# **Bioinformatics Preparatory Course**

Basic Unix

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  - Searching, organizing, transferring large numbers of files
  - Exploring and manipulating plain text files
  - Can be programmed

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  - SSH (secure shell) interface to access a Unix-like system remotely
  - Standard to interact with larger computer infrastructures (e.g. a compute server or cluster)
- The majority of bioinformatics tools have a CLI
  - We work with a high number of large files
  - Crunching that data requires high compute power ⇒ compute server, cluster
  - Most file formats established in Bioinformatics are plain text

## Basic Usage

ullet Enter a command and press the Return  $(\display)$  key

```
$ ls ↓

Desktop Documents Exercise
```

• To get help on how to use a particular command, try

```
$ ls --help ↓
$ man ls ↓
```

• The basic command syntax often follows this structure:

```
COMMAND [OPTIONAL SWITCHES] [MANDATORY ARGUMENTS]
```

## Basic Usage

• Often there are long (--) and short (-) versions of switches

```
$ ls ↓

Desktop Documents Exercise

$ ls -r ↓

Exercise Documents Desktop

$ ls --reverse ↓

Exercise Documents Desktop
```

#### Basic Usage

• Often there are long (--) and short (-) versions of switches

```
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Desktop Documents Exercise

$ ls -r ↓

Exercise Documents Desktop

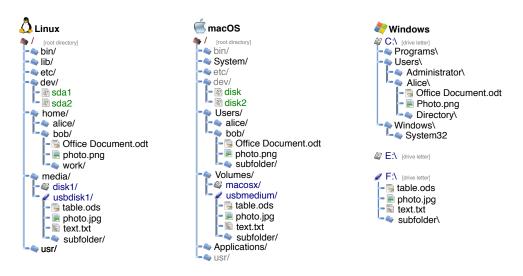
$ ls --reverse ↓

Exercise Documents Desktop
```

Multiple switches can be specified together

```
$ ls -F -r ←
Exercise/ Documents/ Desktop/
$ ls -Fr ←
Exercise/ Documents/ Desktop/
```

#### A typical computer file system is structured like a tree



• Every running computer program, i.e. every process, including the shell, refers to one directory inside the file system tree as its working directory.

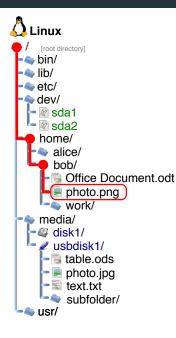
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- When using the shell, one often says "I am in the directory xzy".
- The working directory is not static, it can be changed throughout the runtime of a process.

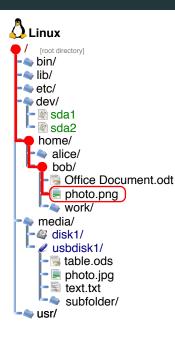


```
[root directory]
🖦 bin/
-⊸ lib/
🗕 🦤 etc/
-—— dev/
 🖢 🖳 sda2
   home/
  🖦 alice/
 bob/
    - Grice Document.odt
    🗕 📃 photo.png
   work/
   media/
  usbdisk1/
    - 🥞 table.ods
       photo.jpg
        text.txt
        subfolder/
 🔷 usr/
```

• Every file or directory in the file system can be described by a path

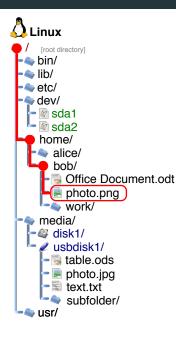


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  - ./bob/photo.png

given that the current working directory is /home.



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- Relative paths start in the current working directory, for example
  - ./bob/photo.png
    given that the current working directory is /home.
- Every character sequence that does not start with a slash / character is interpreted as a relative path.

pwd - print working directory
 Prints the current working directory

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   Prints the current working directory
- ls list
   Displays the contents of folders (directories)
  - ls -l show details
  - 1s -1h human readable file sizes
  - ls -l show hidden files and folders

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- ls list
   Displays the contents of folders (directories)
  - 1s -1 show details
  - 1s -1h human readable file sizes
  - 1s -1 show hidden files and folders
- cd change directory
   Changes the current working directory

mkdir <path> - make directory
 Creates a new directory

- mkdir <path> make directory
   Creates a new directory
- cp <source path> <destination path> copy
   Copies files and folders. Overwriting targets cannot be undone!
  - cp -r recursive mode
     Required to copy directories with their contents.

