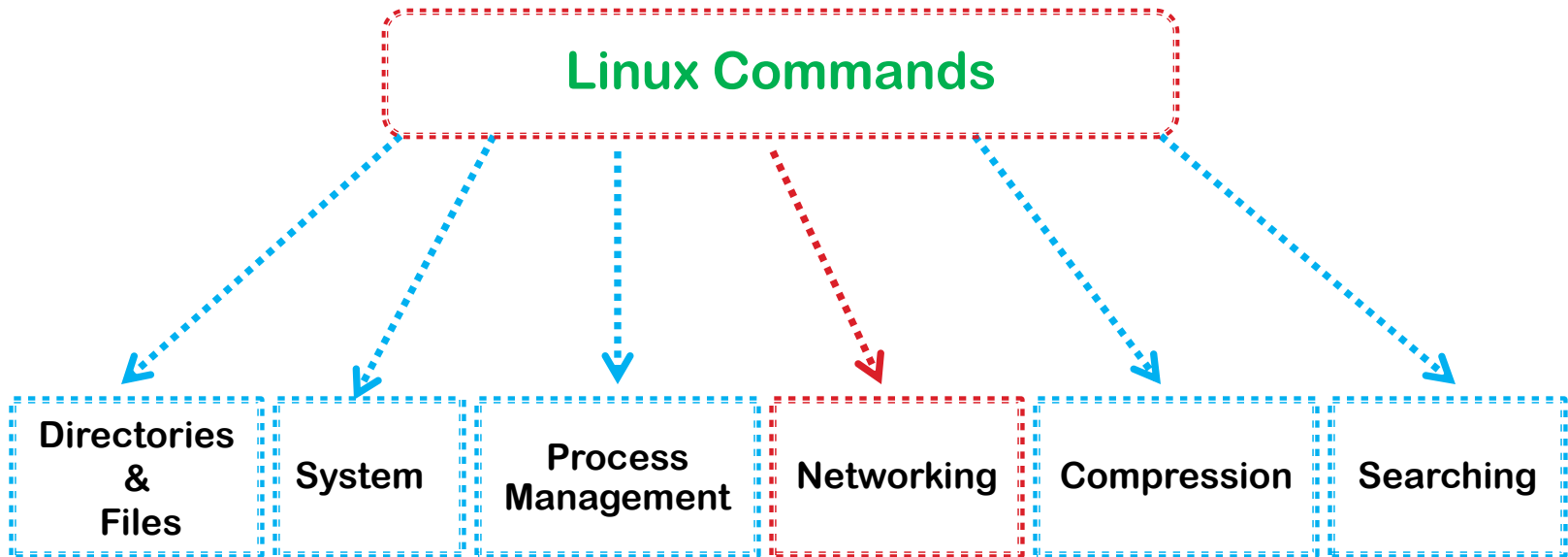
```
echo -e 'Hello \n World'
```

```
echo $( (1+2) )
```

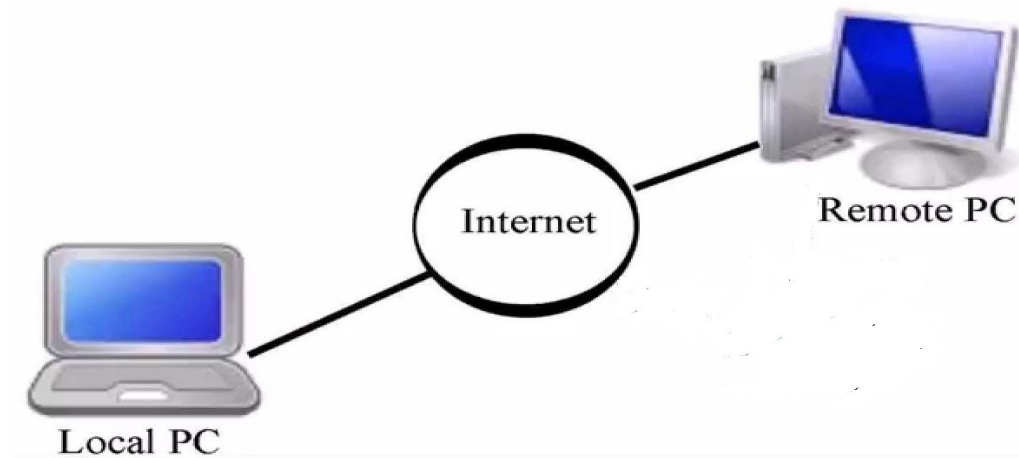

Editing files

- ▶ vi or vim program.
- ▶ press **i** for insert mode.
- ▶ press **Esc** to leave insert mode.
- ▶ type **:wq** and press **Enter** to save changes and quit.
- ▶ type **:q!** and press **Enter** to quit without saving.

