Practical Bioinformatics

Basic Linux

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Why Linux?

In Bioinformatics

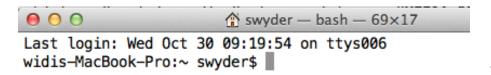
- Many bioinformatics and genomics tools are command-line only and are optimized to run on Linux.
- Very powerful with text files, even large (GBs)

In general

- Experienced Linux users can do more with less effort and much faster
- Multi-user
- Lots of control and customization possibilities
- 0\$
-

Why command line?

- to run bioinformatics software
- you can automate tasks more easily
- to interact with high-performance (high-memory) servers
- offers handy tools to work with text files

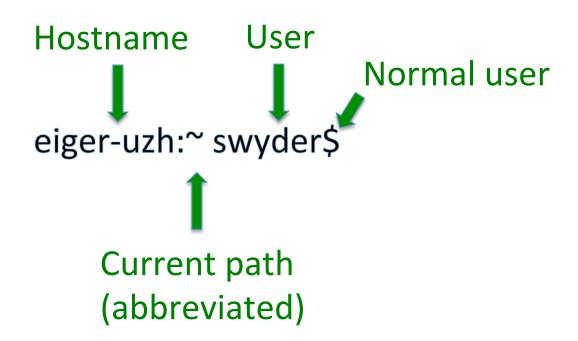


- The shell is an interactive interpreter: it reads commands, finds the corresponding programs, runs them, and displays output
- Today we use the Bash shell (default shell of most Linux distributions and Mac OS X)

The prompt

Prompt

eiger-uzh:~ swyder\$



Commands

- The tools philosophy was to have small programs to accomplish a particular task instead of trying to develop large monolithic programs to do a large number of tasks.
- "Designed to operate together"
 To accomplish more complex tasks, tools can simply be connected together.
- The shell comprises hundreds of commands, but if you know 25 you can achieve many things
- Commands are abbreviations to type less (ls:list, cp:copy, mv:move)
- Common structure:

Command -Option(s) Parameter(s) Is -I /home/swyder/tmp

Setting options

Command -Option(s) Parameter(s)

The order of options does not matter unless they override each other (e.g. sorting)

Options, grep as an example

- \$ grep apple fruitlist.txt apple pineapple
- \$ grep -w apple fruitlist.txt (or grep -x) apple
- \$ grep apple *.txt fruitlist.txt:apple fruitlist.txt:pineapple recipeFruitSalad.txt:1 pineapple recipePinaColada.txt:2oz fresh pineapple juice

\$ grep -v apple fruitlist.txt banana pear peach

• \$ grep apple fruitlist.txt recipeFruitSalad.txt fruitlist.txt:apple fruitlist.txt:pineapple recipeFruitSalad.txt:1 pineapple

Also options to color, to show context, search with compl patterns

Getting help

man <command>

man cp

BSD General Commands Manual CP(1)

CP(1)

NAME

cp -- copy files

SYNOPSIS

cp [-R [-H | -L | -P]] [-fi | -n] [-apvX] source file target file

cp [-R [-H | -L | -P]] [-fi | -n] [-apvX] source file ... target directory

DESCRIPTION

In the first synopsis form, the cp utility copies the contents of the source file to the targe contents of each named source file is copied to the destination target directory. The named source file is copied to the destination target directory. If cp detects an attempt to copy a file to itself, the copy will fail.

The following options are available:

- Same as -pPR options. Preserves structure and attributes of files but not directory st
- -f If the destination file cannot be opened, remove it and create a new file, without pro permissions. (The -f option overrides any previous -n option.)

space: scroll down a page

b: scroll up q: quit man <command> --help cp --help

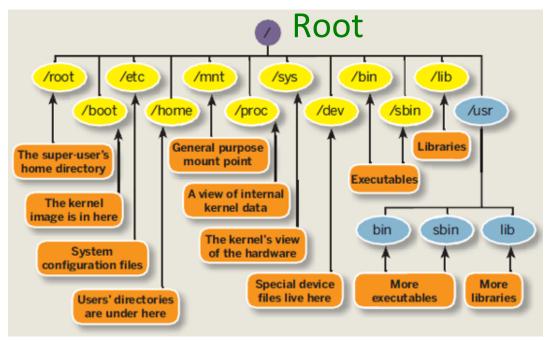
a less verbose help (not working on Mac OS)