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   Move files and folders Overwriting targets cannot be undone!
- rmdir <path> remove directory
   Removes directories, but only if they are empty. Safe to use.

- rm <path> remove
   Remove files and folders. Removing cannot be undone!
  - cp -r recursive mode
     Required to remove directories with their contents.

Getting started with plain text files

• more <path>

Page through text one page at a time

- more <path>
  - Page through text one page at a time
- less <path> the opposite of more
   More powerful than more, bidirectional, provides searching

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  - Page through text one page at a time
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   More powerful than more, bidirectional, provides searching
- cat <path> [<path> ...] concatenate

  Concatenates input files and prints them to the standard output (more on that later).

• head <path>

Output the first n lines of a plain text file.

- head <path>
   Output the first n lines of a plain text file.
- tail <path>
   Output the last n lines of a plain text file.
  - tail -f follow
     Keeps printing new lines as the file grows.

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- Nano is a text editor that does not require a graphical user interface but works on the command line

#### • nano <path>

Opens the file specified by path. Can be used on non-existing paths to create new files.

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#### • CTRL-X

Exits nano

#### Keyboard shortcuts

TAB

Autocomplete the command line

- TAB TAB
   Show possible completions if not unique
- ALT-b and ALT-f
   Move one word backward (or forward) in the current command line
- Pos1 or Home
   Move to the beginning of the current command line
- End

Move to the end of the current command line

### **Keyboard shortcuts**



### Keyboard shortcuts

- Arrow Up
  - Browse the shell history backward in time
- Arrow Down
  - Browse the shell history forward in time
- CTRL-R
  - Search the shell history backward in time
- CTRL-C
  - Interrupt the current program.

- Every process has three input / output streams
  - Standard input (stdin)
  - Standard output (stdout)
  - Standard error (stderr)

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- This output is produced on stdout

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$ ls -l Exercise/data ↓

total 13960
-rw-r--r-- 1 vorkurs vorkurs 14293917 Mar 27
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This output is produced on stderr

```
$ ls -l doesnt.exist ←
ls: cannot access 'doesnt.exist': No such file or directory
```

• We can redirect streams into files using the > character

```
$ ls -l Exercise/data 1>stdout.txt 2>stderr.txt ←
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Try the two examples above and check the contents of the two output files each time!

- The > redirection overwrites the target file!
- Use >> to instead append the contents to the target file

#### Warm up

- Create a folder named repetition
- Create a file in that folder named repetition.txt
- Write the text Hello World into that file
- Copy the file to a new file named repetition2.txt
- Duplicate the content of the file repetition2.txt and write it to the file repetition3.txt
- Print the content of the three files
- Delete the file repetition.txt
- Copy the folder repetition to the folder repetition2 with all its contents
- Rename the folder repetition to repetition\_old
- Remove the contents of the folder repetition\_old
- Remove the folder repetition\_old
- Remove the folder repetition2

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- A user can be a real person or just be an arbitrary logical entity
- Groups are sets of users
- Every file has a single owner and is also associated with a single group

```
$ ls -l Exercise/data ↓

total 13960
-rw-r--r-- 1 vorkurs vorkurs 14293917 Mar 27 13:42 clinvar_20180225.vcf

Group

The file owner
```

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  - read
  - write
  - execute

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- There are three access types:
  - read
  - write
  - execute
- This triplet of permissions (rwx) is defined for three groups of users:
  - The owner
  - Members of the file's group
  - The rest of the world

```
$ ls -l Exercise/data ←
total 13960
-rw-r--r-- 1 vorkurs vorkurs 14293917 Mar 27 13:42 clinvar_20180225.vcf
                     Everyone else can also read (r-)
                     Members of the group vorkurs can read (r-)
                     The owner can read and write (rw-)
```

The interpretation of the permissions for directories is slightly different

- read determines whether the contents of a directory can be seen
- write determines whether files can be created or deleted
- execute determines whether a user can change (cd) into the directory

chown [owner][:[group] <path> - change ownership
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   Changes the ownership and / or group association of a file or directory. Only the root user is allowed to change the owner of a file.
- chmod <permissions> <path> change file mode bits, aka permissions
   Changes the file permissions. Examples:
  - chmod g+w <path>
     Add write permissions to the group
  - chmod o-r <path>
     Remove read permissions for the rest of the world
  - Shortcuts:

     (o)wner, (g)roup, (o)thers, (a)ll at once
     (r)ead, (w)rite, e(x)ecute

