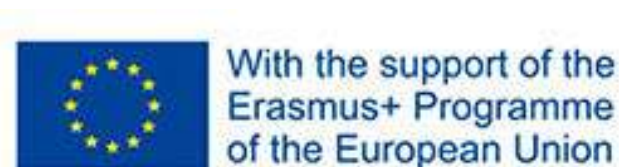


Introduction to GNU/Linux

Day 1: 19/02/2024



Luis J. Chueca

Postdoctoral researcher



@LuisjaChueca

Basque Centre for Climate Change (BC3)



luisjavier.chueca@bc3research.org

Sofia Marcos

Postdoctoral researcher

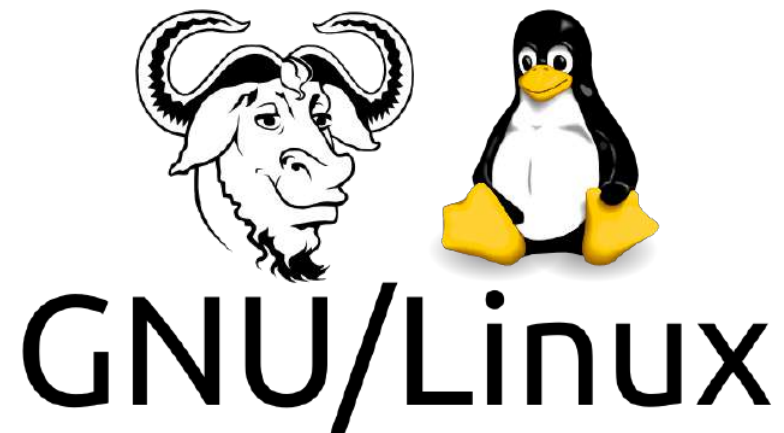


sofia.marcos@ehu.eus

University of the Basque Country (UPV/EHU)



Introduction





Introduction

1969

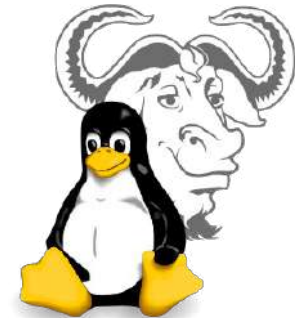
UNIX[®]
An Open Group Standard

1983



GNU's Not Unix (GNU)

1991



GNU/LINUX



debian

1993



Red Hat
Enterprise Linux

2003

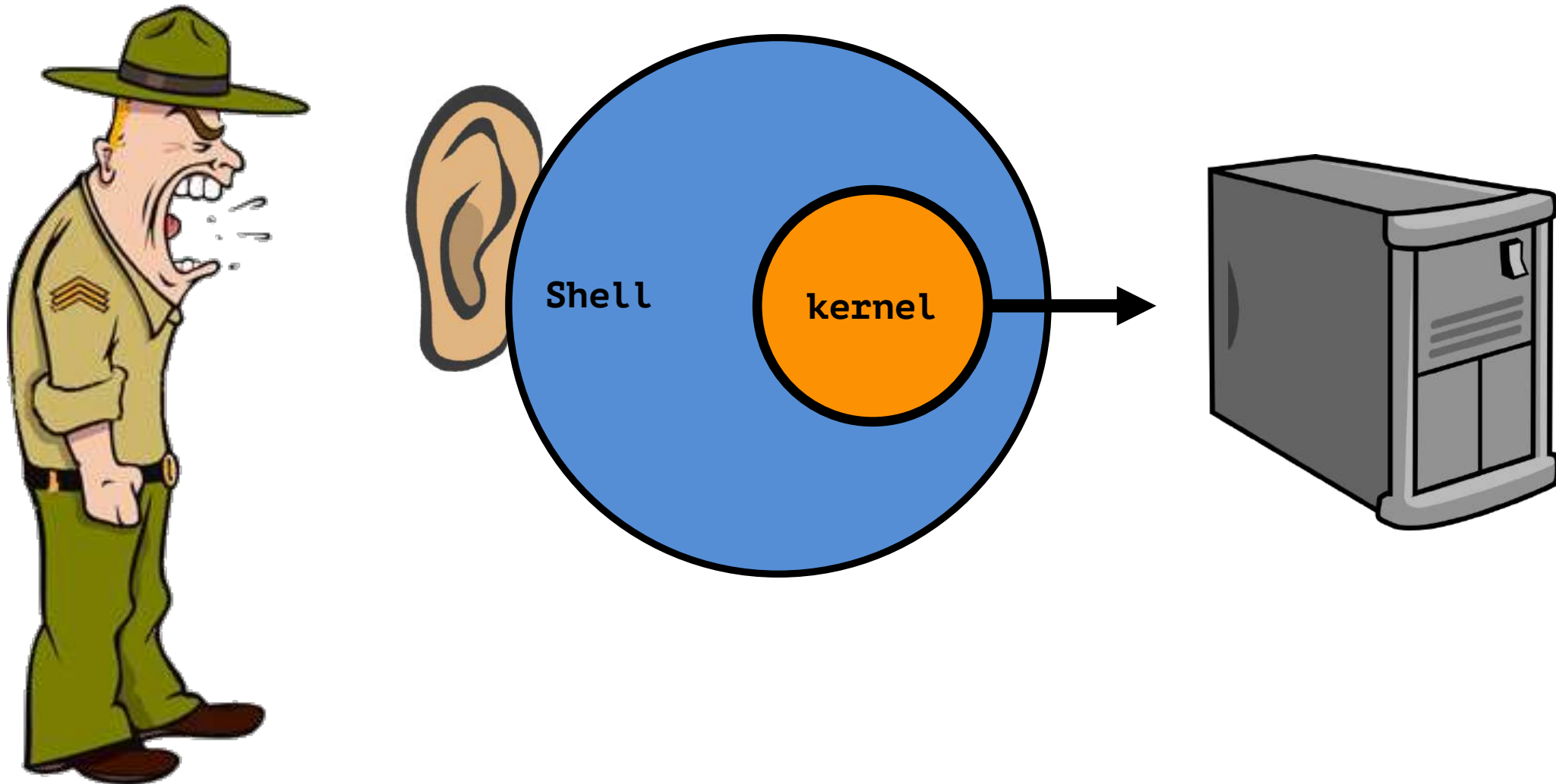


Ubuntu

2004

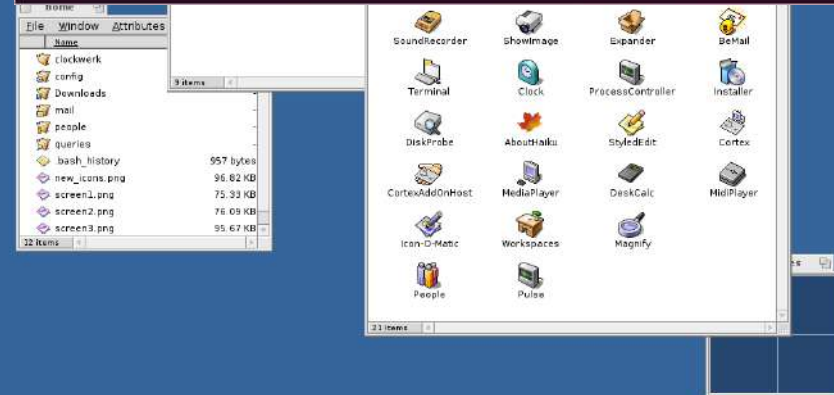
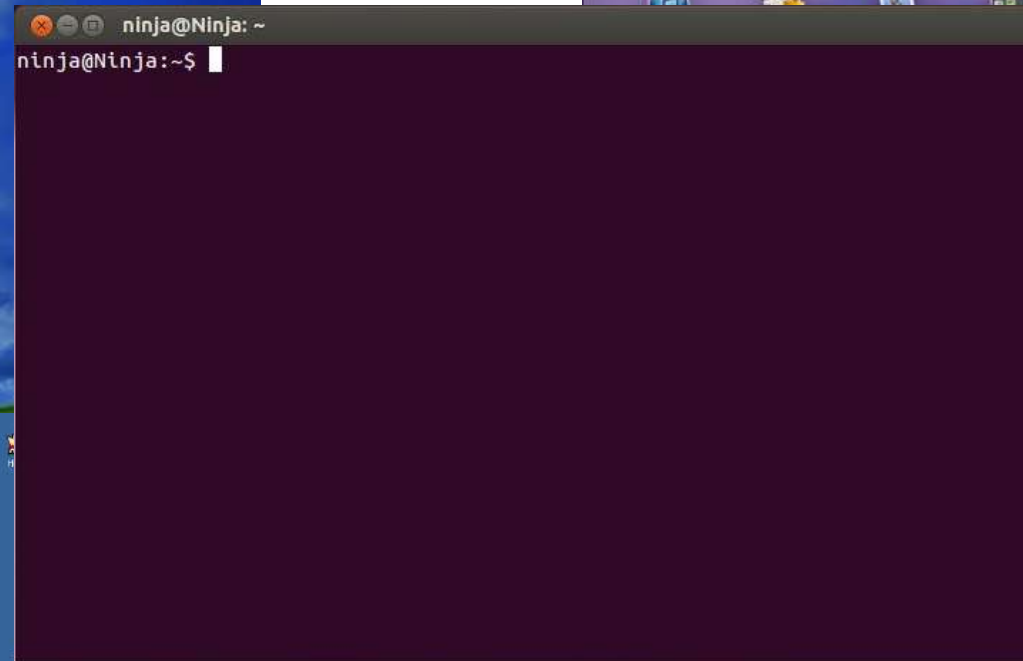


