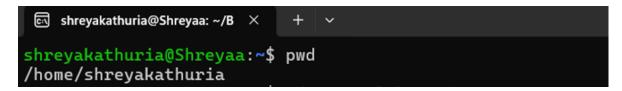
Indian Institute of Technology Gandhinagar BE623 Biocomputing Sem1 2025-2026

Lab session –1 (Shell, Filesystem and Permissions)

1- Open terminal and find out the path of the folder being displayed in the terminal. (Hint- Use pwd command to check the current location in terminal and cd command to change the directory)



2- Change directory to the folder with files uploaded on classroom. Create a new folder named "BE623 labsession 1" (Hint- Use mkdir command)

```
shreyakathuria@Shreyaa:~/B × + ∨

shreyakathuria@Shreyaa:~$ pwd
/home/shreyakathuria
shreyakathuria@Shreyaa:~$ cd BE623_labsession_1
-bash: cd: BE623_labsession_1: No such file or directory
shreyakathuria@Shreyaa:~$ mkdir BE623_labsession_1
```

3- Enter BE623 labsession 1 folder (Hint use cd command)

```
shreyakathuria@Shreyaa:~/B × + v

shreyakathuria@Shreyaa:~$ pwd
/home/shreyakathuria
shreyakathuria@Shreyaa:~$ cd BE623_labsession_1
-bash: cd: BE623_labsession_1: No such file or directory
shreyakathuria@Shreyaa:~$ mkdir BE623_labsession_1
shreyakathuria@Shreyaa:~$ cd BE623_labsession_1
shreyakathuria@Shreyaa:~/BE623_labsession_1
shreyakathuria@Shreyaa:~/BE623_labsession_1$
```

4- Confirm that your current folder is "BE623 labsession 1" (Hint use pwd command)

```
shreyakathuria@Shreyaa:~/B × + v

shreyakathuria@Shreyaa:~$ pwd
/home/shreyakathuria
shreyakathuria@Shreyaa:~$ cd BE623_labsession_1/
shreyakathuria@Shreyaa:~/BE623_labsession_1$ pwd
/home/shreyakathuria/BE623_labsession_1
shreyakathuria@Shreyaa:~/BE623_labsession_1$
shreyakathuria@Shreyaa:~/BE623_labsession_1$
```

- 5- List all the files and folders in the directory (Hint use Is command)
- 6- Copy "sequence.fasta and protein.fasta" file to current folder

```
shreyakathuria@Shreyaa:~/BE623_labsession_1$ Ls
protein.fasta
shreyakathuria@Shreyaa:~/BE623_labsession_1$ cd /mnt/c/Users/SHREYA/Downloads/sequence.fasta .
-bash: cd: too many arguments
shreyakathuria@Shreyaa:~/BE623_labsession_1$ cp /mnt/c/Users/SHREYA/Downloads/sequence.fasta .
shreyakathuria@Shreyaa:~/BE623_labsession_1$ ls
protein.fasta sequence.fasta
shreyakathuria@Shreyaa:~/BE623_labsession_1$ |
```

7- List all the files and folders in the directory (Hint use ls command)

```
shreyakathuria@Shreyaa:~/BE623_labsession_1$ ls
protein.fasta sequence.fasta
shreyakathuria@Shreyaa:~/BE623_labsession_1$ ls
protein.fasta sequence.fasta
shreyakathuria@Shreyaa:~/BE623_labsession_1$ |
```

8- Find out if there is any hidden file or folder in the directory (Hint: ls -al)

```
shreyakathuria@Shreyaa:~/BE623_labsession_1$ ls
protein.fasta sequence.fasta
shreyakathuria@Shreyaa:~/BE623_labsession_1$ ls -al
total 92
drwxr-xr-x 2 shreyakathuria shreyakathuria 4096 Aug 11 07:40 .
drwxr-xr--- 9 shreyakathuria shreyakathuria 4096 Aug 11 07:30 ..
-rwxrwxrwx 1 shreyakathuria shreyakathuria 467 Aug 9 13:05 protein.fasta
-rwxr-xr-x 1 shreyakathuria shreyakathuria 79551 Aug 11 07:40 sequence.fasta
shreyakathuria@Shreyaa:~/BE623_labsession_1$
```

