

To run a command as administrator (user "root"), use "sudo <command>".
See "man sudo_root" for details.

```
user@DESKTOP-A5PNEA5:~$ pwd
/home/user
user@DESKTOP-A5PNEA5:~$
```

To run a command as administrator (user "root"), use "sudo <command>".
See "man sudo_root" for details.

```
user@DESKTOP-A5PNEA5:~$ pwd
/home/user
user@DESKTOP-A5PNEA5:~$ cd
user@DESKTOP-A5PNEA5:~$
```

To run a command as administrator (user "root"), use "sudo <command>".
See "man sudo_root" for details.

```
user@DESKTOP-A5PNEA5:~$ pwd  
/home/user
```

```
user@DESKTOP-A5PNEA5:~$ cd
```

```
user@DESKTOP-A5PNEA5:~$ mkdir "BE623_labsession_1"
```

```
user@DESKTOP-A5PNEA5:~$
```

To run a command as administrator (user "root"), use "sudo <command>".
See "man sudo_root" for details.

```
user@DESKTOP-A5PNEA5:~$ pwd
/home/user
user@DESKTOP-A5PNEA5:~$ cd
user@DESKTOP-A5PNEA5:~$ mkdir "BE623_labsession_1"
user@DESKTOP-A5PNEA5:~$ cd "BE623_labsession_1"
user@DESKTOP-A5PNEA5:~/BE623_labsession_1$
```

```
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ pwd  
/mnt/c/Users/user  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ cd "BE623_labsession_1"  
-bash: cd: BE623_labsession_1: No such file or directory  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ mkdir "BE623_labsession_1"  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ cd "BE623_labsession_1"  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ |
```

```
Windows PowerShell
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PS C:\Users\user> pwd

Path
----
C:\Users\user

PS C:\Users\user> wsl

To run a command as administrator (user "root"), use "sudo <command>".
See "man sudo_root" for details.

user@DESKTOP-A5PNEA5:/mnt/c/Users/user$
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ pwd
/mnt/c/Users/user
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ cd "BE623_labsession_1"
-bash: cd: BE623_labsession_1: No such file or directory
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ mkdir "BE623_labsession_1"
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ cd "BE623_labsession_1"
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ pwd
/mnt/c/Users/user/BE623_labsession_1
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$
```

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PS C:\Users\user> pwd

Path

C:\Users\user

PS C:\Users\user> wsl

To run a command as administrator (user "root"), use "sudo <command>".
See "man sudo_root" for details.

```
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ pwd  
/mnt/c/Users/user  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ cd "BE623_labsession_1"  
-bash: cd: BE623_labsession_1: No such file or directory  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ mkdir "BE623_labsession_1"  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ cd "BE623_labsession_1"  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ pwd  
/mnt/c/Users/user/BE623_labsession_1  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$
```

```
Path
----
C:\Users\user
```

```
PS C:\Users\user> wsl
```

To run a command as administrator (user "root"), use "sudo <command>".
See "man sudo_root" for details.

```
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ pwd
/mnt/c/Users/user
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ cd "BE623_labsession_1"
-bash: cd: BE623_labsession_1: No such file or directory
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ mkdir "BE623_labsession_1"
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ cd "BE623_labsession_1"
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ pwd
/mnt/c/Users/user/BE623_labsession_1
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp sequence.fasta protein.fasta
cp: cannot stat 'sequence.fasta': No such file or directory
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp sequence.fasta
cp: missing destination file operand after 'sequence.fasta'
Try 'cp --help' for more information.
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp /mnt/c/Users/user/Downloads/sequence.fasta /mnt/c/Users/us
er/BE623_labsession_1
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls
sequence.fasta
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$
```



```
PS C:\Users\user> wsl
```

To run a command as administrator (user "root"), use "sudo <command>".
See "man sudo_root" for details.

```
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ pwd  
/mnt/c/Users/user  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ cd "BE623_labsession_1"  
-bash: cd: BE623_labsession_1: No such file or directory  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ mkdir "BE623_labsession_1"  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ cd "BE623_labsession_1"  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ pwd  
/mnt/c/Users/user/BE623_labsession_1  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp sequence.fasta protein.fasta  
cp: cannot stat 'sequence.fasta': No such file or directory  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp sequence.fasta  
cp: missing destination file operand after 'sequence.fasta'  
Try 'cp --help' for more information.  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp /mnt/c/Users/user/Downloads/sequence.fasta /mnt/c/Users/user/BE623_labsession_1  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls  
sequence.fasta  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp /mnt/c/Users/user/Downloads/protein.fasta /mnt/c/Users/user/BE623_labsession_1  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls  
protein.fasta sequence.fasta  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$
```