Shell concept



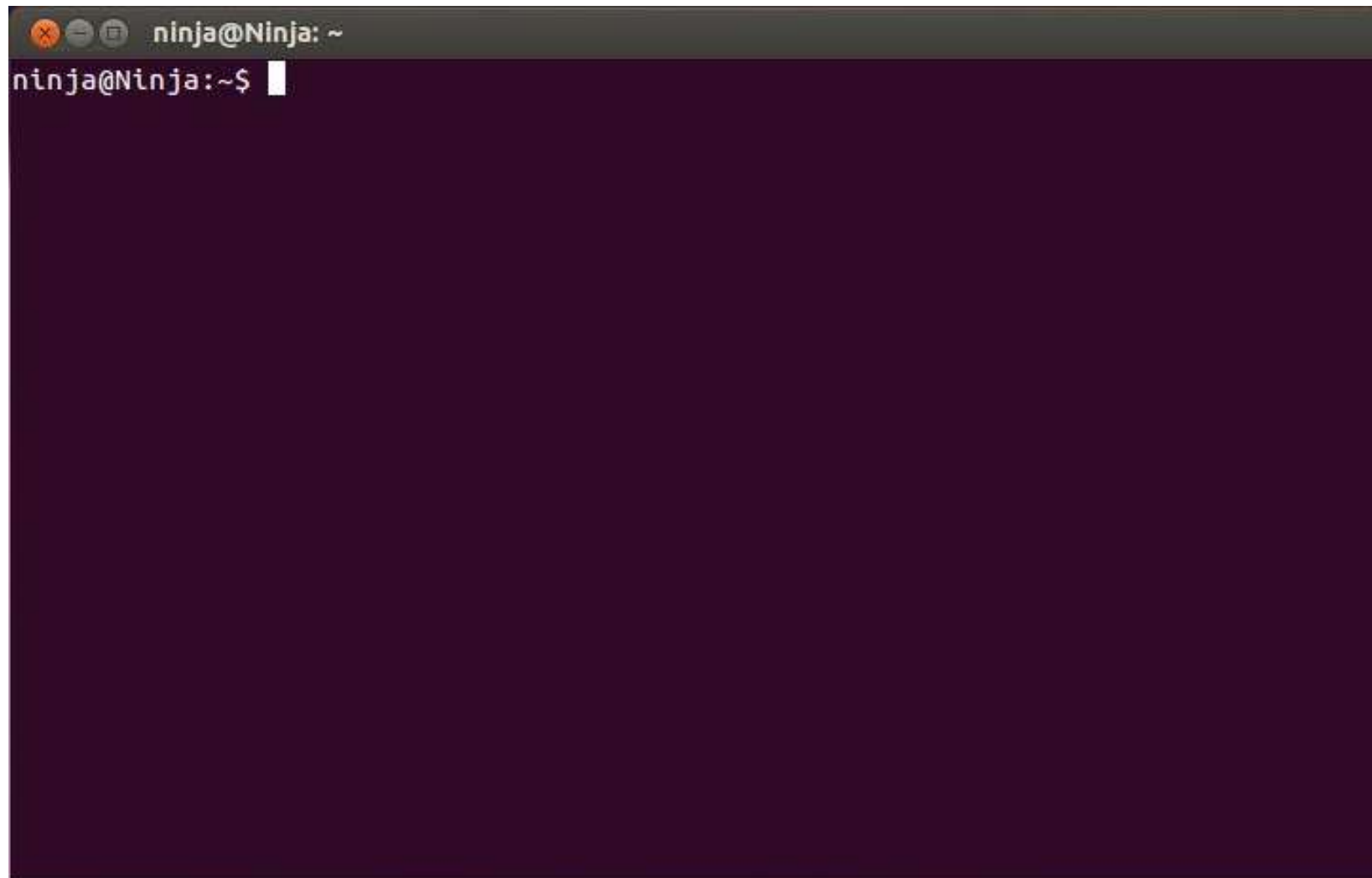


Examples of shells





The prompt



```
ninja@Ninja: ~  
ninja@Ninja:~$
```

A terminal window with a dark purple background. The title bar shows 'ninja@Ninja: ~'. The prompt 'ninja@Ninja:~\$' is displayed on the first line, followed by a white cursor block.

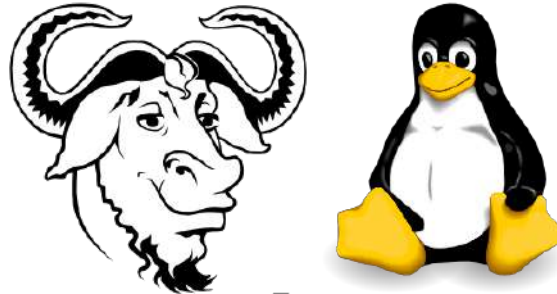
The prompt is highly configurable. Your prompt will look different than mine



Advantages of GNU/Linux

Free

Huge pool of free software



Multitasking


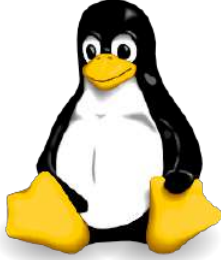
open source

GNU/Linux

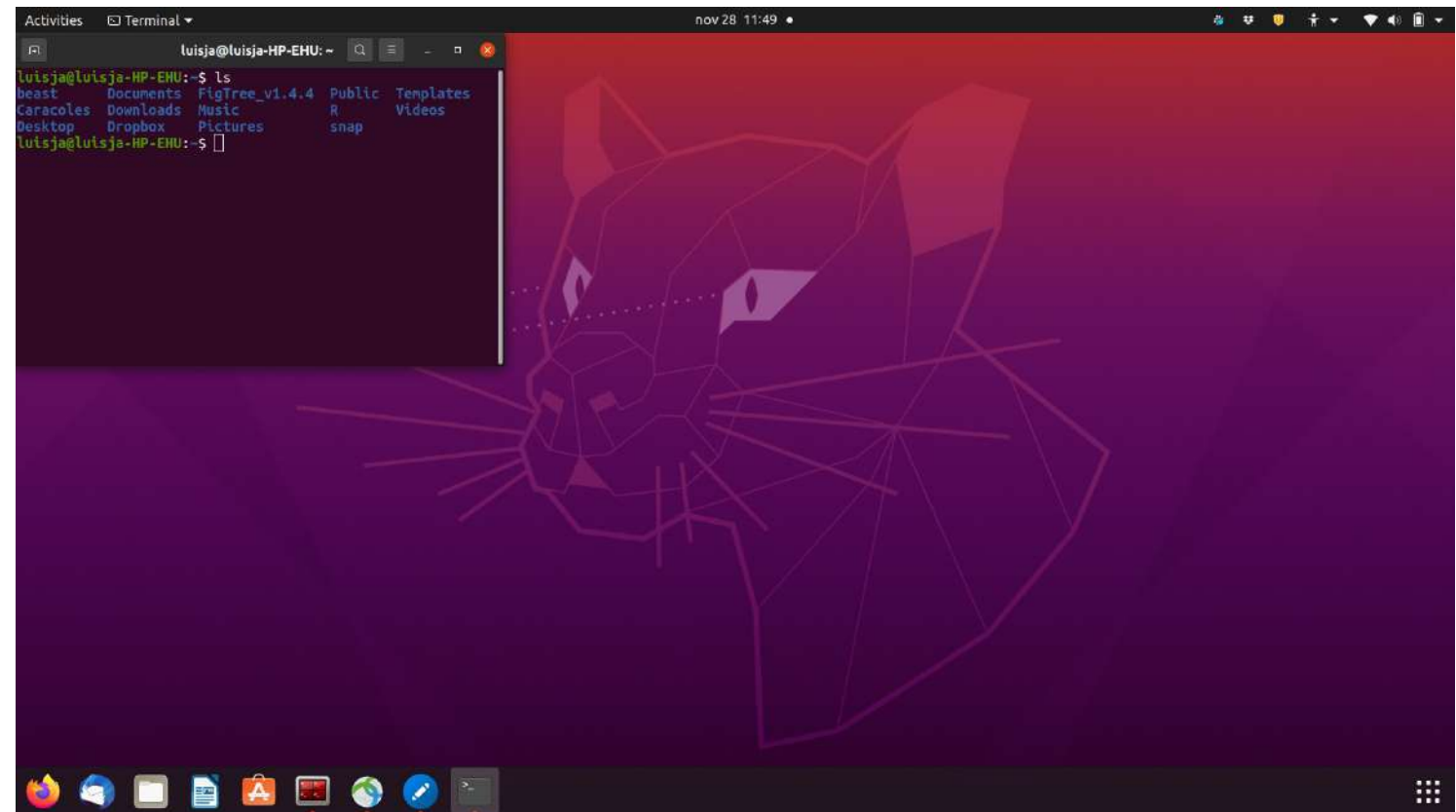
High stability

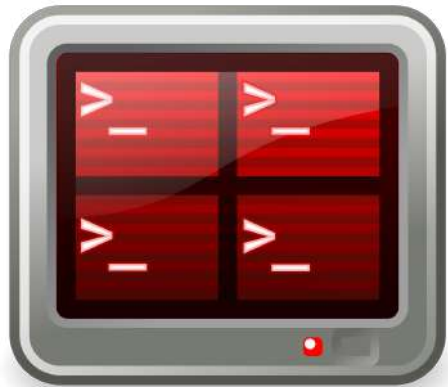
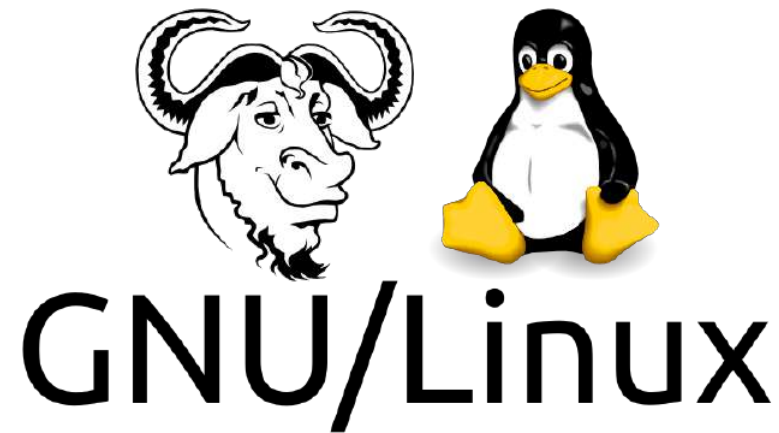
Easy administration

Multi



GNU/Linux



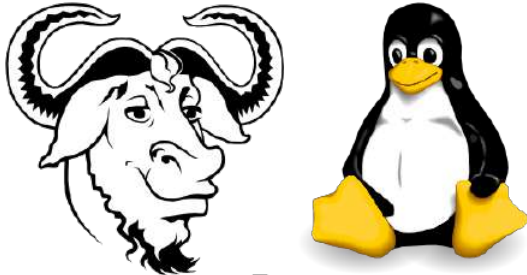


GNOME Terminator

Debian / Ubuntu
Fedora
Gentoo
(...)