Getting Started !!



Networking (Connect to a Remote Machine)

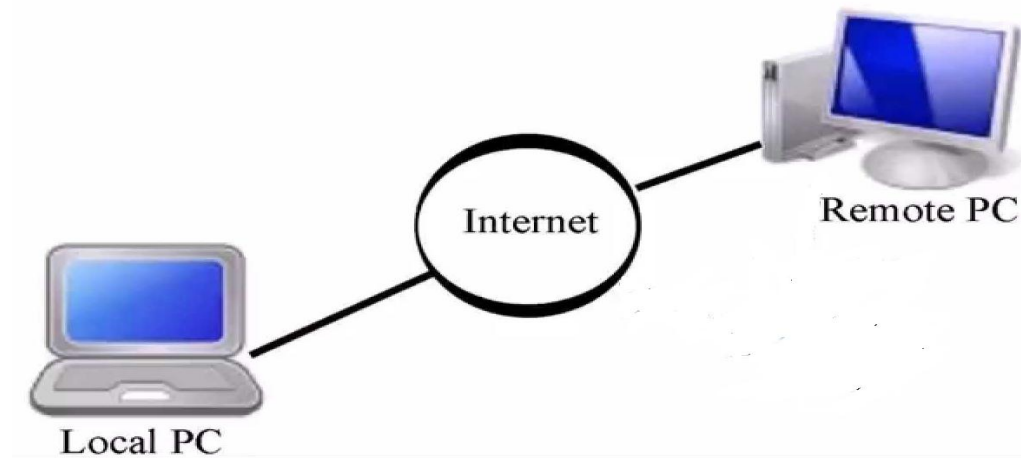


```
ssh username@ageri.sci.eg
```

"Securely connect me to a shell on a remote machine."

<https://phoenixnap.com/kb/ssh-to-connect-to-remote-server-linux-or-windows>

Networking (logging out of a remote machine)



exit or **Ctrl-d**

“close current terminal window.”

Networking (copy files (to | from) a remote machine)

```
scp username@ageri.sci.eg:file Bio
```

Secure copy a file from remote server to the Bio directory on your machine.

```
scp file username@ageri.sci.eg:Bio
```

secure copy a file from your machine to the Bio directory on a remote server.




Networking (copy files (to | from) a remote machine)

```
scp -r username@ageri.sci.eg:Bio Bio
```

Secure copy the directory **Bio** from remote server to the directory **Bio** on your machine.

```
scp username@ageri.sci.eg:~/myfile.txt ./
```



Your home directory on the remote system.



Current directory in your computer.

Networking

```
wget file_link
```

Download a file.

```
curl file_link
```

Download a file.

```
sudo apt-get install mailutils
```

Download and install packages.

Networking

sudo apt-get install mailutils

```
toshiba@ubuntu:~/Documents/agri_training$ mail
The program 'mail' is currently not installed. You can install it by typing:
sudo apt-get install mailutils
toshiba@ubuntu:~/Documents/agri_training$ sudo apt-get install mailutils
[sudo] password for toshiba:
Reading package lists... Done
Building dependency tree
```


Shell Scripting

- ▶ **SHELL SCRIPTING** is writing a series of commands for the shell to execute.
- ▶ It can combine lengthy and repetitive sequences of commands into a single and simple script, which can be stored and executed anytime.
- ▶ This reduces the effort required by the end user.

<https://www.guru99.com/introduction-to-shell-scripting.html>



References

- ▶ Paul Stothard, An Introduction to Linux for bioinformatics , 2016.
- ▶ Robert Bukowski, Linux for Biologists- Part 1.
- ▶ Steve Pederson, Introduction To Linux/Ubuntu & Shell Scripting, 2014.
- ▶ <https://bioinformatics.uconn.edu/unix-basics/#>
- ▶ <https://learn.gencore.bio.nyu.edu/ngs-file-formats/quality-scores/>
- ▶ <https://coding4medicine.com/Members/pages/home/>
- ▶ <https://open.oregonstate.education/computationalbiology/chapter/patterns-regular-expressions/>
- ▶ <https://bioinformaticsworkbook.org/Appendix/Unix/unix-basics-3grep.html#gsc.tab=0>
- ▶ <https://datacarpentry.org/shell-genomics/04-redirection/>

Thanks!

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