```
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total 13960
-rw-r--r-- 1 vorkurs vorkurs 14293917 Mar 27 13:42 clinvar_20180225.vcf
```

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$ ls -l Exercise/data ↓
total 13960
-rw-r--r-- 1 vorkurs vorkurs 14293917 Mar 27 13:42 clinvar_20180225.vcf
```

#### Add the write permission to everyone:

```
$ chmod a+w Exercise/data/clinvar_20180225.vcf.gz ↓
$ ls -l Exercise/data ↓
total 13960
-rw-rw-rw- 1 vorkurs vorkurs 14293917 Mar 27 13:42 clinvar_20180225.vcf
```

- Permissions can also be specified numerically
  - 4: read (r)
  - 2: write (w)
  - 1: execute (x)

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  - 4: read (r)
  - 2: write (w)
  - 1: execute (x)
- To combine permissions, the numbers are added
- Three digits specify the permissions for all three groups (owner, group, rest of the world) simultaneously:

#### chmod 750 <path>

- 7=4+2+1: owner has all permissions (rwx)
- 5=4+1: group members have read and execute permissions (rx)
- 0: rest of the world has no permissions

• wget <url>

Downloads a remote file. Supports HTTP(S) and FTP.

wget -c or wget --background
 Continue downloading a file that was already partially downloaded

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- curl -L0 <url>

   curl is another command line tool to download remote files, similar to wget.

## Example **wget** invocation:

```
$ wget https://compbio.de/vorkurs/material.zip ←
--2020-10-20 20:20:43-- https://compbio.de/vorkurs/material.zip
Resolving compbio.de (compbio.de)... 188.40.166.122
Connecting to compbio.de (compbio.de)|188.40.166.122|:443... connected.
HTTP request sent, awaiting response... 200 OK
Length: 10586 (10K) [application/zip]
Saving to: 'material.zip'
material.zip 100\%[========>] 10.34K --.-KB/s in 0.001s
2020-10-20 20:20:43 (15.3 MB/s) - 'material.zip' saved [10586/10586]
```

## Example **curl** invocation:

```
$ curl -LO https://compbio.de/vorkurs/material.zip ←
 % Total
           % Received % Xferd Average Speed
                                            Time
                                                   Time
                                                           Time
                                                                Current
                             Dload Upload
                                           Total
                                                   Spent
                                                           Left Speed
100 10586
         100 10586
                     0
                             57532
                                       0 --:--:- 57532
```

File compression and file archives

# File compression and file archives

- The most common compressed archive file formats are .tar.gz and zip
- tar is an archive format to reversibly combine multiple files into one file, gz (gzip) is a data compression tool
- zip is an "all in one" solution, i.e. does archiving and compression

- unzip -l <path>Shows the contents of a zip file
- unzip <path>Unpacks the contents of a zip file
- zip -r <outputpath> <contentpath> [<contentpath> ...]