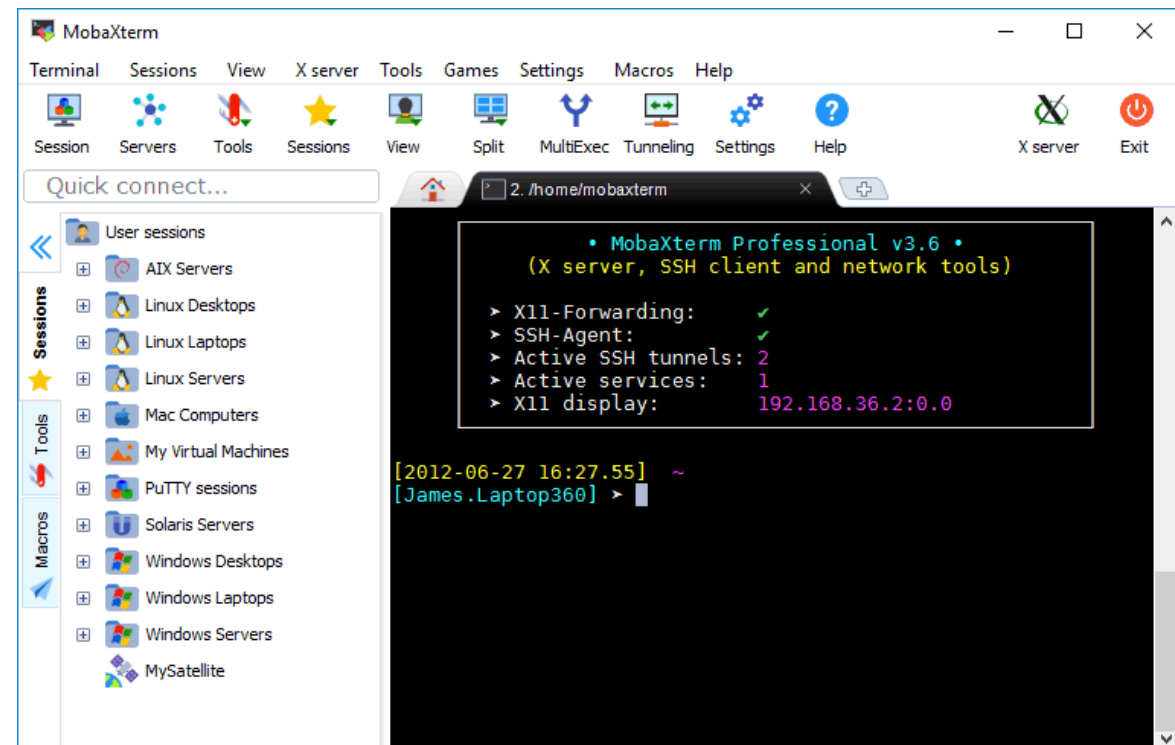
MacOS X

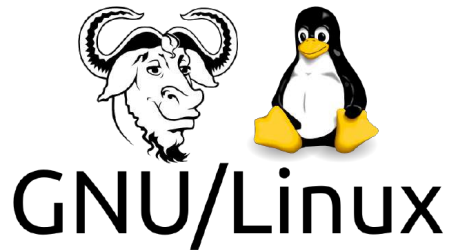


GNU/Linux



MobaXterm





Connect to another computer



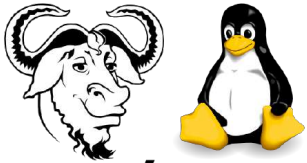
Gaming-PC:

- 8 Cores
- 16 GB RAM
- 250 GB SSD + 2 TB HDD
- Fancy graphic card
- ~ 2.000€



High-performance computer:

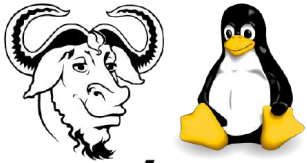
- 72 Cores
- 750 GB RAM
- 15 TB HDD im RAID
- ~ 15.000€



GNU/Linux

ssh connexion

```
$ ssh -Y ikasleXX@kalk2020.lgp.ehu.es  
$ password:
```



GNU/Linux

ssh connexion

```
$ ssh -Y ikasleXX@kalk2020.lgp.ehu.es
```

```
$ password:
```

```

Kaixo ikasle11, Ongi etorri Arinara
Hola ikasle11, Bienvenido a Arina
Hi ikasle11, Welcome to Arina

```

```
=== HOST ===
kalk2020
```

```
=== DISK USAGE ===
```

You are using: 0.3 GB.

```
Your group is using: 6.6 GB.
```

Info updated at night.

Above 3 GB of disk usage it is charged. Rates: <http://www.ehu.es/sgi/tarifas>

Cluster Administrators:

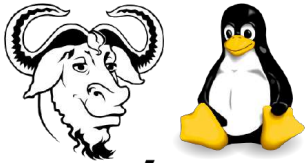
Joaquim Jornet Somoza

Luca Bergamini

For any doubt or question contact us at:

izo-sgi@ehu.eus

```
[ikasle11@kalk2020 ~]$
```



GNU/Linux

ssh connexion

```
$ ssh -Y ikasleXX@kalk2020.lgp.ehu.es
```

```
$ password:
```

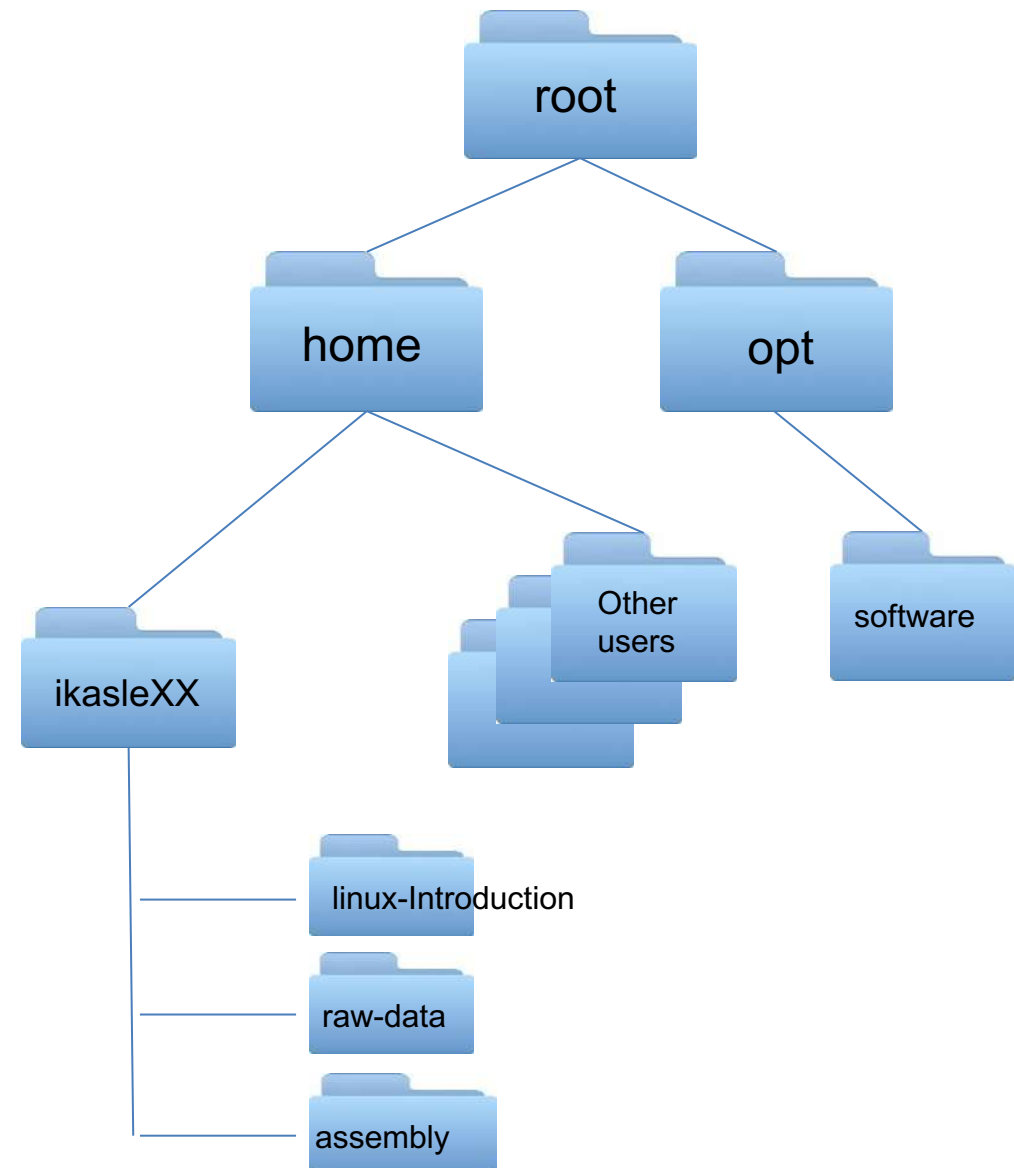
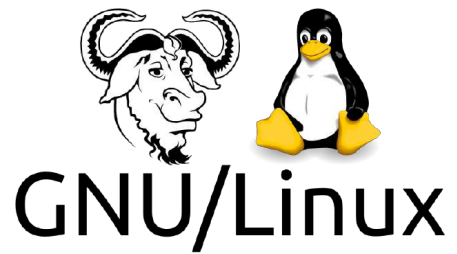
```

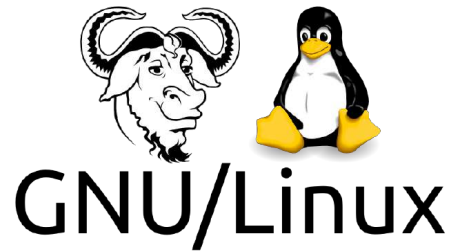
Kaixo ikasle11, Ongi etorri Arinara
Hola ikasle11, Bienvenido a Arina
Hi ikasle11, Welcome to Arina

