# Bacterial WGS training: Exercise 1

| Title             | Linux command line.  |
|-------------------|--|
| Training dataset: | None   |
| Questions:        | <ul> <li>How do I use the command line?</li> <li>How do I navigate the file system?</li> </ul>   |
| Objectives:       | <ul> <li>Learn/Remember how to use the command line.</li> <li>Learn/Remember how to navigate through the Linux file system.</li> </ul> |
| Time estimation:  | 30 min   |
| Key points:       | Remeber the shell basic commands: pwd cd Is mkdir rmdir mv nano cat less more head tail rm   |

## :bangbang: Important things to remnenber:

- Use Tab to automatically complete file names and paths, so it can be easiert to write in the terminal
- Use keyboard arrows (:arrow\_up: :arrow\_down:) to move through your terminal's history, so you don't have to write the commands again.
- Try not to use spaces, accents or special characters like "Ñ" letter, when writting directory of file names.
- Basic commands you should always remember: pwd cd ls mkdir mv rm rmdir less nano

# Answering to main questions

How do I use the command line?

Open a terminal by clicking in the icon or typing Ctrl+Alt+T. Now you can type in the prompt.

How do I navigate the file system?

Let's remember the basics: pwd cd ls mkdir mv rm rmdir less nano . We are going to use those commands to:

Checking the working directory (pwd)

Check our working directory:

pwd
#Output: /home/alumno

Moving between directories (cd)

Move to our Desktop folder:

cd ~/Escritorio

pwd

#Output: /home/alumno/Escritorio

Move to the course folder:

```
cd
pwd

#Output: /home/alumno
cd wgs
pwd

#Output: /home/alumno/wgs
cd RAW/FULL_DATA
pwd

#Output: /home/alumno/wgs/RAW/FULL_DATA
cd .
pwd

#Output: /home/alumno/wgs/RAW/FULL_DATA
cd .
pwd

#Output: /home/alumno/wgs/RAW/FULL_DATA
cd .
#Output: /home/alumno/wgs/RAW/FULL_DATA
cd ..
#Output: /home/alumno/wgs/RAW/
cd ../..
#Output: /home/alumno/
```

#### Questions:

- Which is the meaning of the "~" symbol?
- What does de cd command without arguments do?
- What does "." mean?
- What does ".." mean?

Listing directories (Is)

```
cd wgs
ls
#Output: ANALYSIS bacterial_wgs_training RAW REFERENCES RESULTS
```

This is the folder structure we will use for this training. Now we are going to list the files in the REFERENCE folder:

```
ls REFERENCES
```

This command will output a big list of files, which are the files that we will usea as REFERENCE through the different exercises of the training. Now wi will run this other command:

```
ls /home/alumno/wgs/REFERENCES/
```

#### Questions:

- Which is the difference between this las command and the previous one?
- Do they display the same information?
- Which one is relative path?
- Which one is absolute path?

Let's see different parameters for the ls command. Write:

```
ls REFERENCES
ls -l REFERENCES
ls -a REFERENCES
ls -la REFERENCES
```

## Questions:

- What does de different arguments of \(\begin{aligned} \subsection \\ \text{ls} \\ \does \end{aligned}
- What does the new file special?

 $\triangle$  **REMINDER:**  $\triangle$  EVERY TIME YOU CHANGE DIRECTORY (cd) YOU HAVE TO CHECK YOUR PATH (pwd) AND LIST THE FILES INSIDE (Is) TO CHECK YOU DIDN'T MAKE MISTAKES

Creating and removing directories (mkdir & rmdir)

Now we are going to move to the ANALYSIS folder which is the folder were we will run all the exercises

```
cd ANALYSIS
pwd
#Output: /home/alumno/wgs/ANALYSIS
ls
```

As you can see the folder is empty, so now we will fill this folder. Create a directory for this handson: **Remember:** Linux is case sensitive and does not like white spaces in names

```
mkdir 01-handsonLinux
ls
#Output: 01-handsonLinux
```

#### Now type:

```
mkdir 01-handsonlinux 01-HandsOnLinux
ls
#Output: 01-handsonlinux 01-handsonLinux 01-HandsOnLinux
```

### Questions:

- Is it possible to create more than one directory at the same time?
- If the names of the folders are the same, why it creates three different directories?

Now we will remove the extra directories:

```
rmdir 01-handsonLinux 01-HandsOnLinux
ls
#Output: 01-handsonlinux
```

Moving and renaming files (mv)

Move to the new folder

```
cd 01-handsonlinux
pwd
#Output: /home/alumno/wgs/ANALYSIS/01-handsonlinux
```

We are going to move the hidden file in REFERENCE folder to this directory and then rename it:

### Questions:

- Which is the difference between the two mv commands?
- Do you remember what "." mean from the first questions?
- And ".."?

## △ REMINDER: △ LINUX DOES NOT REQUIRE FILE EXTENSIONS

Editing files and displaying them (nano & cat)

We are going to read the file and edit it:

```
cat NowImNotHidden
#Output: I'm a hidden file.
```

This is not true, so we are going to edit it:

```
nano NowImNotHidden
```

Write: I'm not a hidden file.

And save it: Ctrl + O Intro

Close the new file: Ctrl + X

Now read the new file:

```
cat NowImNotHidden
#Output: I'm not a hidden file.
ls
#Output: NowImNotHidden
cd ../../
```

Read files other ways (less & more & head & tail)

And now we will read this file:

```
cat REFERENCES/bacterial_wgs_training_initial.tree
less REFERENCES/bacterial_wgs_training_initial.tree
#Remember: To close less press "q"
```

```
more REFERENCES/bacterial_wgs_training_initial.tree
#Remember: To close more press "q"
```

```
head REFERENCES/bacterial_wgs_training_initial.tree
tail REFERENCES/bacterial_wgs_training_initial.tree
```

```
head -n4 REFERENCES/bacterial_wgs_training_initial.tree
tail -n3 REFERENCES/bacterial_wgs_training_initial.tree
```

## Questions:

- Which is the difference between head and tail?
- What does the argument -nX do to tail and head?

Removing a file (rm)

Now we will learn how to remove files:

```
cd ANALYSIS/01-handsonlinux/
pwd
ls
#Output: NowImNotHidden
mv ../../REFERENCES/bacterial_wgs_training_initial.tree .
ls
#Output: bacterial_wgs_training_initial.tree NowImNotHidden
rm NowImNotHidden
ls
#Output: bacterial_wgs_training_initial.tree
```