  Add content to a zip file recursively. Creates the zip file if it does not exist.

```
$ unzip -l material.zip
Archive: material.zip
  Length
              Date
                      Time
                             Name
        0 03-29-2017 09:36
                             material/
          03-22-2017 10:59
                             material/test_file_1.txt
          03-22-2017 10:59
                             material/test_file_2.txt
                             material/large.fasta
    34753
          03-22-2017 12:36
          03-22-2017 12:39
                             material/small.fasta
                             material/duplicated_file.txt
      595
          03-28-2017 08:58
          03-29-2017 09:29
                             material/my_diff_2.txt
                             material/my_diff_1.txt
          03-29-2017 09:29
          03-29-2017 09:34
                             material/my_sort_1.txt
          03-29-2017 09:36
                             material/tmp.txt
                             material/my_sort_2.txt
      557
          03-29-2017 09:36
    36765
                              11 files
```

```
$ unzip material.zip ←
Archive: material.zip
   creating: material/
  inflating: material/test_file_1.txt
  inflating: material/test_file_2.txt
  inflating: material/large.fasta
  inflating: material/small.fasta
  inflating: material/duplicated_file.txt
 extracting: material/my_diff_2.txt
  inflating: material/my_diff_1.txt
  inflating: material/my_sort_1.txt
  inflating: material/tmp.txt
  inflating: material/my_sort_2.txt
```

```
$ unzip material.zip ←
Archive: material.zip
   creating: material/
  inflating: material/test_file_1.txt
  inflating: material/test_file_2.txt
  inflating: material/large.fasta
  inflating: material/small.fasta
  inflating: material/duplicated_file.txt
 extracting: material/my_diff_2.txt
  inflating: material/my_diff_1.txt
  inflating: material/my_sort_1.txt
  inflating: material/tmp.txt
  inflating: material/my_sort_2.txt
$ ls -F ⊿
Desktop/ Documents/ Exercise/ material/ material.zip
```

```
$ unzip material.zip ←
Archive: material.zip
   creating: material/
  inflating: material/test_file_1.txt
  inflating: material/test_file_2.txt
  inflating: material/large.fasta
  inflating: material/small.fasta
  inflating: material/duplicated_file.txt
 extracting: material/my_diff_2.txt
  inflating: material/my_diff_1.txt
  inflating: material/my_sort_1.txt
  inflating: material/tmp.txt
  inflating: material/my_sort_2.txt
$ ls -F ⊿
Desktop/ Documents/ Exercise/ material/
                                           material.zip
$ 1s material
duplicated_file.txt my_diff_1.txt my_sort_1.txt small.fasta
                                                                   test_file_2.txt
large.fasta
                    my_diff_2.txt my_sort_2.txt test_file_1.txt
                                                                   tmp.txt
```

```
$ zip -r new_material.zip material/ 
adding: material/ (stored 0%)
adding: material/my_diff_2.txt (stored 0%)
adding: material/test_file_2.txt (deflated 41%)
adding: material/duplicated_file.txt (deflated 92%)
adding: material/test_file_1.txt (deflated 63%)
adding: material/my_sort_1.txt (deflated 3%)
adding: material/small.fasta (deflated 45%)
adding: material/tmp.txt (deflated 52%)
adding: material/my_sort_2.txt (deflated 51%)
adding: material/my_diff_1.txt (deflated 4%)
adding: material/large.fasta (deflated 77%)
```

• tar tzf <path>

Shows the contents of a gziped tar file

- tar tzf <path>
   Shows the contents of a gziped tar file
- tar xvzf <path>
   Unpacks the contents of a gziped tar file. Overwrites existing output file paths!

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- tar xvzf <path>
   Unpacks the contents of a gziped tar file. Overwrites existing output file paths!
- tar cvzf <outputpath> <contentpath> [<contentpath> ...]
   Add content to a zip file recursively. Overwrites an existing output zip file if it exists.

```
$ tar tzf material.tar.gz 
material/
material/test_file_2.txt
material/tmp.txt
material/my_sort_1.txt
material/test_file_1.txt
material/small.fasta
material/my_diff_2.txt
material/my_sort_2.txt
material/my_diff_1.txt
material/my_diff_1.txt
material/duplicated_file.txt
material/large.fasta
```

```
$ tar xvzf material.tar.gz 
material/
material/test_file_2.txt
material/tmp.txt
material/my_sort_1.txt
material/test_file_1.txt
material/small.fasta
material/small.fasta
material/my_diff_2.txt
material/my_sort_2.txt
material/my_diff_1.txt
material/duplicated_file.txt
material/large.fasta
```

```
$ tar cvzf new_material.tar.gz material/ 
material/
material/my_diff_2.txt
material/test_file_2.txt
material/duplicated_file.txt
material/test_file_1.txt
material/my_sort_1.txt
material/small.fasta
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material/large.fasta
```

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- TAR files are sometimes compressed with compression tools other than gzip.
- For .tar.bz2 files, use the j switch instead of the z switch
- For .tar.xz files, use the J switch instead of the z switch
- In fact, recent versions of tar will detect the compression algorithm automatically:

\$ tar xf material.tar.gz  $\rightleftarrows$ 

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Compresses the given file into <path>.gz and removes the original file