```

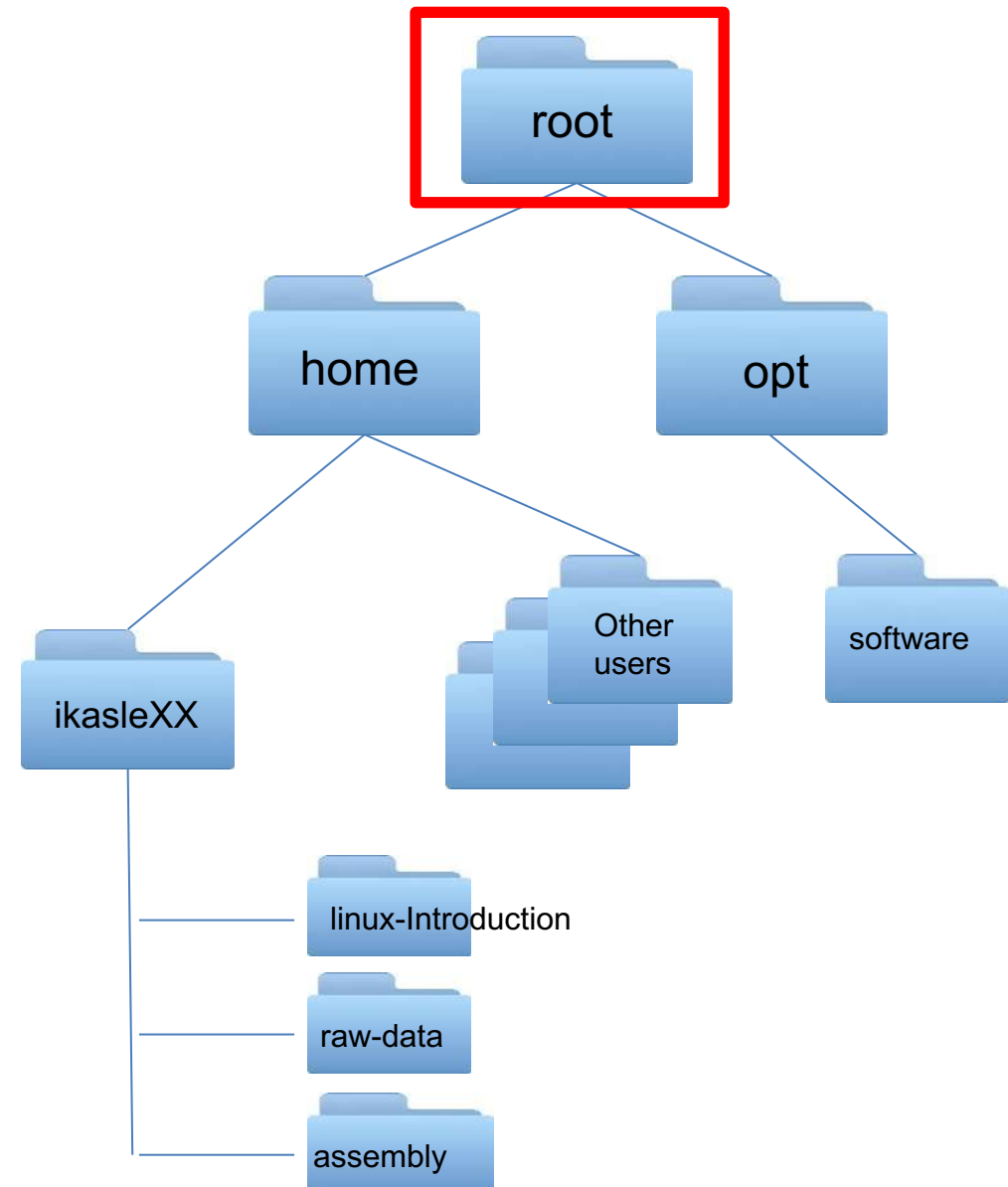
```
$ interactive -n 4 -p vfast -r mer1
```

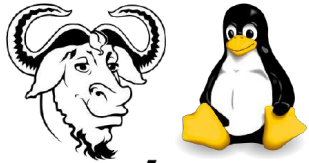
\$





Path of root directory:
/





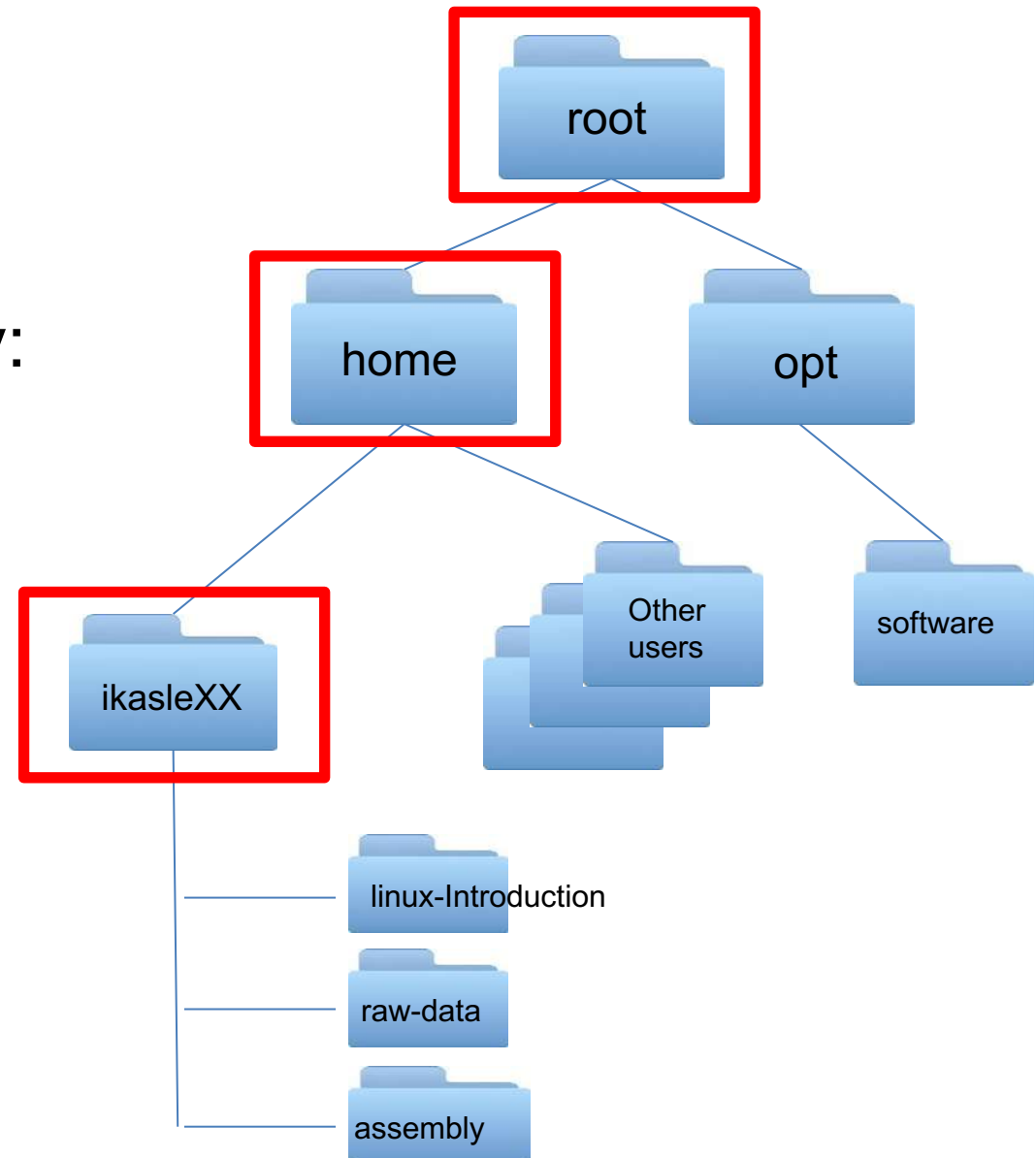
GNU/Linux

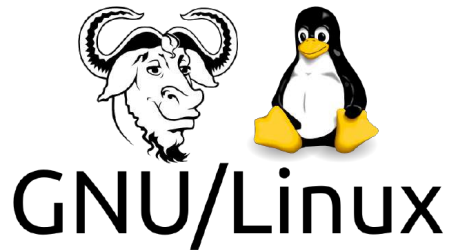
Path of your home-directory:

/home/ikasleXX/

or:

~





Main commands

Linux cheat sheet

Cheatography

Linux Command Line Cheat Sheet

by Dave Child (DaveChild) via cheatography.com/1/cs/49/

Bash Commands

<code>uname -a</code>	Show system and kernel
<code>head -n1 /etc/issue</code>	Show distribution
<code>mount</code>	Show mounted filesystems
<code>date</code>	Show system date
<code>uptime</code>	Show uptime
<code>whoami</code>	Show your username
<code>man command</code>	Show manual for <i>command</i>

Bash Shortcuts

CTRL-c	Stop current command
CTRL-z	Sleep program
CTRL-a	Go to start of line
CTRL-e	Go to end of line

Bash Variables (cont)

<code>export NAME=value</code>	Set \$NAME to <i>value</i>
<code>\$PATH</code>	Executable search path
<code>\$HOME</code>	Home directory
<code>\$SHELL</code>	Current shell

IO Redirection

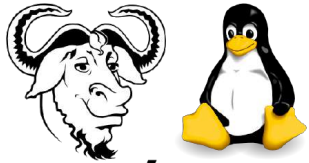
<code>cmd < file</code>	Input of <i>cmd</i> from <i>file</i>
<code>cmd1 <(cmd2)</code>	Output of <i>cmd2</i> as file input to <i>cmd1</i>
<code>cmd > file</code>	Standard output (stdout) of <i>cmd</i> to <i>file</i>
<code>cmd > /dev/null</code>	Discard stdout of <i>cmd</i>
<code>cmd >> file</code>	Append stdout of <i>cmd</i> to <i>file</i>

Command Lists

<code>cmd1 ; cmd2</code>	Run <i>cmd1</i> then <i>cmd2</i>
<code>cmd1 && cmd2</code>	Run <i>cmd2</i> if <i>cmd1</i> is successful
<code>cmd1 cmd2</code>	Run <i>cmd2</i> if <i>cmd1</i> is not successful
<code>cmd &</code>	Run <i>cmd</i> in a subshell

Directory Operations

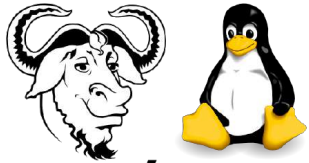
<code>pwd</code>	Show current directory
<code>mkdir dir</code>	Make directory <i>dir</i>
<code>cd dir</code>	Change directory to <i>dir</i>
<code>cd ..</code>	Go up a directory
<code>ls</code>	List files



