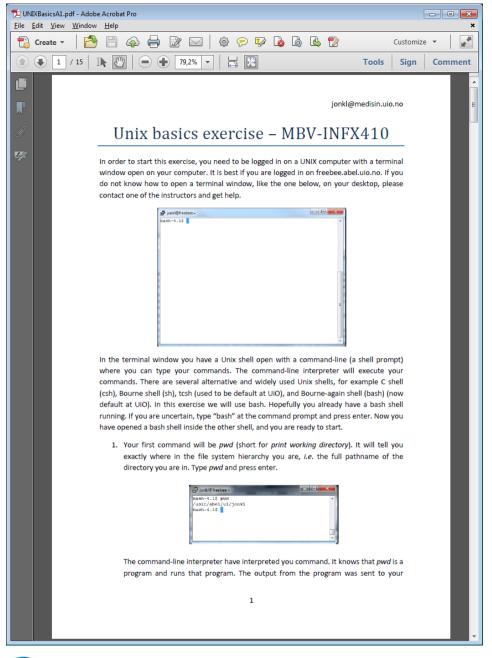
# Summing up Unix, Python, etc.





Very important to be able to use basic Unix

Make your own Unix scripts is not very central

Most likely you will instead use Python for writing small programs/scripts



jonkl@medisin.uio.no

## A brief practical Unix/Python exercise – MBV-INFX410

We will in this exercise work with a practical task that, it turns out, can easily be solved by using basic Unix and Python.

Let us pretend that an engineer in your group has spent several weeks growing and Sanger sequencing several hundred clones. All the successful sequencing data has been "packed" in one single file called allChroms.tar.gz.

 Log onto freebee.abel.uio.no and create a new directory in your home area. Let us call it "chromatograms".



- 2. Download the file allChroms.tar.gz from the wiki page and put it in the new directory. How you do this will depend on your laptop. When you have done this, make sure you understand what you did. We will do similar operations more times during the course (and very likely for the exam...). This is important!
- 3. The file has a double ending, ".tar.gz". This indicates that this is a compressed file that has been compressed, or packed to save space, by the gzip software application (hence the ".gz"). It is also a "tar file", also known as a "tarball", which usually means that many files have been packed into a single file. This is often done to make file transfer and/or file storage easier. Use Is -I allChroms.tar.gz to see the size of the

### Main points:

- Demonstrate a practical task where making a script is absolutely necessary
- Show you that some tasks cannot easily be solved by using «databases and web servers» including Galaxy/LifePortal





## A brief practical Unix/Python exercise – MBV-INFX410

We will in this exercise work with a practical task that, it turns out, can easily be solved by using basic Unix and Python.

Let us pretend that an engineer in your group has spent several weeks growing and Sanger sequencing several hundred clones. All the successful sequencing data has been "packed" in one single file called allChroms.tar.gz.

- Log onto freebee.abel.uio.no and create a new directory in your home area. Let us call it "chromatograms".
  - After this exercise you *must* be able to:
    - In a Unix shell, create directories, move around between the directories and copy and move files. Use *ls* with options to see which files and directories you have in a directory.
    - Use cat and more to view the contents of a file, and use nano to write text into a file or create a new one.
    - Delete files
    - o Transfer files from your laptop to freebee.abel.uio.no and back
    - Download a compressed tarball file from somewhere and store it on your laptop and on freebee.abel.uio.no
    - Uncompress and extract the files from the tarball
    - Make a tarball archive file with tar and compress with gzip
    - Make programs executable and files visible or hidden from other users with chmod
    - o Know how to use grep, "pipe" (|), and how to redirect output with ">"
    - If you obtain a little Python script, you should be able to run it and, at least if it is not too complicated, be able to figure out what it is doing

# This is most important!



## Python

- Steep (or not?) learning curve
- Day 1
- Day 2
- Classes, objects and Biopython
- You are not expected to become experienced programmers
- The exercise on MSAs and "the Python script for moving files" illustrates what I expect you to be able to do with Python



