

Quality of Life

a better terminal experience

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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
 - Finding files and contents of files
- Also covered: How to transfer files to/from Rackham



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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 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



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 ~/ $
```



```
user@rackham5 ~/ $ cd /proj/g2020009
```



```
user@rackham5 ~/ $ cd /proj/g2020009 user@rackham5 /proj/g2020009 $
```



```
user@rackham5 ~/ $ cd /proj/g2020009 user@rackham5 /proj/g2020009 $ cd -
```



```
user@rackham5 ~/ $ cd /proj/g2020009
user@rackham5 /proj/g2020009 $ cd -
user@rackham5 ~/ $
```



```
user@rackham5 ~/ $ cd /proj/g2020009
user@rackham5 /proj/g2020009 $ cd -
user@rackham5 ~/ $ cd -
```



```
user@rackham5 ~/ $ cd /proj/g2020009
user@rackham5 /proj/g2020009 $ cd -
user@rackham5 ~/ $ cd -
user@rackham5 /proj/g2020009 $
```



- Copy files between computers
- Similar syntax as cp

rsync 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



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

rsync -aP 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
- Copy files either direction

```
rsync -aP user@rackham.uppmax.uu.se:/home/user/t.txt .
rsync -aP 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
- Copy files either direction

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



Find

Find files based on name

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



Find

Find files based on name

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



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
$ cd /sw/courses/ngsintro/linux/qol
$ grep MYNAME protein seq.fa
```



Searches content of files

```
$ grep texttofind filetolookin.txt
$ grep "text to find" /path/to/files/*.txt
$ cd /sw/courses/ngsintro/linux/qol
$ grep MYNAME protein_seq.fa
$ grep -r "found" filetree/*
```





- Kills whatever your terminal is currently running
 - Sometimes we want to end programs prematurely, for whichever reason.
- \$ cd /sw/courses/ngsintro/linux/qol \$ python friendly counter.py





- 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



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 file:
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 on
command > file	Write output of a command to a file
command >> file	Append (add to end) output to file
less file Show con	tents 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	as it grows, starting with 10 last line:
	with a simple command line text editor
File Permissions	
chmod permission fi	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=	r file
chmod a-x file	x: Remove execute permission for al

Arch barrern rrres	Find lines in files, containing pattern
grep -r pattern di	E Recursively do same as above in dir
command grep pat	tern Run grep on command output
find dir grep pa	ttern Find filepaths matching pattern
Compression	
tar cf file.tar fi	les Create tar archive, adding files
tar xf file.tar	Extract tar archive
tar czf file.tar.g	z files With gzip compression
tar xzf file.tar.q	z Extract gzip compressed tar archive
gzip file	Compress file with gzip
gunzip file.gz	Decompress file with gzip
UPPMAX modules	THE AMERICAN
module avail	List available modules
module load module	name Load module
Showing user and project	ct info
uquota	Show current user's disk usage
projinfo Show use	ed core hours for current user's projects
view details of a specific p	
View details of a specific p egrep '^b2011999'	/etc/slurm/grantfile
	/etc/slurm/grantfile
egrep '^b2011999' Running jobs with the SI	/etc/slurm/grantfile urm resource manager
egrep '^b2011999' Running jobs with the SI	/etc/slurm/grantfile lurm resource manager Il running and waiting jobs in the queue
egrep '^b2011999' Running jobs with the SI jobinfo Show a	/etc/slurm/grantfile urm resource manager Il running and waiting jobs in the queue Show jobs for specific user
egrep '^b2011999' Running jobs with the Si jobinfo Show a jobinfo -u user interactive -A pro	/etc/slurm/grantfile urm resource manager Il running and waiting jobs in the queue Show jobs for specific user ject Start interactive job
egrep '^b2011999' Running jobs with the Si jobinfo Show a jobinfo -u user interactive -A pro Start batch job (see user g	/etc/slurm/grantfile urm resource manager Il running and waiting jobs in the queue Show jobs for specific user
egrep '^b2011999' Running jobs with the Si jobinfo Show a jobinfo -u user interactive -A pro Start batch job (see user g	/etc/slurm/grantfile lurm resource manager Il running and waiting jobs in the queue Show jobs for specific user ject Start interactive job uide on the web for more info): -t d-hh:mm:ss -n cores \
egrep '^b2011999' Running jobs with the Si jobinfo Show a jobinfo -u user interactive -A pro Start batch job (see user g sbatch -A project -p partition jobs	/etc/slurm/grantfile lurm resource manager Il running and waiting jobs in the queue Show jobs for specific user ject Start interactive job uide on the web for more info): -t d-hh:mm:ss -n cores \
egrep '^b2011999' Running jobs with the SI jobinfo Show a jobinfo -u user interactive -A pro Start batch job (see user g sbatch -A project -p partition jobs Ex: Running for 7 days on	/etc/slurm/grantfile lurm resource manager Il running and waiting jobs in the queue Show jobs for specific user ject Start interactive job uide on the web for more info): -t d-hh:mm:ss -n cores \ ccript_file
egrep '^b2011999' Running jobs with the Si jobinfo Show a jobinfo -u user interactive -A pro Start batch job (see user g sbatch -A project -p partition jobs Ex: Running for 7 days on	/etc/slurm/grantfile turm resource manager Il running and waiting jobs in the queue Show jobs for specific user ject Start interactive job uide on the web for more info): -t d-hh:mm:ss -n cores \ cript_file 16 cores (2 nodes) on node partition: -t 7-00:00:00 -n 16 \
egrep '^b2011999' Running jobs with the SI jobinfo Show a jobinfo -u user interactive -A pro Start batch job (see user g sbatch -A project -p partition jobs Ex: Running for 7 days on sbatch -A b2011999	/etc/slurm/grantfile turm resource manager Il running and waiting jobs in the queue Show jobs for specific user ject Start interactive job uide on the web for more info): -t d-hh:mm:ss -n cores \ ccript file 16 cores (2 nodes) on node partition: -t 7-00:00:00 -n 16 \ ipt_file
egrep '^b2011999' Running jobs with the SI jobinfo Show a jobinfo -u user interactive -A pro Start batch job (see user g sbatch -A project -p partition jobs Ex: Running for 7 days on sbatch -A b2011999 -p node my_jobscr	/etc/slurm/grantfile lurm resource manager Il running and waiting jobs in the queue Show jobs for specific user ject Start interactive job uide on the web for more info): -t d-hh:mm:ss -n cores \ cript_file 16 cores (2 nodes) on node partition: -t 7-00:00:00 -n 16 \ ipt_file Cancel a single job





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, check the results in a third...
- Just like you might have several tabs in your browser or have several documents open at once