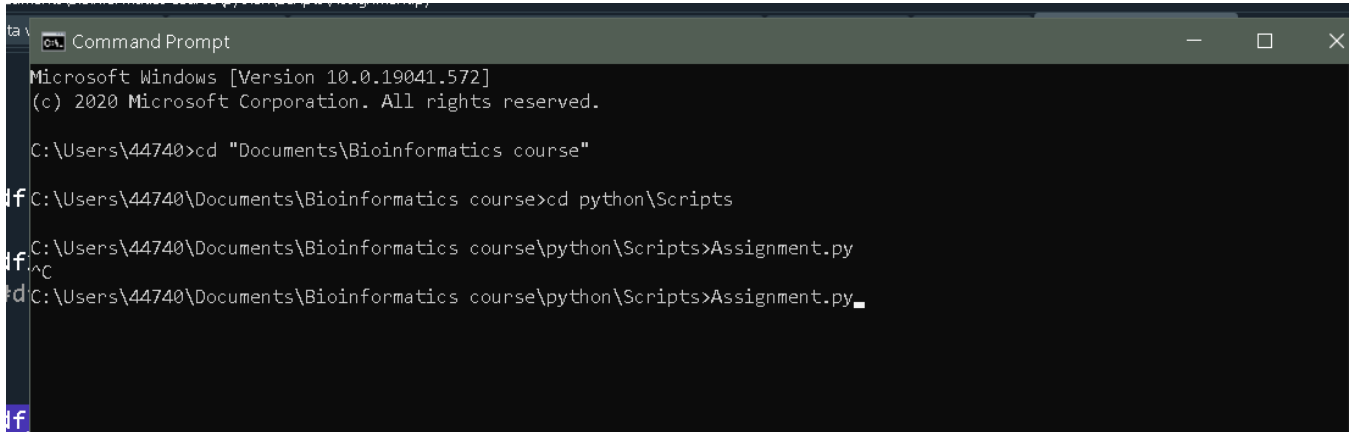


User Manual

1. Go to the command prompt and give the directory path ("script path\python\scriptname.py") for the py script (Assignment.py) from the desktop like below screenshot:



```
Microsoft Windows [Version 10.0.19041.572]
(c) 2020 Microsoft Corporation. All rights reserved.

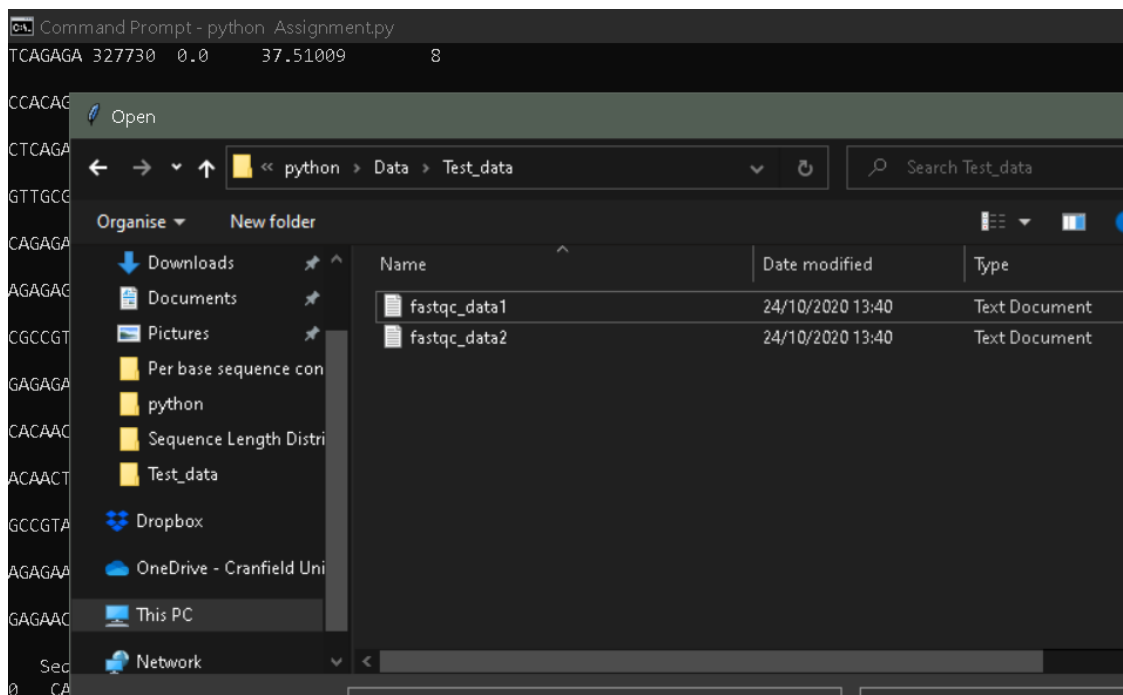
C:\Users\44740>cd "Documents\Bioinformatics course"

C:\Users\44740\Documents\Bioinformatics course>cd python\Scripts

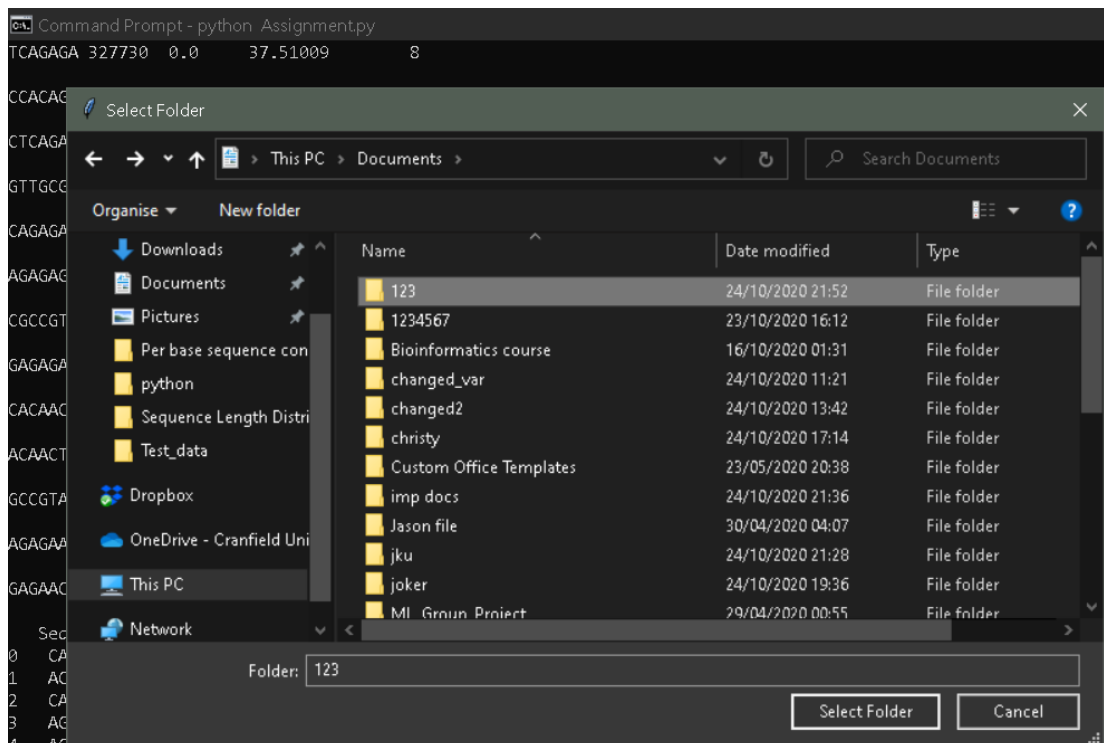
C:\Users\44740\Documents\Bioinformatics course\python\Scripts>Assignment.py

C:\Users\44740\Documents\Bioinformatics course\python\Scripts>Assignment.py
```

