

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 Dardel



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
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Remembers previous directory

user@login1 ~/ \$



• Remembers previous directory

user@login1 ~/ \$ cd /proj/g2020009



• Remembers previous directory

user@login1 ~/ \$ cd /proj/g2020009 user@login1 /proj/g2020009 \$



• Remembers previous directory

user@login1 ~/ \$ cd /proj/g2020009 user@login1 /proj/g2020009 \$ cd -



• Remembers previous directory

user@login1 ~/ \$ cd /proj/g2020009 user@login1 /proj/g2020009 \$ cd user@login1 ~/ \$



• Remembers previous directory

user@login1 ~/ \$ cd /proj/g2020009 user@login1 /proj/g2020009 \$ cd user@login1 ~/ \$ cd -



• Remembers previous directory

user@login1 ~/ \$ cd /proj/g2020009 user@login1 /proj/g2020009 \$ cd user@login1 ~/ \$ cd user@login1 /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@dardel.pdc.kth.se:/cfs/klemming/home/u/user/t.txt \



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

```
rsync -aP \
user@dardel.pdc.kth.se:/cfs/klemming/home/u/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@dardel.pdc.kth.se:/cfs/klemming/home/u/user/t.txt \
.
```

```
rsync -aP t.txt \
user@dardel.pdc.kth.se:/cfs/klemming/home/u/user/
```



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

```
rsync -aP \
user@dardel.pdc.kth.se:/sw/courses/ngsintro/linux/gol/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



- 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