GNU/Linux

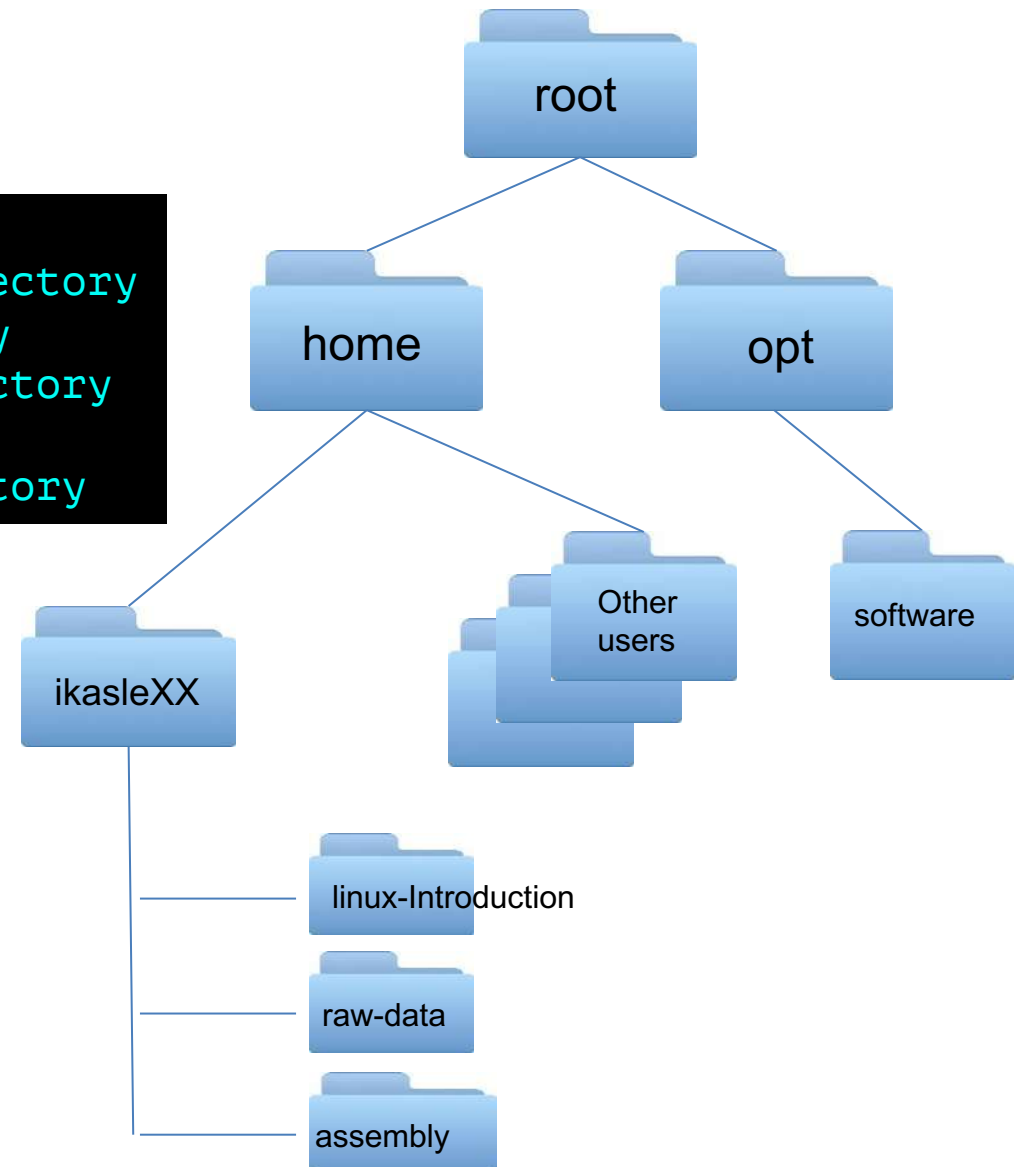
Main commands

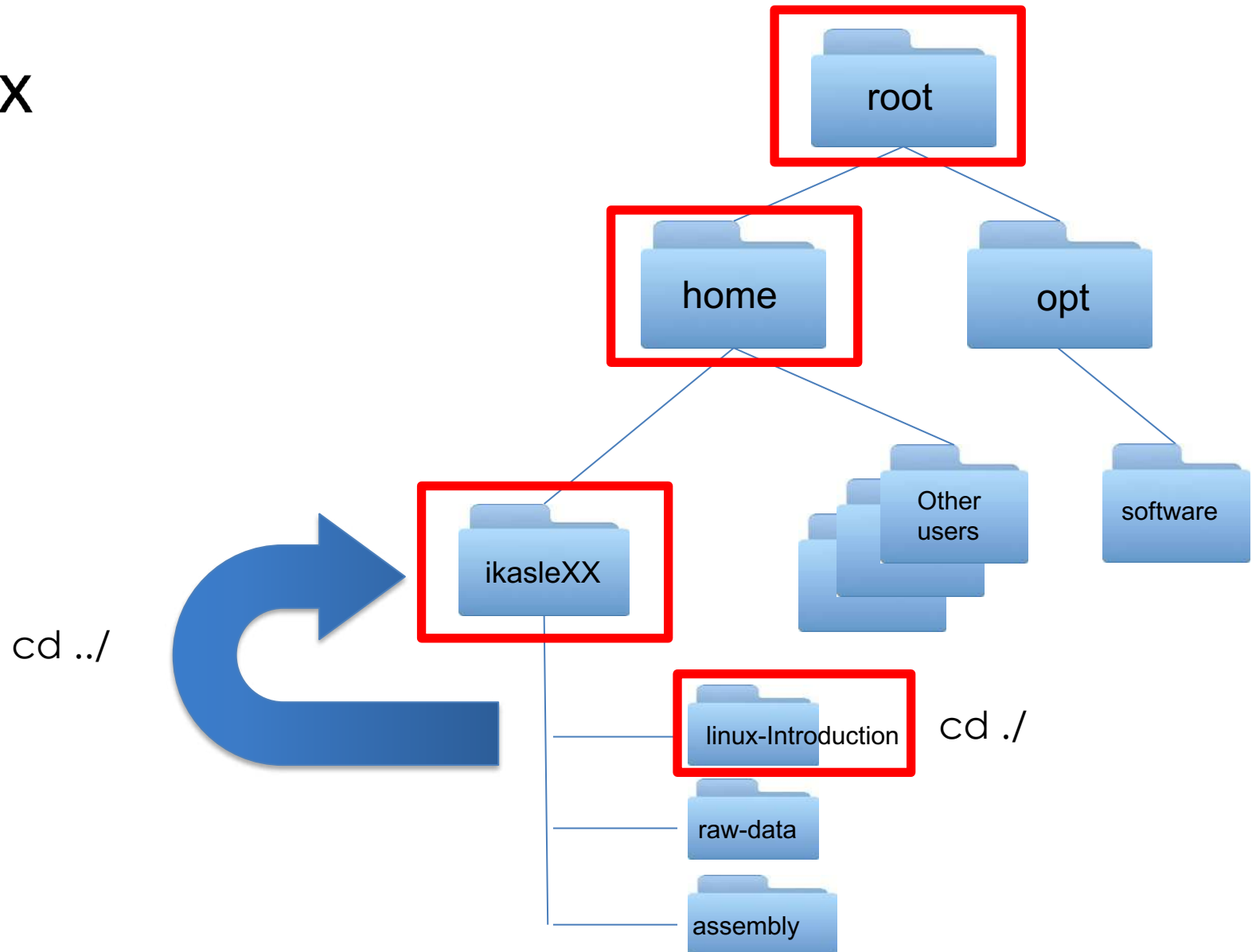
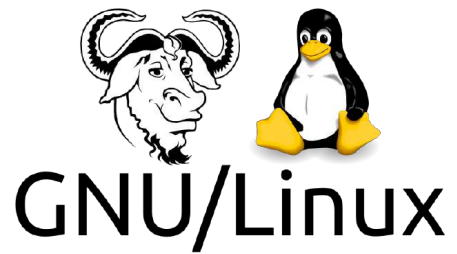
```
$ cd  
$ cd ../  
$ cd -  
$ mkdir  
$ w  
$ history  
$ top  
$ df  
$ mv  
$ vim  
$ less example.txt  
$ ln -s
```

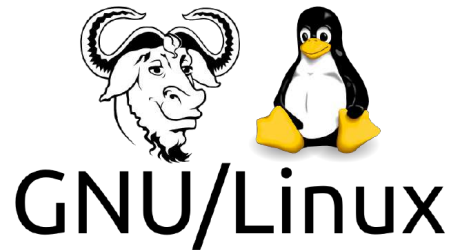


GNU/Linux

# Abbreviation	# Meaning
cd . or cd ./	# Current working directory
cd .. or cd ../	# Next upper directory
cd -	# Last "visited" directory
cd ~	# Home directory
pwd	# print working directory