To run a command as administrator (user "root"), use "sudo <command>".
See "man sudo_root" for details.

```
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ pwd  
/mnt/c/Users/user  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ cd "BE623_labsession_1"  
-bash: cd: BE623_labsession_1: No such file or directory  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ mkdir "BE623_labsession_1"  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ cd "BE623_labsession_1"  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ pwd  
/mnt/c/Users/user/BE623_labsession_1  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp sequence.fasta protein.fasta  
cp: cannot stat 'sequence.fasta': No such file or directory  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp sequence.fasta  
cp: missing destination file operand after 'sequence.fasta'  
Try 'cp --help' for more information.  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp /mnt/c/Users/user/Downloads/sequence.fasta /mnt/c/Users/user/BE623_labsession_1  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls  
sequence.fasta  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp /mnt/c/Users/user/Downloads/protein.fasta /mnt/c/Users/user/BE623_labsession_1  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls  
protein.fasta sequence.fasta  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ pwd  
/mnt/c/Users/user/BE623_labsession_1  
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$
```

```

/mnt/c/Users/user
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ cd "BE623_labsession_1"
-bash: cd: BE623_labsession_1: No such file or directory
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ mkdir "BE623_labsession_1"
user@DESKTOP-A5PNEA5:/mnt/c/Users/user$ cd "BE623_labsession_1"
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ pwd
/mnt/c/Users/user/BE623_labsession_1
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp sequence.fasta protein.fasta
cp: cannot stat 'sequence.fasta': No such file or directory
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp sequence.fasta
cp: missing destination file operand after 'sequence.fasta'
Try 'cp --help' for more information.
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp /mnt/c/Users/user/Downloads/sequence.fasta /mnt/c/Users/user/BE623_labsession_1
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls
sequence.fasta
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp /mnt/c/Users/user/Downloads/protein.fasta /mnt/c/Users/user/BE623_labsession_1
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls
protein.fasta sequence.fasta
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ pwd
/mnt/c/Users/user/BE623_labsession_1
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls -al
total 80
drwxrwxrwx 1 user user 512 Aug 11 11:40 .
drwxrwxrwx 1 user user 512 Aug 11 08:30 ..
-rwxrwxrwx 1 user user 467 Aug 11 11:40 protein.fasta
-rwxrwxrwx 1 user user 79551 Aug 11 11:38 sequence.fasta
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$

```

```

/mnt/c/Users/user/BE623_labsession_1
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp sequence.fasta protein.fasta
cp: cannot stat 'sequence.fasta': No such file or directory
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp sequence.fasta
cp: missing destination file operand after 'sequence.fasta'
Try 'cp --help' for more information.
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp /mnt/c/Users/user/Downloads/sequence.fasta /mnt/c/Users/us
er/BE623_labsession_1
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls
sequence.fasta
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ cp /mnt/c/Users/user/Downloads/protein.fasta /mnt/c/Users/use
r/BE623_labsession_1
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls
protein.fasta  sequence.fasta
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ pwd
/mnt/c/Users/user/BE623_labsession_1
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls -al
total 80
drwxrwxrwx 1 user user  512 Aug 11 11:40 .
drwxrwxrwx 1 user user  512 Aug 11 08:30 ..
-rwxrwxrwx 1 user user  467 Aug 11 11:40 protein.fasta
-rwxrwxrwx 1 user user 79551 Aug 11 11:38 sequence.fasta
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls -al
total 80
drwxrwxrwx 1 user user  512 Aug 11 11:40 .
drwxrwxrwx 1 user user  512 Aug 11 08:30 ..
-rwxrwxrwx 1 user user  467 Aug 11 11:40 protein.fasta
-rwxrwxrwx 1 user user 79551 Aug 11 11:38 sequence.fasta
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$

```


try 'ls --help' for more information.

