

Quality of Life tips

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Enabler for Life Science











Terminal improvements

- There are a lot of small tips that will improve your experience greatly.
- This lecture will cover some of them:
 - Navigating the terminal
 - Shortcuts for programs, files and directories
 - Finding files and contents of files
- Also covered: How to transfer files with rsync, scp



Create shortcuts to files and catalogs

```
$ ln -s /link/to/folder/or/file
```



Create shortcuts to files and catalogs

```
$ ln -s /link/to/folder/or/file
```

You can create shortcuts to access your project catalog without the entire path

```
$ cd ~
$ ln -s /proj/g2019015/nobackup/yourusername g2019015
```

Create shortcuts to files and catalogs

```
$ ln -s /link/to/folder/or/file
```

You can create shortcuts to access your project catalog without the entire path

```
$ cd ~
$ ln -s /proj/g2019015/nobackup/yourusername g2019015
```

Now you can go to the project directory from your home folder:

```
$ cd g2019015
```

Create shortcuts to files and catalogs

```
$ ln -s /link/to/folder/or/file
```

Now create a shortcut to this lab's folder!

```
$ ln -s /sw/courses/ngsintro/qol/
```



```
$ alias sc="program -gRe /proj/g209999/test"
```



```
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$ sc
```



```
$ alias sc="program -gRe /proj/g209999/test"
$ sc
```

```
$ alias ngsintro="module load bioinfo-tools;
module load samtools; module load bwa; cd
/proj/g2019015/nobackup/yourusername"
```



```
$ alias sc="program -gRe /proj/g209999/test"
$ sc

$ alias ngsintro="module load bioinfo-tools;
module load samtools; module load bwa; cd
/proj/g2019015/nobackup/yourusername"

$ ngsintro
```



Command history

Up & down arrow to step through history



Command history

Up & down arrow to step through history

Ctrl+r to search through previous commands



Command history

Up & down arrow to step through history

- Ctrl+r to search through previous commands
 - Ctrl+r again to search further back



Quicker cursor

 Navigating the command line can be done quicker than with just left and right arrows



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ctrl+a and ctrl+e places the cursor at the beginning and end of command line respectively



Quicker cursor

 Navigating the command line can be done quicker than with just left and right arrows

ctrl+a and ctrl+e places the cursor at the beginning and end of command line respectively

alt+b to go back a word, alt+f to go forward



Cursor position

- No need to be at end of line when pressing enter
- Only determines where you type or erase

```
$ echo "Position not important"
```



Cursor position

- No need to be at end of line when pressing enter
- Only determines where you type or erase

```
$ echo "Position not important"
```

```
user@rackham5 ~/ $
```



Remembers previous directory

user@rackham5 ~/ \$ cd /proj/g2099004



```
user@rackham5 ~/ $ cd /proj/g2099004 user@rackham5 /proj/g2099004 $
```



```
user@rackham5 ~/ $ cd /proj/g2099004 user@rackham5 /proj/g2099004 $ cd -
```



```
user@rackham5 ~/ $ cd /proj/g2099004
user@rackham5 /proj/g2099004 $ cd -
user@rackham5 ~/ $
```



```
user@rackham5 ~/ $ cd /proj/g2099004
user@rackham5 /proj/g2099004 $ cd -
user@rackham5 ~/ $ cd -
```



```
user@rackham5 ~/ $ cd /proj/g2099004
user@rackham5 /proj/g2099004 $ cd -
user@rackham5 ~/ $ cd -
user@rackham5 /proj/g2099004 $
```



UPPMAX cheat sheet

• https://uppmax.uu.se/support/getting-started/uppmax-cheat-sheet/

SNIC-UPPMAX CHEAT SHEET

ssh -AX user@host	Connect to host with ssl
Getting Help	
man command	Read manual for command
apropos keyword	Find commands related to keyword
File commands	
ls	List contents of current di
ls -al	Detailed listing with hidden files
cd dir Go to direc	ctory (if dir is not given, go to home dir
pwd	Show the current directory
mkdir dir	Create directory
rm file	Remove file
rm -r dir	Recursively remove directory
rm -f file	Force remove file
rm -rf dir	Force recursively remove directory
cp -i file1 file2	Copy file1 to file:
cp -r dirl dir2	Recursively copy director
mv -i filel file2	Rename or move file/director
(if file	2 is a directory, it places file1 inside it
ln -s file link	Create a symbolic lini
touch file Create fi	le or update timestamp of existing one
command > file	Write output of a command to a file
command >> file	Append (add to end) output to file
less file Show con	itents of file, with scrolling (quit with 'q
head file	Show 10 first lines of file
tail file	Show 10 last lines of file
tail -f file Show file	e as it grows, starting with 10 last lines
nano file Edit file	with a simple command line text editor
File Permissions	
chmod permission fi	ile Change file permission
chmod -R perm dir	Recursively change permission for di
Ex: Allow read/write/exec for	or user, r/w for group and r for others:
chmod u=rwx,g=rw,o=	· ·
chmod a-x file	x: Remove execute permission for al