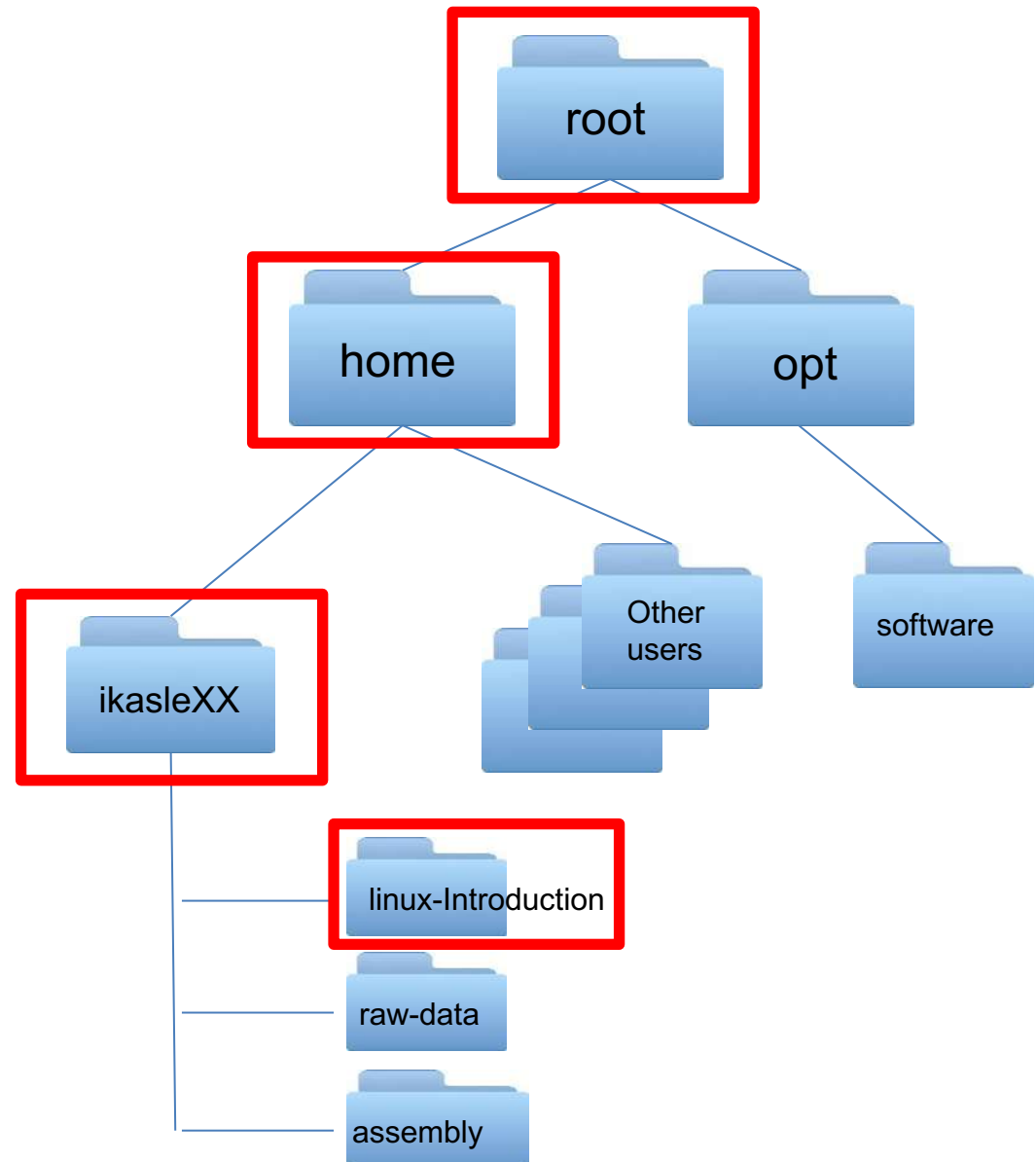


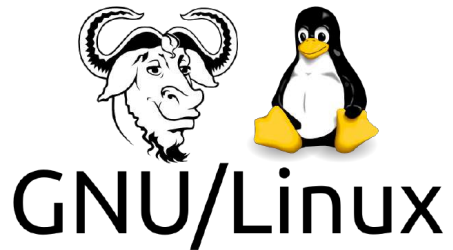


Working directory:
`/home/ikastleXX/linux-Intro`

ABSOLUTE PATH:
`/home/ ikastleXX/linux-Intro`

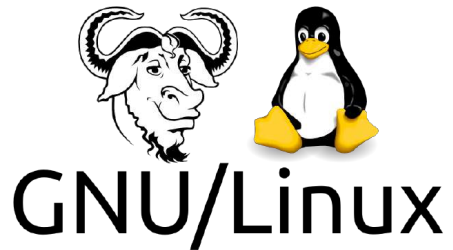
RELATIVE PATH:
`../ikastleXX/linux-Intro`





Paths

- **Absolute path:** The absolute path of a file always starts at the root directory represented by "/" :
/home/studentx/linux_introduction/exercise
- **Relative path:** The relative path of a file describes its location always relative to where the user is right now:
./linux_introduction/exercise



Your first commands!

Fundamental structure:

command [options] [argument]

Example 1:

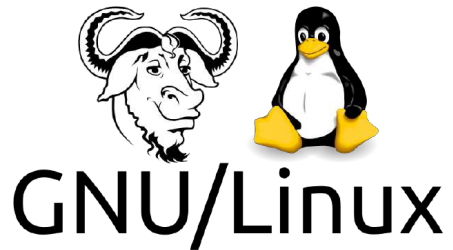
ls

Example 2:

ls -sh

Example 3:

ls -sh ../gscratch



Your first commands!

Fundamental structure:

command [options] [argument]

Example 1:

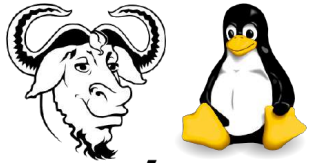
```
$ wc
```

Example 2:

```
$ wc -l
```

Example 3:

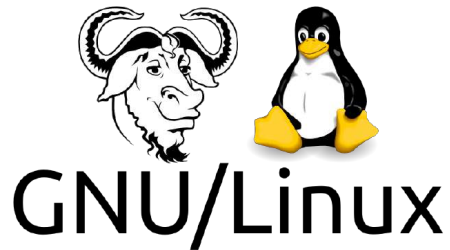
```
$ wc -l filename.txt
```



GNU/Linux

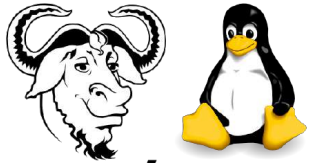
Exercise 1:

1. Open your terminal
2. What's the path to your current directory?
3. How many folders does this directory contain?
4. Navigate to `/home/username/ehu_genomic_data`
5. Navigate to your home directory
6. Make a new directory called „linux_intro“
7. Navigate into the „linux_intro“ directory



File handing commands

Command	Use
<code>cp [file1] [file2]</code>	Copy file1 and call it file2
<code>mv [file1] [file2]</code>	Rename file1 to file2
<code>mv [file] /path/to/directory</code>	Move file to directory
<code>rm [file]</code>	remove file
<code>rmdir [directory]</code>	remove directory
<code>cat [file]</code>	display file
<code>more [file] , less [file]</code>	display a file a page at a time
<code>head [file]</code>	display first 10 lines of a file
<code>tail [file]</code>	display last 10 lines of a file
<code>grep "[keyword]" [file]</code>	display all lines in file containing the keyword
<code>wc [file]</code>	count number of lines/words/characters in file



GNU/Linux

File naming conventions

Why “uebung” rather than “übung”?

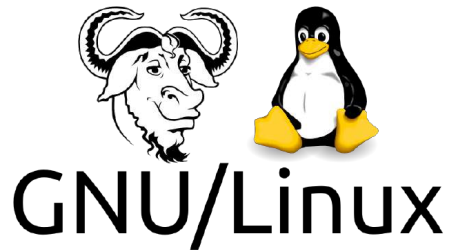
Why “ron_aniejo” rather than “ron_Añejo”?

Why “testfile” rather than “Testfile”?

Fundamentals:

- **Avoid** characters such as / * & %
- **NO WHITE SPACES!**
- **ONLY** use letters and numbers and replace whitespaces with the following special characters: _ . + -

e.g. chironomus-riparius+piger.fasta
 thats_my_logfile.txt
 thatsMyLogfile.txt



Create files

Example of built-in text editors on Unix command line:

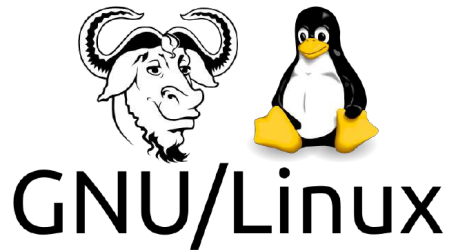
```
11LERRD). ;jD666666D}
.LG1t1000D, f6j}jL8000E;
1E :8888E1, ,G8888,
;1 E888, ,8888,
D888, :8888; 888
D888, :8888; 888 888Y88b 888 888 888
D888, :8888; 888 888888 888 Y888888 888 888
D888, :8888; 888 888 888 Y888888 888 888
D888, :8888; Y88b d88P 888 Y8888 Y88b, ,d88P
D888, :8888; "Y8888P88 888 Y888 "Y88888P"
W88W, :8888;
W88W, :8888;
D888, :8888; 888888, 888888, ,d888,
D888, :8888; 888 "88b "88b 888 "88b d88"888
D888, :8888; 888 888 ,d888888 888 888 888 888
W888, :8888; 888 888 888 888 888 888 Y88, d8P
D888, :8888; 888 888 "Y888888 888 888 "Y88P"
E888, :8888;
T8888
```

The

- nano



- vim



Create files

Create files in a plain text editor (not Word):



<https://notepad-plus-plus.org/>

Most popular more advanced code editors



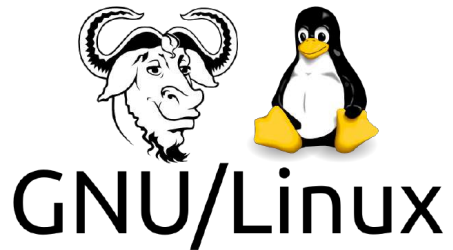
<https://code.visualstudio.com>



<https://atom.io>



<https://www.sublimetext.com>



Files inspection

See the whole file

```
$ cat <file>
```

Inspecting the file

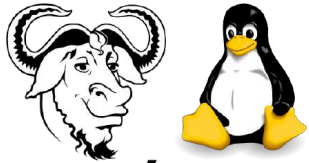
```
$ less <file>      # Within less search for an specific word typing "/"word"  
  
                  # Within less, exit typing q
```

Seeing the top of the file

```
$ head <file>      # head [- number] <file>
```

Seeing the end of the file

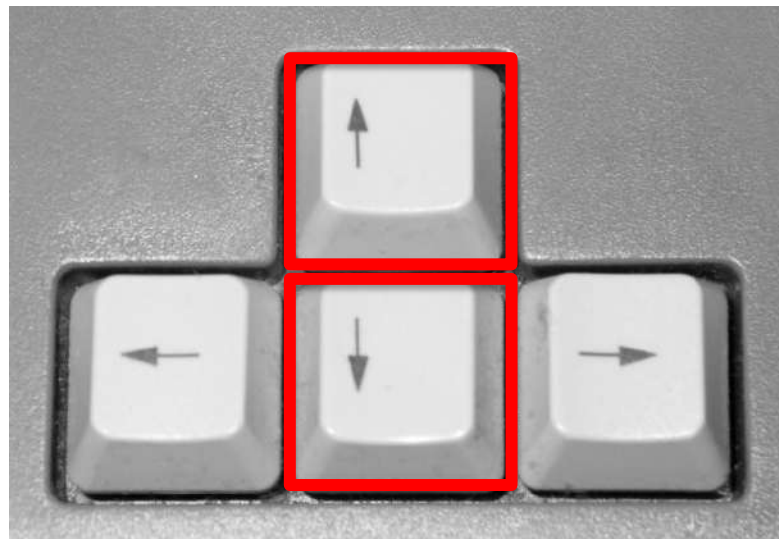
```
$ tail <file>      # tail [- number] <file>
```

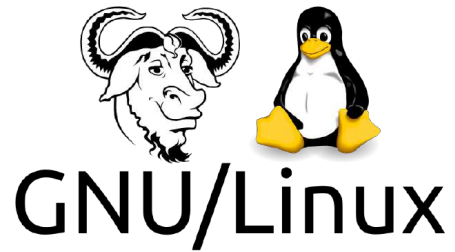


GNU/Linux

- Used commands → history

#gives you all the commands previously run in your session

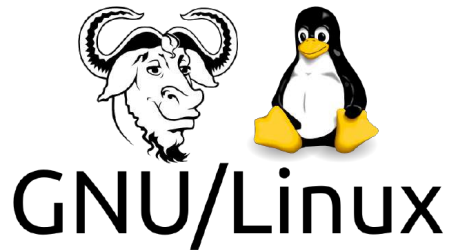




Autofill



What happens if you write/press **head Te[tab]**?