- You must be able to move files between laptop and freebee.abel.uio.no!
- You must be able to uncompress and get all files out of a gzipped tarball, for example bigfile.tar.gz
  - And make this kind of archive file yourself



- Three ways to run a Python script
- 1: In an interactive shell

```
jonkl@freebee:~/MBV-INFX410_Python
                                                                         bash-4.1$ python
Python 2.6.6 (r266:84292, May 27 2013, 05:35:12)
[GCC 4.4.7 20120313 (Red Hat 4.4.7-3)] on linux2
Type "help", "copyright", "credits" or "license" for more information.
>>> print "Hello World"
Hello World
>>> print len("HelloWorld")
10
>>>
```



- Three ways to run a Python script
- 2: As a script sent as input to Python

```
jonkl@freebee:~/MBV-INFX410_Python
                                                                         _ - X
bash-4.1$ cat myScript.py
import sys
first word = sys.argv[1]
print first word, "has", len(first word), "letters"
bash-4.1$ python myScript.py Banana
Banana has 6 letters
bash-4.1$ python myScript.py Bananarama
Bananarama has 10 letters
bash-4.1$
```



- Three ways to run a Python script
- 3: As a script with #! at first line

```
jonkl@freebee:~/MBV-INFX410_Python
                                                                                        bash-4.1$ cat myScript2.py
#!/usr/bin/pvthon
import sys
first word = sys.argv[1]
print first word, "has", len(first word), "letters"
bash-4.1$ ./mvScript2.pv Apple
Apple has 5 letters
bash-4.1$ ls -1 myScript2.py
-rwx----- 1 jonkl jonkl 108 Nov 17 12:04 myScript2.py
bash-4.1$ chmod u-x myScript2.py
bash-4.1$ ls -1 myScript2.py
-rw----- 1 jonkl jonkl 108 Nov 17 12:04 myScript2.py
bash-4.1$ ./myScript2.py Apple
bash: ./myScript2.py: Permission denied
bash-4.1$
```



 Why can we do Is, cat, rmdir etc, but have to do ./myScript2

- Environment variables in the shell
- \$PATH, \$HOME etc
- \$HOME is same as ~/

```
jonkl@freebee:~/MBV-INFX410 Python
bash-4.1$ ./myScript2.py Hi
Hi has 2 letters
bash-4.1$ myScript2.py Hi
bash: myScript2.py: command not found
bash-4.1$ ls myScript2.py
myScript2.py
bash-4.1$ head -n 1 myScript2.py
#!/usr/bin/python
bash-4.1$ echo $PATH
/usr/lib64/gt-3.3/bin:/usr/local/bin:/bin:/usr/bin:/opt/bin
bash-4.1$ which ls
/bin/ls
bash-4.1$ echo $HOME
/usit/abel/u1/jonkl
bash-4.1$
```

## Non-printing characters

 Use cat -v to see "invisible" control codes in the text

```
jonkl@freebee:~/MBV-INFX410_Python
                                                                                      bash-4.1$ python parse blast.py
  File "parse blast.py", line 1
SyntaxError: Non-ASCII character '\xe2' in file parse blast.py on line 1, but no encoding decl
ep-0263.html for details
bash-4.1$ cat parse blast.pv | head -3
fh = open("blastout2.txt", "r")
lines =
fh.close()
bash-4.1$ cat -v parse blast.py | head -3
fh = open(M-bM-^@M-^\blastout2.txtM-bM-^@M-^], M-bM-^@M-^\rM-bM-^@M-^])
lines =
fh.close()
bash-4.1$
```



