Introduction to GNU Linux CLI

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What is CLI?

CLI stands for **C**ommand **L**ine **I**nterface, usually shortened to *command line* or, by historical reasons, into *terminal*. The former dates back to the days when **UNIX** ran on large mainframes accessed via remote terminals.

What is shell?

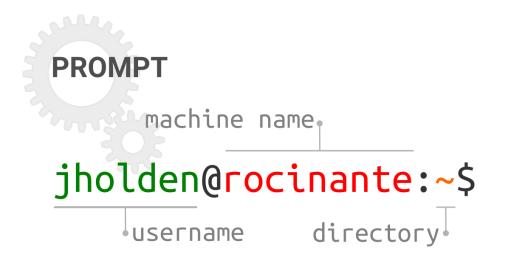
Shell is a program which serves as an interface between you and the **o**perating **s**ystem. It runs in your **terminal emulator** and allows you to interact with the *kernel* of your system.

There are *many* such programms, which can be qualified as *shells*: **GUI**, **sh**, **csh**, **ksh**, **zsh**, **c**, **bash**.

The **bash** shell is the most popular shell used across most of the Linux distributions.

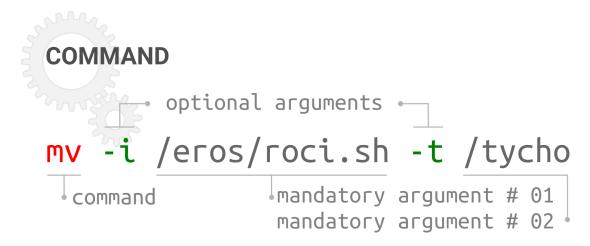
What is prompt?

The *prompt* is what you see after you've logged into the server. The prompt is called *prompt* because it *prompts* you for your *command*:



What is command?

Command is a single-purpose *tool*, a *program* which is designed to do a *single task*, and do it well.



Command accepts <u>zero or more</u> mandatory and <u>zero or more</u> optional arguments which allows you to get the desired result, and fine-tune the output of the command.

Echo

Intro

echo simply prints into the terminal window anything you give it as an *argument*:

```
echo Hello, bash!
echo echo
```

In the example above Hello, bash! and second echo are arguments the command echo evaluates.

Hands-on

Print a string O Romeo, Romeo! Wherefore art thou Romeo? into your screen.

Brackets expansion

Intro

A *brackets expansion* is a nice feauture of the bash which you might find very interesting. To use brackets expansion, separate the arguments you are going to *expand* by the comma, and enclose them to the curly brackets {}.

```
echo file_{a,b,c}.txt
echo file_{1..5}.txt
echo file_{1..5}{a,b,c}.txt
echo {a..z..2}
```

Hands-on

Find the way to print all odd numbers within the interval [1, 50]

Print working directory

Intro

pwd stands for **p**rint **w**orking **d**irectory. When you enter this command **bash** will print your *current or working directory*.

```
pwd
# /home/jholden/
```

Look at your prompt, you are probably seeing \sim (*tilde*) symbol instead of your home directory *path* I have just found with pwd. In Linux CLI \sim is a short-hand for the *home directory*.

Hands-on

Print your working directory into the terminal.

Moving between directories

Intro

cd command stands for **c**hange **d**irectory. With cd you can move from your *current* directory to any directory you want, if you provide a *path* to the target directory as an argument.

```
cd /data3  # change your working directory to data3
pwd  # print your working directory
```

Hands-on

- Move from your working directory to the /data3/ directory;
- Move back to your home directory using ~ instead of the full path to your home directory;

Relative vs absolute paths

Intro

Use . to shorthand your *current* directory, and . . to shorthand the *parent* directory of your current directory.

```
cd ./projects # Move to the 'projects' directory inside your working directory
cd ../rawdata # Move to the 'rawdata' which is located in the parent directory
```

Hands-on

Move one directory above (parent directory) from your current directory. Use pwd to veryfy your location, and move back to your home directory using ~ instead of the *full path*.