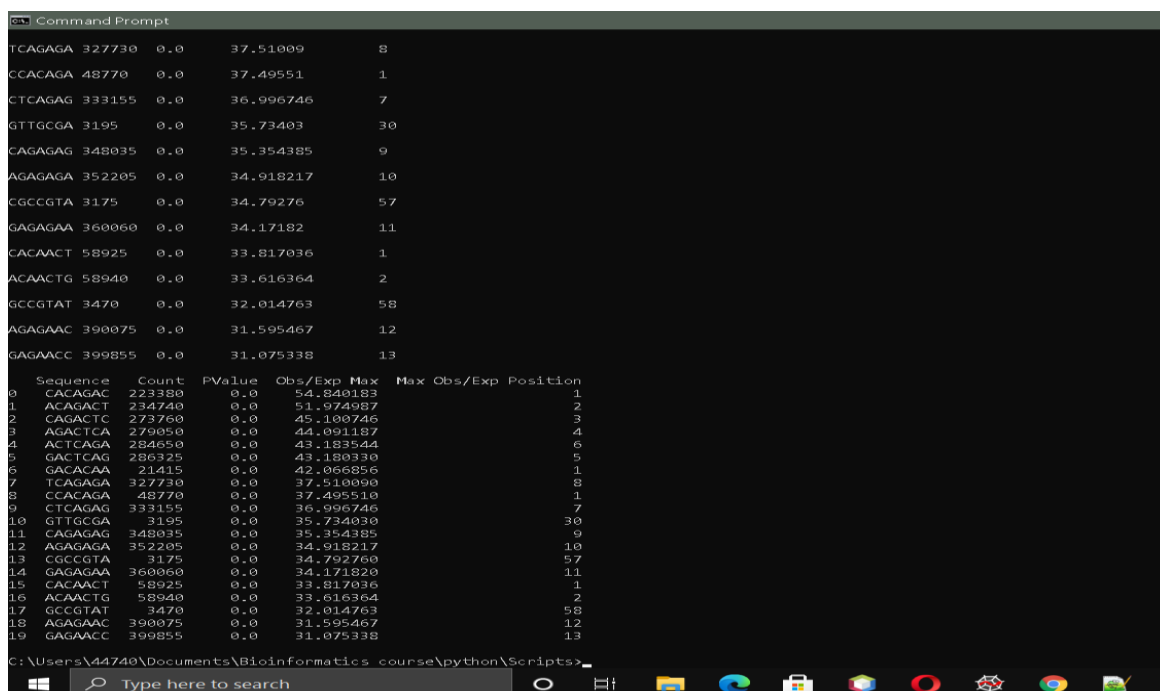
2. A dialog box appears and the user can go to the specific directory where the data is present, then select any of the two fastq file and select folder.



3. Instantly another dialog box appears to select the output folder (make an output folder) where all the files and graphs would be saved and enter.



4. The script will now be run in the command line and the window will appear like below screen and the report can be seen.



5. Now to view the report files and graph for each module, please go to the earlier selected output directory and you may find all the files here in the folder “output directory”.