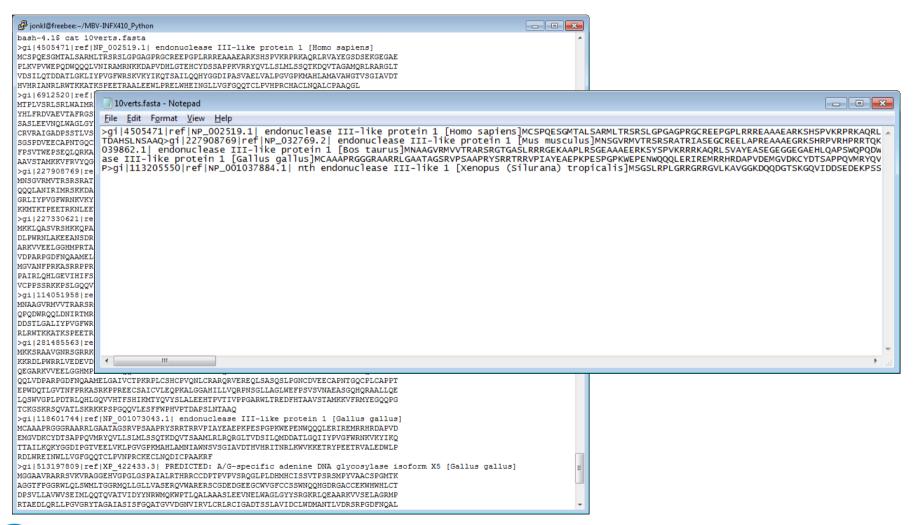
## End-of-line markers (EOLs) are *not* the same in Unix and MS Windows!

- Use cat -v to see "invisible" control codes, for example Windows type EOLs
- Use dos2unix command to go from MS Windows to Unix
- Use unix2dos command to do the reverse

```
- - X
jonkl@freebee:~/MBV-INFX410_Python
bash-4.1$ cat myScript2.py
#!/usr/bin/python
import svs
first word = sys.argv[1]
print first word, "has", len(first word), "letters"
bash-4.1$ ./myScript2.py
bash: ./myScript2.py: /usr/bin/python^M: bad interpreter: No such file or directory
bash-4.1$ cat -v myScript2.py
#!/usr/bin/python^M
import sys^M
first word = sys.argv[1]^M
print first word, "has", len(first word), "letters"^M
bash-4.1$ dos2unix mvScript2.pv
dos2unix: converting file mvScript2.pv to UNIX format ...
bash-4.1$ cat -v mvScript2.pv
#!/usr/bin/python
import sys
first word = sys.argv[1]
print first word, "has", len(first word), "letters"
bash-4.1$ ./mvScript2.pv Hei
Hei has 3 letters
bash-4.1$
```

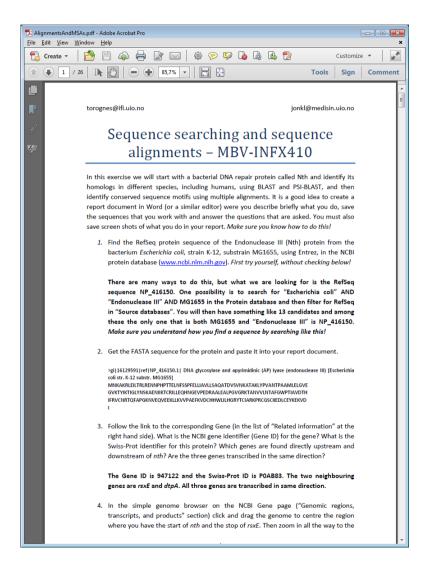


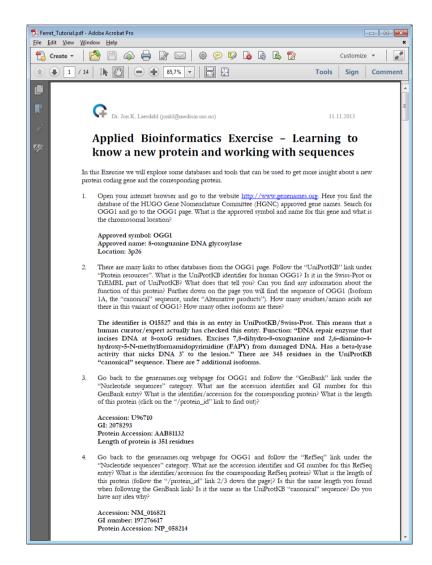
# End-of-line markers (EOLs) are *not* the same in Unix and MS Windows!