Listing the directories

Intro

ls stands for **list**. The command general format is ls [OPTION] ... [FILE]... By default, ls will list the content of your current directory.

- -a, --all do not ignore *hidden* files;
- -1 use a long listing format;
- -h human-readable (e.g. show file size in Kb or Mb);

```
ls -alh
```

Hands-on

List content of your home directory.

Making directories

Intro

To make a directory use mkdir command (stands for make directory). It accepts the path to the directory you are going to create, and several optional arguments.

```
mkdir sandbox  # create a new directory 'sandbox'
ls -lh  # list files and directories
cd sandbox  # change directory to 'sandbox'
pwd  # print working directory
mkdir -p ./data/in  # create subdirectory
mkdir -p ./src/r  # create subdirectory
tree  # explore the dir structure
```

Hands-on

Create a subdirectory sandbox with data subdirectory in in your home directory with one command.

Removing files and directories

Intro

The command rm is used for removing *files* or *directories*. To remove directories containing other files and directories, use -r or -R switch.

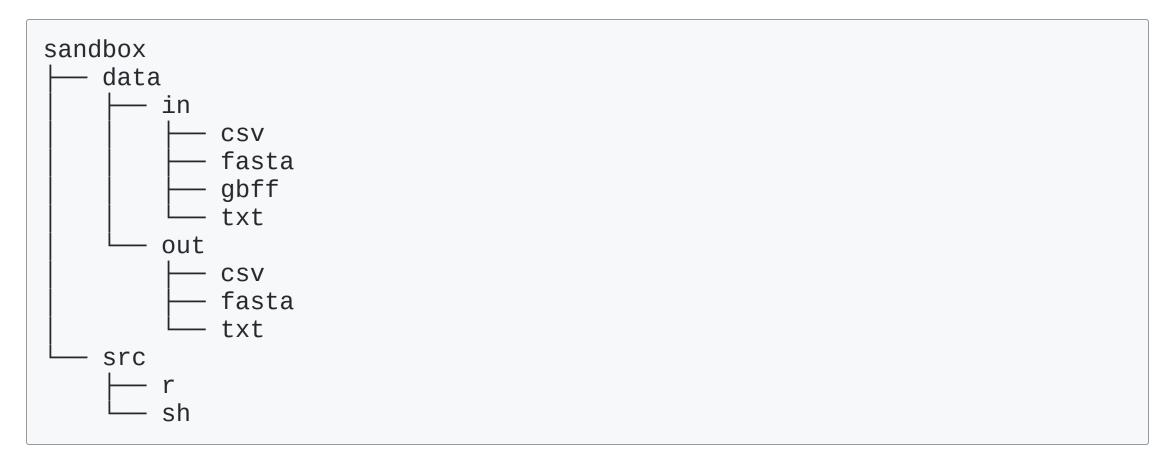
```
rm ./sandbox # Returns an error since directory is not empty rm -rv ./data/raw02 # Succesfully removes directory with its subdirectories
```

Above we use -v switch to make rm verbose and -r switch to recursively remove the content of the directory.

Hands-on

Remove whole sandbox directory and it's subdirectories.

Note about the project structure



More information: Vince Buffalo, 2015 and Noble, 2009.

Using brackets expansion with mkdir

Intro

We can use brackets expansion to create a above described directory structure with one command.

```
mkdir -pv sandbox/{src/{sh,r},data/{in/{csv,txt,fasta,gbff},out/{csv,txt,fasta}}}
```

Hands-on

Repeat the example above. After that create a subdirectory <code>img</code> in <code>./sandbox/data/out/</code> directory. <code>img</code> must contain subdirectories <code>png</code>, <code>pdf</code>, <code>svg</code>. Use <code>brackets expansion</code>.