grep pattern files	Find lines in files, containing pattern
grep -r pattern dir	Recursively do same as above in dir
command grep patte	ern Run grep on command output
find dir grep patt	tern Find filepaths matching pattern
Compression	
tar cf file.tar file	Create tar archive, adding files
tar xf file.tar	Extract tar archive
tar czf file.tar.gz	files With gzip compression
tar xzf file.tar.gz	Extract gzip compressed tar archive
gzip file	Compress file with gzip
gunzip file.gz	Decompress file with gzip
UPPMAX modules	
module avail	List available modules
module load modulens	ume Load module
Showing user and project i	info
uquota	Show current user's disk usage
projinfo Show used	core hours for current user's projects
View details of a specific proj	
egrep '^b2011999' /e	
Running jobs with the Slur	m resource manager
jobinfo Show all r	unning and waiting jobs in the queue
jobinfo -u user	Show jobs for specific user
interactive -A proje	Start interactive job
Start batch job (see user guid	de on the web for more info):
	d-hh:mm:ss -n cores \
-p partition jobscz	ript file
	cores (2 nodes) on node partition:
Ex. Pullilling for Tuays on 10	+ 7-00:00:00 -= 16 \
sbatch -A b2011999 -	C /-00:00:00 -n 16 (
sbatch -A b2011999 -	
sbatch -A b2011999 - -p node my_jobscrip	
sbatch -A b2011999 -	ot_file



www.snic.vr.se



- Copy files between computers
- Similar syntax as cp

```
rsync user@host:/path/to/file /local/path
scp user@host:/path/to/file /local/path
```



- Copy files between computers
- Similar syntax as cp

```
rsync user@rackham.uppmax.uu.se:/home/user/t.txt .
scp user@rackham.uppmax.uu.se:/home/user/t.txt .
```



- Copy files between computers
- Similar syntax as cp
- rsync: -a save modification time, -P show progress
- scp: -p save modification time

```
rsync -aP user@rackham.uppmax.uu.se:/home/user/t.txt .
scp -p user@rackham.uppmax.uu.se:/home/user/t.txt .
```



- Copy files between computers
- Similar syntax as cp
- rsync: -a save modification time, -P show progress
- scp: -p save modification time
- Copy files either direction

```
rsync -aP t.txt user@rackham.uppmax.uu.se:/home/user/
scp -p t.txt user@rackham.uppmax.uu.se:/home/user/
```



- Copy files between computers
- Similar syntax as cp
- rsync: -a save modification time, -P show progress
- scp: -p save modification time
- Copy files either direction

```
rsync -aP user@rackham.uppmax.uu.se:/sw/courses/ngsintro/qol/aa.fa .
scp -p user@rackham.uppmax.uu.se:/sw/courses/ngsintro/qol/aa.fa .
```



Find files based on name

\$ find /path/to/look/ -name nametolookfor.txt



```
$ find /path/to/look/ -name nametolookfor.txt
$ find . -name *.txt
```



```
$ find /path/to/look/ -name nametolookfor.txt
$ find . -name *.txt
$ cd ~/g2019015/qol
$ tree
```



```
$ find /path/to/look/ -name nametolookfor.txt
$ find . -name *.txt
$ cd ~/g2019015/qol
$ tree
$ find . -name CARROTCAKE
```



```
$ find /path/to/look/ -name nametolookfor.txt
$ find . -name *.txt
$ cd ~/g2019015/qol
$ tree
$ find . -name CARROTCAKE
$ find . -name CARROT*
```



Searches content of files

\$ grep texttofind filetolookin.txt



Searches content of files

```
$ grep texttofind filetolookin.txt
```

```
$ grep "text to find" /path/to/files/*.txt
```



Searches content of files

```
$ grep texttofind filetolookin.txt
$ grep "text to find" /path/to/files/*.txt
```

\$ grep MYNAME protein seq.fa



Searches content of files

```
$ grep texttofind filetolookin.txt
$ grep "text to find" /path/to/files/*.txt
$ grep MYNAME protein_seq.fa
$ grep -r found filetree
```





- Kills whatever your terminal is currently running
 - Destroys hanged or non-executing commands
 - Cancel commands that are running for too long
 - Cancel commands that you realize are incorrect



Using multiple terminals

- Launch and use several terminals for better overview of your work
 - Write your scripts in one terminal, run your scripts in another, read the output in a third...
- Just like you might have several tabs in your browser or have several documents open at once