## These are very important!









## Articles and written material

- Curriculum comprises
  - All lectures
  - All «handouts»
  - All exercises, demos and computers lab sessions
  - All articles listed on the wiki
  - Obligatory assignment

https://wiki.uio.no/projects/clsi/index.php/MBV-INFX410 2015

	Week 46	Week 47	Week 48	Week 49	Week 50	Week 51
Mon	Lecture/ exercises 1st day of course - Nov 9	Lecture/ exercises	Study day/work on obligatory take-home assignment	Lecture/ exercise	Study day	
Tues	Lecture/ exercises	Lecture/ exercises	Study day/work on oblig take- home assignment	Lecture/ exercises	Study day	
Wed	Lecture/ exercises	Lecture/ exercises	Study day/work on oblig take- home assignment	Lecture/ exercises	Take-home exam start	Take-home exam finish
Thurs	Lecture/ exercises	Lecture/ exercises	Study day/work on oblig take- home assignment	Lecture/ exercises		
Fri	Lecture/ exercises	Lecture/ exercises	Hand in oblig take-home assignment	Study day		
Sat/Sun						



# You will only be able to take the exam if you follow all teaching!

If you cannot finish the exercises during the exercise sessions, finish them at home!

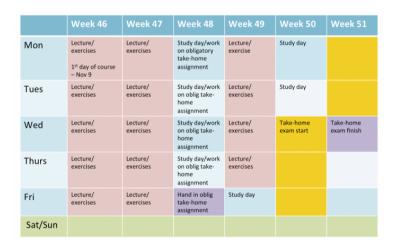
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Sat/Sun						





# You will only be allowed to take the exam if your obligatory assignment has been approved!

- Exercise for oblig will be handed out at the end of week 2
- Will be relatively easy and similar to exercises in weeks 1 and 2
- Must be returned before the first lecture in week 3 (4)
- PhD students (MBV-INF9410) must in addition write an assignment/essay
  - Describe how you would some of the methods covered in the course in your own research
  - > 2500 words







Page 1 of 3

### UNIVERSITETET I OSLO

#### Det matematisk-naturvitenskapelige fakultet

Obligatory assignment: MBV-INF4410 and MBV-INF9410

Deadline: November 30th 2015, at 09:00

Permitted materials: All written material, including all internet resources

Only students that gets this assignment approved will be permitted to take the course home exam.

Your completed assignment must be returned, at the latest, at 9 am, Monday November 30. It should be sent by e-mail to the course coordinator Jon K. Lærdahl (e-mail address: jonkl@medisin.uio.no). Please put the course code and your name in the subject field (e.g. Oblig MBV-INF4410 Dolly Duck").

The <u>oblig</u> must be handed in as a single PDF document (Microsoft Word or an Open Office Document is also acceptable). Please also include your name and course code in the document and in the document title.

You are encouraged to use screenshots and other figures in order to improve your explanations.

#### THE WORK MUST REPRESENT YOUR OWN ANSWERS

Answers should contain only what is asked for. Some questions have multiple parts. Your answers may be given in English or in Norwegian. *Technical questions* about the oblig can be answered by Jon K. Lærdahl (e-mail address: jonkl@medisin.uio.no).

#### EXTRA Obligatory assignment: MBV-INF9410 (only)

Write an essay of at least 2500 words on one of these two topics

- How can some of the methods described in this course be used in your own research?
- A course relevant topic of your own choice. In this case you must get this approved by Jon K. Lærdahl before you write the essay



Jon K. Lærdahl, Structural Bioinformatics

UiO • Department of Informatics
University of Oslo