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- gzip is commonly used to compress all kinds of files
- gzip <path>
   Compresses the given file into <path>.gz and removes the original file
- gunzip <path>.gz
   Decompresses the given file into <path> and removes the compressed file

## Download a compressed FASTQ file

```
$ curl -LO http://compbio.de/vorkurs/SRR6800317.fastq.gz ←
 % Total
           % Received % Xferd Average Speed Time
                                                Time
                                                        Time
                                                             Current
                            Dload Upload Total
                                                Spent
                                                        Left Speed
100
     330
        100
              330
                             6346
                                     0 --:--:- 6470
100 13.5M
         100 13.5M
                    0
                            9403k
                                        0:00:01 0:00:01 --:--: 11.1M
```

## Download a compressed FASTQ file

```
$ curl -LO http://compbio.de/vorkurs/SRR6800317.fastq.gz ←
 % Total
           % Received % Xferd Average Speed
                                           Time
                                                  Time
                                                          Time
                                                               Current
                             Dload Upload
                                           Total
                                                  Spent
                                                          Left Speed
     330
         100
               330
                              6346
                                       0 --:--:- 6470
100
         100 13.5M
100 13.5M
                             9403k
                                         0:00:01 0:00:01 --:--: 11.1M
```

#### Inspect the downloaded file

```
$ ls -lh SRR6800317.fastq.gz ↓
-rw-r--r-- 1 vorkurs vorkurs 14M Apr 3 2018 SRR6800317.fastq.gz
```

## Download a compressed FASTQ file

```
$ curl -LO http://compbio.de/vorkurs/SRR6800317.fastg.gz ←
 % Total
           % Received % Xferd Average Speed
                                          Time
                                                  Time
                                                          Time
                                                               Current
                             Dload
                                  Upload
                                          Total
                                                  Spent
                                                          Left Speed
         100
              330
                             6346
                                      0 --:--:- 6470
100
     330
100 13.5M
         100 13.5M
                            9403k
                                         0:00:01 0:00:01 --:--: 11.1M
```

#### Inspect the downloaded file

```
$ ls -lh SRR6800317.fastq.gz ↓
-rw-r--r-- 1 vorkurs vorkurs 14M Apr 3 2018 SRR6800317.fastq.gz
```

## Decompress the downloaded file

```
$ gunzip SRR6800317.fastq.gz ←
```

## Download a compressed FASTQ file

```
$ curl -LO http://compbio.de/vorkurs/SRR6800317.fastg.gz ←
 % Total
           % Received % Xferd Average Speed
                                           Time
                                                   Time
                                                           Time
                                                                Current
                                   Upload
                                           Total
                             Dload
                                                   Spent
                                                           Left Speed
         100
               330
                              6346
                                       0 --:--:- 6470
100
100 13.5M
         100 13.5M
                             9403k
                                          0:00:01 0:00:01 --:--: 11.1M
```

#### Inspect the downloaded file

```
$ ls -lh SRR6800317.fastq.gz ↓
-rw-r--r- 1 vorkurs vorkurs 14M Apr 3 2018 SRR6800317.fastq.gz
```

## Decompress the downloaded file

```
$ gunzip SRR6800317.fastq.gz ←
```

## Inspect the decompressed file

```
$ ls -lh SRR6800317.fastq ←
-rw-r--r- 1 vorkurs vorkurs 89M Apr 3 2018 SRR6800317.fastq
```

- du <path> disk usage
   Can be used to show the sizes of individual files or the total size of entire directories
  - du -h <path>
     Shows the sizes in a human readable format instead of kb
  - du -a <path>
     When applied on a directory, report sizes of all files, not only folders.

• du and ls may report different sizes

```
$ du -h material.zi ←|p

12K material.zip

$ ls -lh material.zip ←|
-rw-r--r-- 1 vorkurs vorkurs 11K Apr 3 22:05 material.zip
```

• du and ls may report different sizes

- The tools define the size of a file differently
  - du reports how much space the file consumes on the underlying storage
  - **ls** reports the size of the content stored inside the file

#### File sizes

• du and ls may report different sizes

```
$ du -h material.zi ←|p

12K material.zip

$ ls -lh material.zip ←|
-rw-r--r-- 1 vorkurs vorkurs 11K Apr 3 22:05 material.zip
```

- The tools define the size of a file differently
  - du reports how much space the file consumes on the underlying storage
  - 1s reports the size of the content stored inside the file
- When du is invoked with the flag --apparent-size, it too reports the size of the content

# File comparison

### File comparison

#### diff <path1> <path2>

Identifies differences between two plain text files and reports them as the smallest number of insertions and deletions required to transform one file into the other.