The web

http://linuxconfig.org/linux-commands

Working with files and directories

Directory structure



http://www.tuxradar.com/

- Everything the system uses is located somewhere under root '/'.
- Every user has his home directory, e.g. /home/swyder
- Do your work in your home directory
- The system folders can only be modified by the administrator

File and Directory names

- Upper- and lower-case matter
- up to 256 characters long
- Every character except / is allowed. But by convention special characters like \$äéÜ are not used.
- Don't use white spaces. Use underscores (_), hashes (-) and dots (.) to separate words
 Oct2013_RNAseq
 Oct2013\ RNAseq

Main commands

Command	Meaning	
Is	Content of current directory	
cd <i>dir_name</i>	Change to directory	
cd	Change to home directory	
cd ~	Change to home directory	
mkdir	Make a directory	
ср	Copy a file/directory	
rm	Delete a file/directory	

Working with text files

Main text commands

UNIX has an extensive toolkit for text extraction, reporting and manipulation

Task	Commands	
Show	less, more, head, tail, cat	
Search/Extract	grep, cut, awk, uniq	
Manipulate	sort, tr, sed, join, paste	
Replace	tr, sed	
Count	wc, uniq -c	
Compare	comm, diff	



Piping

Philosophy

"filters": simple programs which only do 1 thing the output of a filter is the input of the next

grep "mRNA" test.gff | less

grep -w "gene" test.gff | cut -f 1 | sort | uniq -c

Redirection

- > Writing the output to a file
- >> Appending the output to a file

```
Is > output.txt
grep -w "gene" test.gff | cut -f 1 | sort | uniq -c > output.txt
Is >> output.txt
```

< Reading from file

wc < hello.txt > hello_counts.txt

What we learned in Part 1

- Command -Option(s) Parameter(s)
 Is -rlh ~/data
- Working with files / directories ls, cd, mkdir, cp, mv, rm
- Directory structure everything is under the root: /
- Working with text files head, less, grep, cut, sort, tr, wc, uniq
- Tools can be connected by "|"
- The Mac OS X Shell differs from the typical Linux shell

Sources & Links

Acknowledgements

Some exercises are from von Mering group (IMLS, UZH)

Material

- SIB course http://edu.isb-sib.ch/course/view.php?id=41
- O'reilly Books http://oreilly.com/linux/
- Video tutorials (~100 min) http://software-carpentry.org/v4/shell/index.html
- Cheatsheet http://www.embnet.org/sites/default/files/quickguides/guideUNIX.pdf

Additional topics

- Connecting with Unix/Linux servers
- File/Dir Compression&Extraction
- Installing and running software
- Permissions
- Differences in the shell MA

Server commands

Command	Task
ssh -X user@hostname	Connect to server
scp <what> <towhere></towhere></what>	Transfer file from/to server
sftp user@hostname	Transfer file from/to server (interactive)

Connect to remote computer

```
$ ssh username@mnf-44.uzh.ch

RSA key fingerprint is 71:ed:af:1f:d6:0a:43:05:8d:11:34:68:
2c:2d:79:01.

Are you sure you want to continue connecting (yes/no)?
```

Type "yes" and press ENTER.

Then you will be asked for your password.

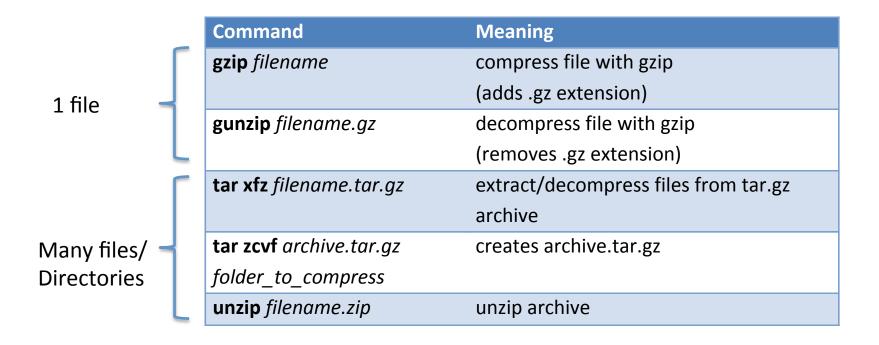
```
$ bash # we will use the bash shell

$ whoami # what is my username on this host

$ uname -a # show basic info of the host OS

$ df -h # displays free disk space
```