Creating files

Intro

You can create an empty file with touch command. The command accept several optional arguments, and mandatory argument(s) - a file name(s) you are going to create:

touch logfile tree

Hands-on

It's usefull to have a ReadMe.md file in a root of your project directory, to document your work. Create a ReadMe.md file in a sandbox directory.

Copying files

Intro

cp stands for **c**o**p**y. It accepts the path to the file or directory you want to copy, the destination path, and several optional arguments (e.g. -R, -r for *recursive*, -t for *destination* path).

```
cp /home/dst20/cli00/sandbox/data/txt/ncbi_links.txt -t ./data/in/txt
```

Hands-on

Repeat the command above to copy the <code>ncbi_links.txt</code> file. But instead of copying it into your <code>./data/in/txt</code> directory, copy it into <code>./data/in/csv</code> directory. Verify that file has been copied succesfully.

Moving files

Intro

Use mv command (stands for move) for moving files and directories. This command accepts path(s) to the files (directories) you are going to move, and destination path(s) as well as several optional arguments.

```
mv -v ./data/in/csv/ncbi_links.txt -t ./data/in/txt
```

We use -v switch to make mv verbose, and -t switch to specify *target* directory.

Hands-on

You realized that you copied a .txt file into the wrong (.csv) directory. Use the example above to move ncbi_links.txt into the directory it belongs to. Don't forget to verify that the operation was sucessful.

Renaming files and directories

Intro

The fact that we can specify the name of our *destination file* allows us not ony to *move* but also to *rename* files and directories with mv:

```
mv -v file_with_old_name.txt file_with_new_name.txt
```

Notice that we use -v switch to make mv verbose.

Hands-on

```
Rename your ./data/in/txt/ncbi_links.txt file into ./data/in/txt/ncbi_pa_assemblies_gbff.txt using example above.
```

Reading the files with cat

Intro

The simplest command you can use for reading the content of the files is cat. Supply the cat with the path to your file, and read the content of the file on the screen.

```
cat ./data/in/txt/ncbi_pa_assemblies_gbff.txt
```

Hands-on

Using command above try to find out if all of the first ten links point to the .gbff.gz files.

Heads and tails

Intro

We can use head and tail commands to print the head and tail of the text file respectively. Commands accept few useful arguments for instance -n argument followed by the number of lines you want to print instead ten first (last) lines (default).

Hands-on

- Print only first 5 lines of ./data/in/txt/ncbi_pa_assemblies_gbff.txt file;
- Print only last 5 lines of ./data/in/txt/ncbi_pa_assemblies_gbff.txt file;

Pipes

Intro

What if want *to chain* commands? Can we *pipe* the output of one command into another command? We can use | known as a *pipe operator* for this purpose. Imagine we are interested only in *last 50 lines* of our

./data/in/txt/ncbi_pa_assemblies_gbff.txt file, however not all of them, but only first 3 lines?

```
tail ./data/in/txt/ncbi_pa_assemblies_gbff.txt -n 50 | head -n 3
```

Hands-on

Repeat the example avbove, but this time, try to find what are the first 2 lines of the last 123 lines of the ./data/in/txt/ncbi_pa_assemblies_gbff.txt file.

more on reading the files

Intro

Instead of cat we can use more to read text files (and more). The basic workaround is the same as for cat, however more accepts more optional arguments, and allows you to navigate through the text document in more comfortable way.

```
more -d ./data/in/txt/ncbi_pa_assemblies_gbff.txt
```

Hands-on

Repeat the command above to see more in action.

less on reading the files

Intro

The less is more! With tones of options, it allows you to scroll up and down through the document, and even to *search* a specific string: just hit /, type your string and hit <ENTER>. more will highlight all the patterns it found in the document, and you can jump from one match to another pushing the n key!

```
less ./data/in/txt/ncbi_pa_assemblies_gbff.txt
```

Hands-on

Browse the ./data/in/txt/ncbi_pa_assemblies_gbff.txt with more , and locate the link to the Pseudomonas_aeruginosa_PAKAF assembly.

To be continued...