- diff -i or diff --ignore-case
  Ignore case differences in the files contents
- diff -w or diff --ignore-all-space
   Ignore all differences that involve only white space
- diff -B or diff --ignore-blank-lines
   Ignore blank lines
- diff -y or diff --side-by-size
   Output a human readable, side-by-side view instead of machine readable output

## File comparison

Use diff to identify the differences between the files  $my\_diff\_1.txt$  and  $my\_diff\_2.txt!$ 

- sort <path> sort plain text
   The sort tool can be used to sort plain text files line by line.
  - sort -b or sort --ignore-leading-blanks
    Ignore leading white space caracters
  - sort -n or sort --numeric-sort
     Interpret digits numerically instead of lexicographically
  - sort -r or sort --reverse
    Sort in reverse order
  - sort -u or sort --unique
     Output redundant lines only once

uniq <path>

Collapse consecutive identical lines into one line.

- uniq -c or uniq --count
   Report the number of occurrences with each line
- uniq -i or uniq --ignore-case
  Ignore case differences when comparing lines

- wc <path> word count
   Count words (or lines, characters, bytes) in plain text
  - wc -c
    Report the number of characters
  - wc -1 Report the number of lines
  - wc -w
    Report the number of words
  - wc -c
    Print all three numbers

- Sort the file my\_sort\_1.txt!
- Sort the file my\_sort\_2.txt in reverse numerical order!
- Which lines in duplicated\_file.txt are duplicated and how often do they occur?
- Are all globally redundant lines collapsed by **uniq**?

• grep <path> <path> [<path> ...]

Prints all lines of the input that match the given regular expression pattern

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- grep <path> <path> [<path> ...]

  Prints all lines of the input that match the given regular expression pattern
  - grep -v <path> (<path> ...)
     Inverse mode, print all lines that do not match the given pattern
  - grep -l <path> (<path> ...)

    Print only the names of files that the pattern matches against
  - grep -n <path> (<path> ...)

    Print the line numbers along with the matches
  - grep -color <path> (<path> ...)
    Highlight the pattern occurrence

```
$ grep --color -n is sortme.txt ←
1:This is an example file with content
2:that may be sorted. If you sort this file
4:because the line order is critical. None
5:of this makes any sense!
```

• The | (pipe) character can be used to redirect the standard output stream of one process to the standard input stream of another process

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- Remember the common structure of command line interfaces that we have seen often today:

COMMAND [OPTIONAL SWITCHES] [INPUT PATH]

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- Remember the common structure of command line interfaces that we have seen often today:

#### COMMAND [OPTIONAL SWITCHES] [INPUT PATH]

 Most programs we have seen today would process data from the standard input stream if no input file path is given!

#### Without stream redirection:

```
$ ls -l 1> temp.txt ↓
$ wc -l temp.txt ↓
15 temp.txt
```

#### Without stream redirection:

```
$ ls -l 1> temp.txt ↓
$ wc -l temp.txt ↓
15 temp.txt
```

#### With stream redirection:

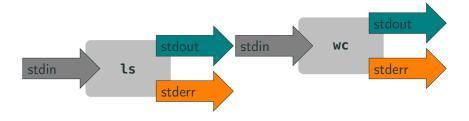
```
$ ls -l | wc -l ←
15
```

#### Without stream redirection:

```
$ ls -l 1> temp.txt ← $ wc -l temp.txt ← 15 temp.txt
```

#### With stream redirection:

```
$ ls -l | wc -l ↓
15
```



#### Recall:

```
$ uniq -c duplicated_file.txt ↓

22 bla

28 blabla

1 this line is not duplicatedbla

21 bla

28 blabla
```

## Combining **sort** and **uniq**:

```
$ sort duplicated_file.txt | uniq -c ↓

43 bla

56 blabla

1 this line is not duplicatedbla
```

What does this do and why?

\$ gunzip -c material.tar.gz | tar t ↓