File/Dir compression



Running programs

- Software are executable files (permissions!)
- Run a bash script

```
chmod +x script1.sh  # make file executable
./script1.sh or bash path/script.sh  # run
```

Its the same for any scripting language (python/perl,...)

```
chmod +x script.py
./script.py or python path/script.py
```

Run a binary

```
chmod +x bowtie
./bowtie
```

Installing Software (binaries)

Packages

Using a package manager - takes also care of dependencies

- Linux:

Ubuntu: via .deb files (e.g. aptitude or apt-get)

Fedora/SUSE: via .rpm files

- Mac OS X: homebrew (my favourite), MacPorts, fink

Compiling from source

Typically open-source software is written in C/C++ -> GCC compiler

- Linux: install gcc using the package manager (apt-get search gcc, then apt-get install gcc-XXXX)
- Mac Os X: install gcc using homebrew (brew search gcc, ...) or via XCode

```
./configure
make
make install  # optional
make clean  # optional
```

To install an executable

Bash only looks at certain directories for commands/software/programs \$PATH is a variable

echo \$PATH /usr/bin:/usr/sbin:/usr/local/bin:/opt/X11/bin:/usr/texbin

- 1. You copy it into one of the folders in \$PATH
- 2. You add its directory to \$PATH
- 3. You create a symbolic link to it into a folder contained in \$PATH

Details

- 2. export PATH=\$PATH:directory
- 3. sudo ln -s executable *directory*

Summary: executables

- Come in two flavours: Scripts / Binaries
- Execute permissions must be set:

```
chmod +x programname
```

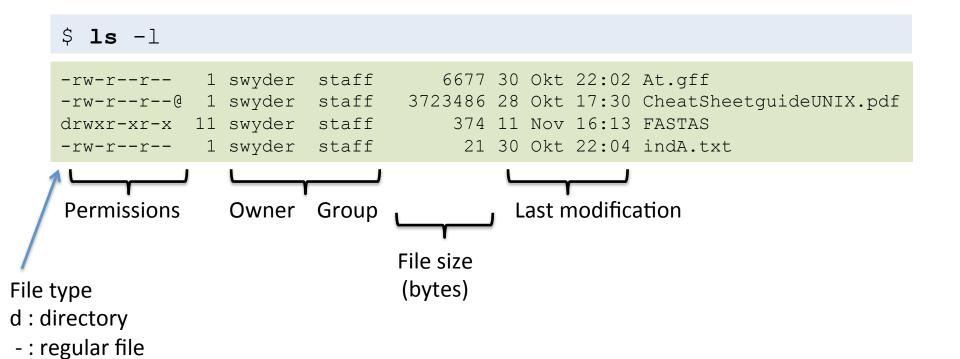
 Scripts mostly start with the shebang line, telling the shell which interpreter to use. E.g.

```
#!/usr/bin/perl
```

Executing of executables

```
./prg
./prg.sh (bash prg.sh)
python prg.py
perl prg.pl
```

Permissions



Changing Permissions

```
r: read
w: write
x: execute

$ chmod [ugo][+-][rwx] file

u: user
g: group
o: other/world
```

Make a file executable (for you)

\$ chmod +x file
chmod ug+rx my_file

Setting exact permission (only reading, for you)

\$ chmod =r file

Removing permissions (for you)

\$ chmod -wx file

Permission code

Owner	Group	Other
r w x	r w x	r - x
4+2+1	4+2+1	4+0+1
4	_	4
7	7	5

r: read w: write

x: execute

chmod 775 file

Differences in the Shell Linux - Mac OS

Differences Linux - Mac OS X

Mac line breaks are '\r\n' (and Windows: '\r') instead
of the standard Linux '\n'
When working in the command line make sure the files have the right format
perl -pi -e 's/\r\n?/\n/g' <filename>
which you could alias and put in the .bashrc

 Mac has no GNU tools by default less options – less powerful

for installation install homebrew then: brew install coreutils All GNU commands are then installed with the prefix 'g': gls, gcp, gsed,...

 Mac has non-standard folder structure e.g. home (~) is /Users/swyder