user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1\$ ls -al /

total 2748

drwxr-xr-x	22	root	root	4096	Aug	11	08:27	.
drwxr-xr-x	22	root	root	4096	Aug	11	08:27	..
lrwxrwxrwx	1	root	root	7	Apr	22	2024	bin -> usr/bin
drwxr-xr-x	2	root	root	4096	Feb	26	2024	bin.usr-is-merged
drwxr-xr-x	2	root	root	4096	Apr	22	2024	boot
drwxr-xr-x	15	root	root	3860	Aug	11	08:27	dev
drwxr-xr-x	87	root	root	4096	Aug	11	08:27	etc
drwxr-xr-x	3	root	root	4096	Aug	11	05:33	home
-rwxrwxrwx	1	root	root	2724480	Jul	31	14:56	init
lrwxrwxrwx	1	root	root	7	Apr	22	2024	lib -> usr/lib
drwxr-xr-x	2	root	root	4096	Apr	8	2024	lib.usr-is-merged
lrwxrwxrwx	1	root	root	9	Apr	22	2024	lib64 -> usr/lib64
drwx-----	2	root	root	16384	Aug	11	05:25	lost+found
drwxr-xr-x	2	root	root	4096	Jan	6	2025	media
drwxr-xr-x	8	root	root	4096	Aug	11	05:25	mnt
drwxr-xr-x	2	root	root	4096	Jan	6	2025	opt
dr-xr-xr-x	214	root	root	0	Aug	11	08:27	proc
drwx-----	4	root	root	4096	Aug	11	05:25	root
drwxr-xr-x	19	root	root	560	Aug	11	08:27	run
lrwxrwxrwx	1	root	root	8	Apr	22	2024	sbin -> usr/sbin
drwxr-xr-x	2	root	root	4096	Mar	31	2024	sbin.usr-is-merged
drwxr-xr-x	2	root	root	4096	Aug	11	05:25	snap
drwxr-xr-x	2	root	root	4096	Jan	6	2025	srv
dr-xr-xr-x	13	root	root	0	Aug	11	08:27	sys
drwxrwxrwt	8	root	root	4096	Aug	11	08:28	tmp
drwxr-xr-x	12	root	root	4096	Jan	6	2025	usr
drwxr-xr-x	13	root	root	4096	Aug	11	05:25	var

```

lrwxrwxrwx 1 root root 7 Apr 22 2024 bin -> usr/bin
drwxr-xr-x 2 root root 4096 Feb 26 2024 bin.usr-is-merged
drwxr-xr-x 2 root root 4096 Apr 22 2024 boot
drwxr-xr-x 15 root root 3860 Aug 11 08:27 dev
drwxr-xr-x 87 root root 4096 Aug 11 08:27 etc
drwxr-xr-x 3 root root 4096 Aug 11 05:33 home
-rwxrwxrwx 1 root root 2724480 Jul 31 14:56 init
lrwxrwxrwx 1 root root 7 Apr 22 2024 lib -> usr/lib
drwxr-xr-x 2 root root 4096 Apr 8 2024 lib.usr-is-merged
lrwxrwxrwx 1 root root 9 Apr 22 2024 lib64 -> usr/lib64
drwx----- 2 root root 16384 Aug 11 05:25 lost+found
drwxr-xr-x 2 root root 4096 Jan 6 2025 media
drwxr-xr-x 8 root root 4096 Aug 11 05:25 mnt
drwxr-xr-x 2 root root 4096 Jan 6 2025 opt
dr-xr-xr-x 214 root root 0 Aug 11 08:27 proc
drwx----- 4 root root 4096 Aug 11 05:25 root
drwxr-xr-x 19 root root 560 Aug 11 08:27 run
lrwxrwxrwx 1 root root 8 Apr 22 2024 sbin -> usr/sbin
drwxr-xr-x 2 root root 4096 Mar 31 2024 sbin.usr-is-merged
drwxr-xr-x 2 root root 4096 Aug 11 05:25 snap
drwxr-xr-x 2 root root 4096 Jan 6 2025 srv
dr-xr-xr-x 13 root root 0 Aug 11 08:27 sys
drwxrwxrwt 8 root root 4096 Aug 11 08:28 tmp
drwxr-xr-x 12 root root 4096 Jan 6 2025 usr
drwxr-xr-x 13 root root 4096 Aug 11 05:25 var
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ chmod ugo + rwx protein.fasta
chmod: invalid mode: 'ugo'
Try 'chmod --help' for more information.
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ chmod ugo+rwx protein.fasta
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$