9- Find out permissions associated with files in the folder (Hint: ls -al)

```
shreyakathuria@Shreyaa:~/BE623_labsession_1$ ls -l
total 84
-rwxrwxrwx 1 shreyakathuria shreyakathuria 467 Aug 9 13:05 protein.fasta
-rwxr-xr-x 1 shreyakathuria shreyakathuria 79551 Aug 11 07:40 sequence.fasta
shreyakathuria@Shreyaa:~/BE623_labsession_1$
```

10- Find out the permissions of the files and folders in /

```
shreyakathuria@Shreyaa: ~ × + ~
 hreyakathuria@Shreyaa:~$ cd BE623_labsession_1/
 hreyakathuria@Shreyaa:~/BE623_labsession_1$ cd
hreyakathuria@Shreyaa:~$ ls -l /
total 2740
                                          7 Apr 22 2024 bin -> usr/bin
4096 Feb 26 2024 bin.usr-is-merged
4096 Apr 22 2024 boot
                  1 root root
2 root root
2 root root
lrwxrwxrwx
drwxr-xr-x
drwxr-xr-x
                  drwxr-xr-x 15 root root
drwxr-xr-x 88 root root
drwxr-xr-x
-rwxrwxrwx
lrwxrwxrwx
drwxr-xr-x
lrwxrwxrwx
drwx----
drwxr-xr-x
drwxr-xr-x
drwxr-xr-x
                                         4096 Feb 15 08:09 opt
0 Aug 9 07:49 proc
4096 Feb 15 08:11 root
560 Aug 9 07:49 run
8 Apr 22 2024 sbin -> usr/sbin
4096 Mar 31 2024 sbin.usr-is-merged
4096 Aug 6 17:56 snap
4096 Feb 15 08:09 srv
0 Aug 11 04:23 sys
4096 Aug 11 03:58 tmp
4096 Feb 15 08:09 usr
4096 Aug 6 17:56 var
dr-xr-xr-x 258 root root
drwx----
                  3 root root
drwxr-xr-x 19 root root
                  1 root root
2 root root
lrwxrwxrwx
drwxr-xr-x
                   2 root root
drwxr-xr-x
drwxr-xr-x 2 root root
dr-xr-xr-x 13 root root
                  8 root root
drwxrwxrwt
drwxr-xr-x 12 root root
drwxr-xr-x 13 root root
                                          4096 Aug
                                                        6 17:56 var
 hreyakathuria@Shreyaa:~$
```

11- Change the permissions of the file "protein.fasta" so that it is readable, writable and executable by everyone (Hint use chmod).

```
shreyakathuria@Shreyaa:~$ cd BE623_labsession_1/
shreyakathuria@Shreyaa:~/BE623_labsession_1$ chmod a+rwx protein.fasta
shreyakathuria@Shreyaa:~/BE623_labsession_1$ ls -al
total 92
drwxr-xr-x 2 shreyakathuria shreyakathuria 4096 Aug 11 07:40 .
drwxr-x--- 9 shreyakathuria shreyakathuria 4096 Aug 11 07:30 ..
-rwxrwxrwx 1 shreyakathuria shreyakathuria 467 Aug 9 13:05 protein.fasta
-rwxr-xr-x 1 shreyakathuria shreyakathuria 79551 Aug 11 07:40 sequence.fasta
shreyakathuria@Shreyaa:~/BE623_labsession_1$
```

Change the permissions back to what they were.

```
shreyakathuria@Shreyaa:~/BE623_labsession_1$ chmod a+rwx protein.fasta
shreyakathuria@Shreyaa:~/BE623_labsession_1$ ls-l
ls-l: command not found
shreyakathuria@Shreyaa:~/BE623_labsession_1$ ls -l
total 84
-rwxrwxrwx 1 shreyakathuria shreyakathuria 467 Aug 9 13:05 protein.fasta
-rwxr-xr-x 1 shreyakathuria shreyakathuria 79551 Aug 11 07:40 sequence.fasta
shreyakathuria@Shreyaa:~/BE623_labsession_1$ |
```