```

```

drwxr-xr-x 87 root root 4096 Aug 11 08:27 etc
drwxr-xr-x 3 root root 4096 Aug 11 05:33 home
-rwxrwxrwx 1 root root 2724480 Jul 31 14:56 init
lrwxrwxrwx 1 root root 7 Apr 22 2024 lib -> usr/lib
drwxr-xr-x 2 root root 4096 Apr 8 2024 lib.usr-is-merged
lrwxrwxrwx 1 root root 9 Apr 22 2024 lib64 -> usr/lib64
drwx----- 2 root root 16384 Aug 11 05:25 lost+found
drwxr-xr-x 2 root root 4096 Jan 6 2025 media
drwxr-xr-x 8 root root 4096 Aug 11 05:25 mnt
drwxr-xr-x 2 root root 4096 Jan 6 2025 opt
dr-xr-xr-x 214 root root 0 Aug 11 08:27 proc
drwx----- 4 root root 4096 Aug 11 05:25 root
drwxr-xr-x 19 root root 560 Aug 11 08:27 run
lrwxrwxrwx 1 root root 8 Apr 22 2024 sbin -> usr/sbin
drwxr-xr-x 2 root root 4096 Mar 31 2024 sbin.usr-is-merged
drwxr-xr-x 2 root root 4096 Aug 11 05:25 snap
drwxr-xr-x 2 root root 4096 Jan 6 2025 srv
dr-xr-xr-x 13 root root 0 Aug 11 08:27 sys
drwxrwxrwt 8 root root 4096 Aug 11 08:28 tmp
drwxr-xr-x 12 root root 4096 Jan 6 2025 usr
drwxr-xr-x 13 root root 4096 Aug 11 05:25 var
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ chmod ugo + rwx protein.fasta
chmod: invalid mode: 'ugo'
Try 'chmod --help' for more information.
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ chmod ugo+rwx protein.fasta
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls -l
total 80
-rwxrwxrwx 1 user user 467 Aug 11 11:40 protein.fasta
-rwxrwxrwx 1 user user 79551 Aug 11 11:38 sequence.fasta
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$

```

```

lrwxrwxrwx 1 root root      9 Apr 22  2024 lib64 -> usr/lib64
drwx----- 2 root root 16384 Aug 11 05:25 lost+found
drwxr-xr-x 2 root root  4096 Jan  6  2025 media
drwxr-xr-x 8 root root  4096 Aug 11 05:25 mnt
drwxr-xr-x 2 root root  4096 Jan  6  2025 opt
dr-xr-xr-x 214 root root    0 Aug 11 08:27 proc
drwx----- 4 root root  4096 Aug 11 05:25 root
drwxr-xr-x 19 root root   560 Aug 11 08:27 run
lrwxrwxrwx 1 root root      8 Apr 22  2024 sbin -> usr/sbin
drwxr-xr-x 2 root root  4096 Mar 31  2024 sbin.usr-is-merged
drwxr-xr-x 2 root root  4096 Aug 11 05:25 snap
drwxr-xr-x 2 root root  4096 Jan  6  2025 srv
dr-xr-xr-x 13 root root    0 Aug 11 08:27 sys
drwxrwxrwt 8 root root  4096 Aug 11 08:28 tmp
drwxr-xr-x 12 root root  4096 Jan  6  2025 usr
drwxr-xr-x 13 root root  4096 Aug 11 05:25 var
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ chmod ugo + rwx protein.fasta
chmod: invalid mode: 'ugo'
Try 'chmod --help' for more information.
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ chmod ugo+rwx protein.fasta
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls -l
total 80
-rwxrwxrwx 1 user user  467 Aug 11 11:40 protein.fasta
-rwxrwxrwx 1 user user 79551 Aug 11 11:38 sequence.fasta
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ chmod ugo-rwx protein.fasta
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$ ls -l
total 80
-r-xr-xr-x 1 user user  467 Aug 11 11:40 protein.fasta
-rwxrwxrwx 1 user user 79551 Aug 11 11:38 sequence.fasta
user@DESKTOP-A5PNEA5:/mnt/c/Users/user/BE623_labsession_1$

```