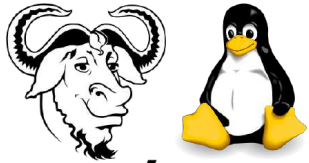
Deleting files and directories

Attention!!



The **system** always **assumes** the **user knows** what she/he is doing

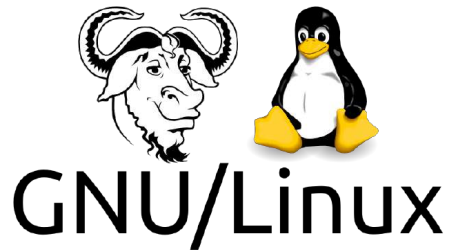
- No query whether you **REALLY** want to delete something ...
- No recycle bin – **Deleted means Deleted**
- Particular caution needed when using wildcards



GNU/Linux

Exercise 2:

1. Create a textfile called „delete_me.txt“ using *`touch [filename]`*.
2. Create a directory called „delete_me“.
3. Move „delete_me.txt“ to „delete_me“.
4. Delete the directory



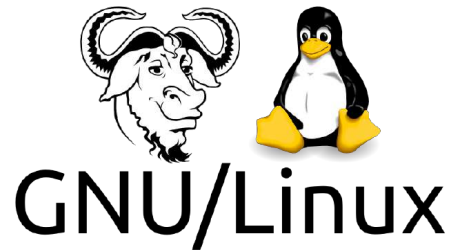
File permissions

```
ikasle01@kalk2020:~/gscratch/TEST2$ ll
total 9
drwxr-xr-x 1 ikasle01 pomaster 24 Nov 23 15:03 File.txt
```

d**r****w****x****r** - **x****r** - **x**

Type User Group Others

r	read
w	write
x	execute
-	denied



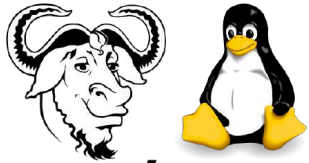
Save output

Instead of printing the output to the terminal, one can save it in a new file

Command (options) file > new_file

For example:

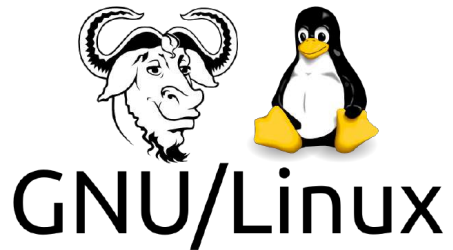
head inputfile.txt > outputfile.txt



GNU/Linux

Exercise 3:

- 1) Save the first line of the file
„WatsonCrick1953.txt“ as „papertitle.txt“
- 2) Save the remaining lines as „mainbody.txt“.

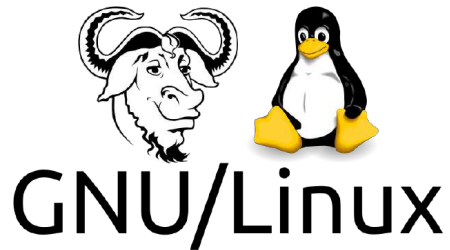


Piping commands

Unix offers the possibility to pipe commands. By doing this, the output of the first command serves as the input of the following one

How many sequences does twogenes.fa contain?

```
grep ">" twogenes.fa | wc -l
```



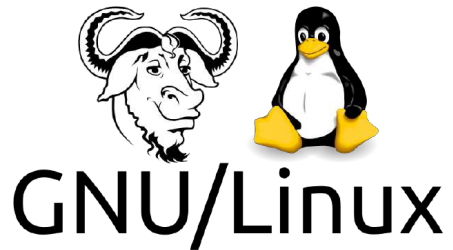
Piping commands

| head – check the output

“| head” offers the possibility to check the command/output

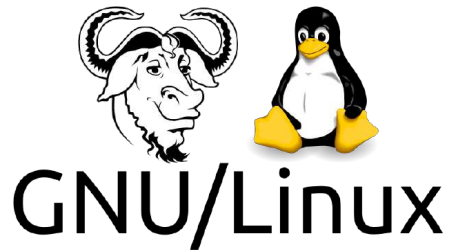
Avoids (unnecessarily) huge amounts of output files

Especially useful when working on huge files or using complex (piped) commands



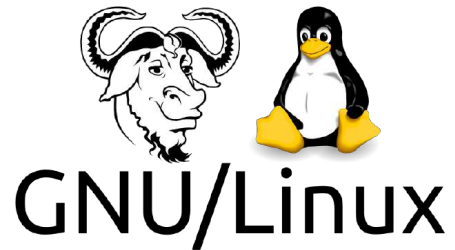
Listing 1. Example of cat to create a file:

```
ikasle01@kalk2020:~$ cat > species.list  
snail  
octopus  
mussel  
wolf  
<ctrl d>
```



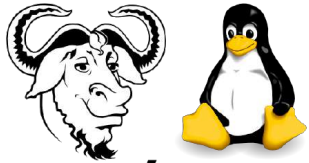
Listing 2. Example of cat to append a file:

```
ikastle01@kalk2020:~$ cat >> species.list  
wasp  
jellyfish  
<ctrl d>
```



Listing 3. Example of cat without flags:

```
ikasle01@kalk2020:~$ cat species.list  
snail  
octopus  
mussel  
wolf  
wasp  
jellyfish  
ikasle01@kalk2020:~$
```



GNU/Linux

Listing 4. Example of cat to count lines:

```
ikastle01@kalk2020:~$ cat -n species.list
```

```
1      snail
```

```
2      octopus
```

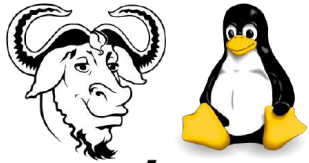
```
3      mussel
```

```
4      wolf
```

```
5      wasp
```

```
6      jellyfish
```

```
ikastle01@kalk2020:~$
```

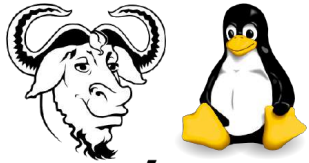


GNU/Linux

Use of 'wc'

The wc (wordcount) command counts the number of lines, words (separated by whitespace), and byte count in specified files, or from stdin.

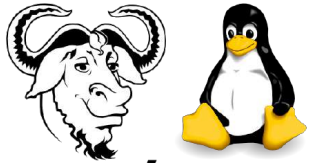
```
ikasle01@kalk2020:~$ wc species.list
6 6 41 species.list
ikasle01@kalk2020:~$ wc -l species.list
6 species.list
ikasle01@kalk2020:~$ wc -w species.list
6 species.list
ikasle01@kalk2020:~$ wc -c species.list
41 species.list
```



GNU/Linux

paste

```
$ paste species.list2 species.list > species.list3
$ cat species.list3
Gastropoda      snail
Cephalopoda     octopus
Bivalvia        mussel
Carnivora       wolf
Hymenoptera     wasp
Cnidaria        jellyfish
```

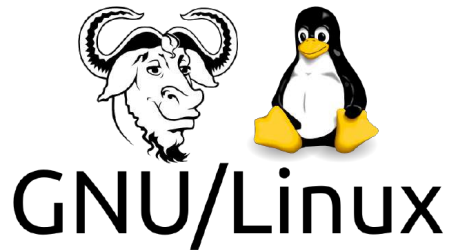


GNU/Linux

Use of 'grep'

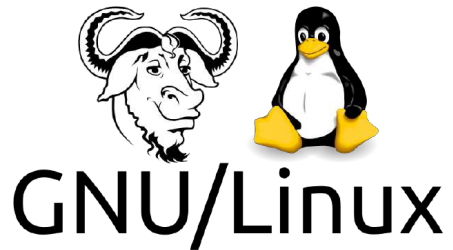
The grep command searches specified files or stdin for patterns matching a given expression(s). Output from grep is controlled by various option flags.

```
ikasle01@kalk2020:~$ grep Carnivora species.list3 species.list4
species.list3:Carnivora wolf
species.list4:Carnivora wolf
species.list4:Carnivora Tiger
species.list4:Fox Carnivora
species.list4:Carnivoras seal
```



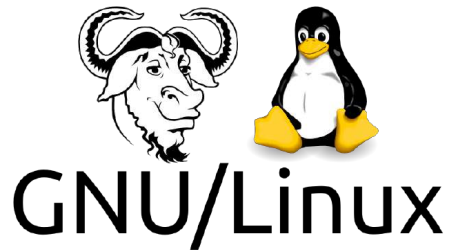
Example of grep – counting number of matches in files:

```
ikasle01@kalk2020:~$ grep -c Carnivora species.list3 species.list4
species.list3:1
species.list4:4
```

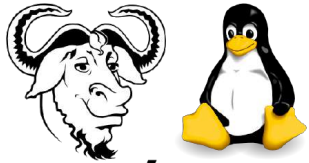
Example of grep – suppress filename in output:

```
ikasle01@kalk2020:~$ grep -h Carnivora species.list3 species.list4  
Carnivora    wolf  
Carnivora    wolf  
Carnivora    Tiger  
Fox    Carnivora  
Carnivoras  seal
```



Example of grep – case insensitive:

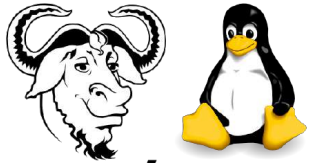
```
ikasle01@kalk2020:~$ grep -i Carnivora species.list3 species.list4
Carnivora    wolf
Carnivora    wolf
Carnivora    Tiger
Fox    Carnivora
carnivora    dingo
carnivoras   bear
Carnivoras   seal
```



GNU/Linux

Example of grep – inverted matching:

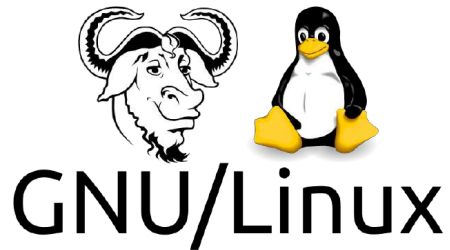
```
ikasle01@kalk2020:~$ grep -v Carnivora species.list3 species.list4
species.list3:Gastropoda      snail
species.list3:Cephalopoda    octopus
species.list3:Bivalvia      mussel
species.list3:Hymenoptera    wasp
species.list3:Cnidaria      jellyfish
species.list4:Gastropoda      snail
species.list4:Cephalopoda    octopus
species.list4:Bivalvia      mussel
species.list4:Hymenoptera    wasp
species.list4:Cnidaria      jellyfish
species.list4:Gastropoda      slug
species.list4:Helicoidea     Snail
species.list4:carnivora      dingo
species.list4:carnivoras     bear
```



GNU/Linux

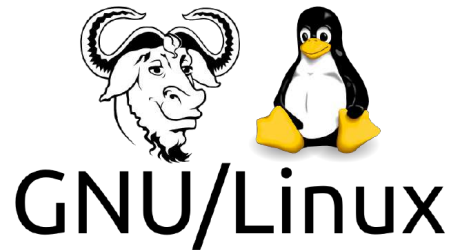
Example of grep – combining flags:

```
ikasle01@kalk2020:~$ grep -v -i Carnivora species.list3 species.list4
species.list3:Gastropoda snail
species.list3:Cephalopoda octopus
species.list3:Bivalvia mussel
species.list3:Hymenoptera wasp
species.list3:Cnidaria jellyfish
species.list4:Gastropoda snail
species.list4:Cephalopoda octopus
species.list4:Bivalvia mussel
species.list4:Hymenoptera wasp
species.list4:Cnidaria jellyfish
species.list4:Gastropoda slug
species.list4:Helicoidea Snail
```



Example of grep – word matching:

```
ikasle01@kalk2020:~$ grep -w Carnivora species.list3 species.list4
species.list3:Carnivora wolf
species.list4:Carnivora wolf
species.list4: Carnivora      Tiger
species.list4: Fox           Carnivora
```



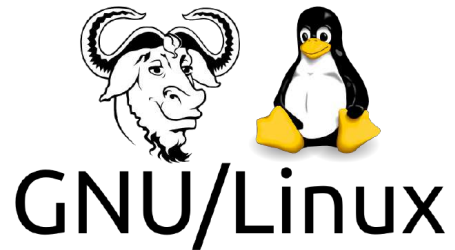
Use of 'awk'

Actual programming language

Input file is not changed

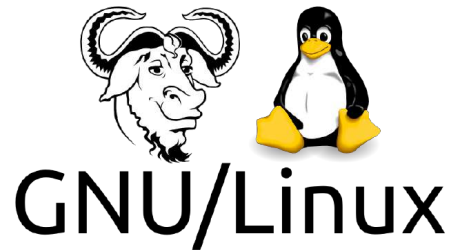
Especially useful when working with tables

Field-oriented



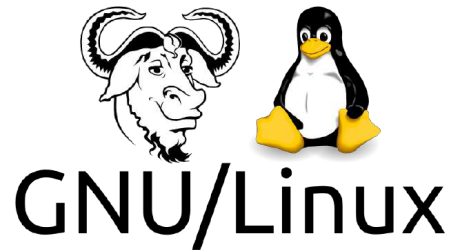
Use of 'awk'

```
ikasle01@kalk2020:~$ awk '{print$1}' species.list4 | head
Gastropoda
Cephalopoda
Bivalvia
Carnivora
Hymenoptera
Cnidaria
Gastropoda
Helicoidea
Carnivora
Fox
```



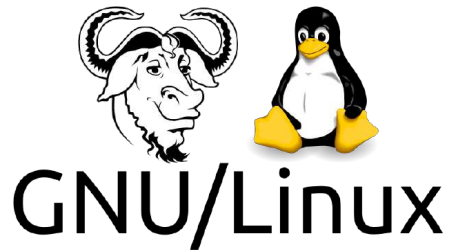
Use of 'awk'

```
ikasle01@kalk2020:~$ awk '{print$2}' species.list4 | head
snail
octopus
mussel
wolf
wasp
jellyfish
slug
Snail
Tiger
Carnivora
```

Use of 'awk'

```
ikasle01@kalk2020:~$ awk '{print$2,$1}' species.list4 | head
snail Gastropoda
octopus Cephalopoda
mussel Bivalvia
wolf Carnivora
wasp Hymenoptera
jellyfish Cnidaria
slug Gastropoda
Snail Helicoidea
Tiger Carnivora
Carnivora Fox
```



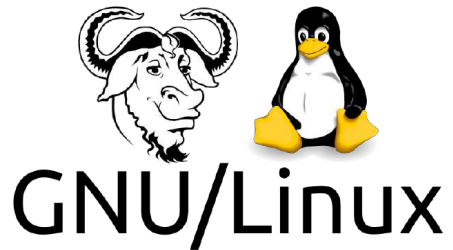
Other interesting commands

cut

Cuts parts of files based on certain criteria

```
cut -f [column_number]
```

```
cut -c [number_of_characters]
```



Other interesting commands

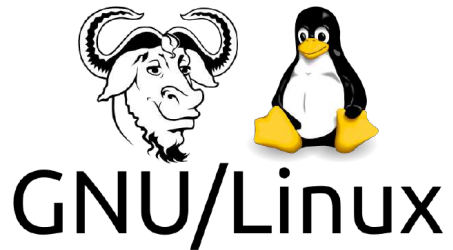
sort

Sorts files according to set criteria

`sort -k [beginning of Column, end of Column]`

`sort -n`

`sort -u`



Other interesting commands

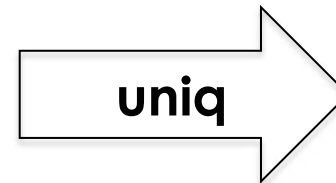
uniq

- Uniq combines identical neighboring lines into one

Example:

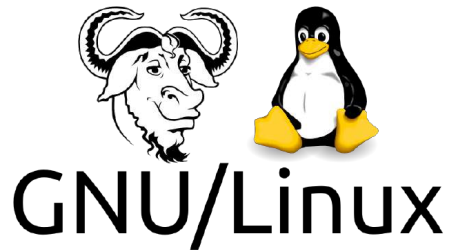
original

1
1
2
3
3
3
4



After uniq

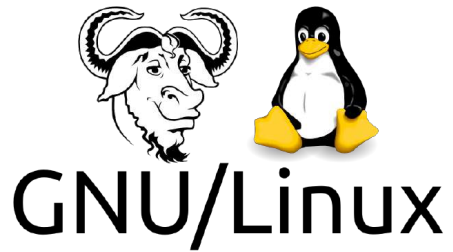
1
2
3
4



Other interesting commands

sed

- Stream **ED**itor
- Actual (simple and compact) programming language
- Parses and transforms text
- Line-oriented
- Input file is not changed
- Most famous application: text substitution

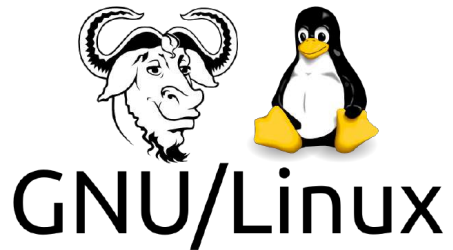


Other interesting commands

sed

substitution

`sed 's/original/substitution/' [file]`



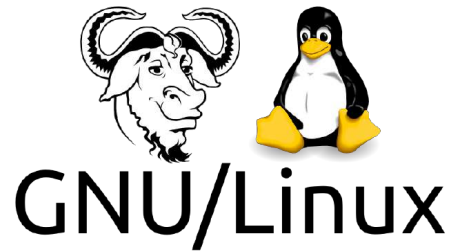
loops

'for' loop

for variable **in** list
do

commands **to** be executed for each item in the list

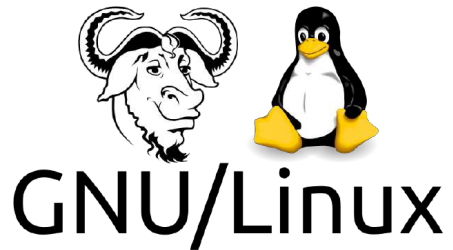
done



loops

'for' loop

```
for i in seq 5  
do  
    echo Wellcome $i times  
done
```

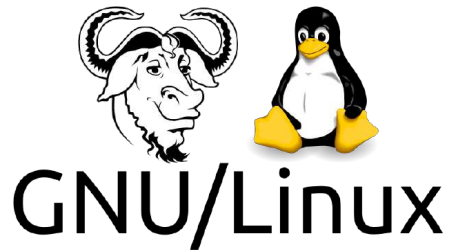



loops

'for' loop

```
for i in seq 5  
do  
    echo Wellcome $i times  
done
```

```
for i in $(seq 5); do echo Wellcome $i times; done
```

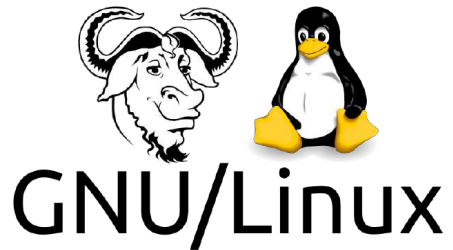


loops

‘while’ loop

```
while [ condition ]  
do
```

```
# commands to be executed while the condition is true  
done
```

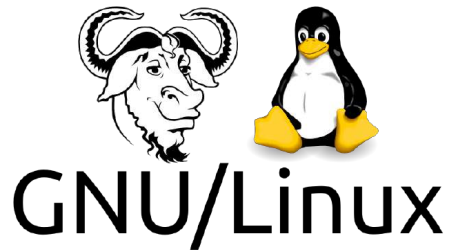


loops

‘while’ loop

```
count=1  
while [ $count -le 5 ]  
do  
    echo $count  
    count=$((count + 1))  
done
```

```
count=1; while [ $count -le 5 ]; do echo $count $(( count++ )); done
```



Things to remember

- Document all steps (will save you time on the long run)
- Save all your commands in one, clearly structured folder
- Document all errors and their solutions
- Search online for error solutions; likely someone else experienced the same problem before
- 99.99% of the errors are user errors
- No error does not mean the output is useful
- Always check your output files (file size and content)
- Find the right balance between understanding the software and accepting you can not know everything



With the support of the
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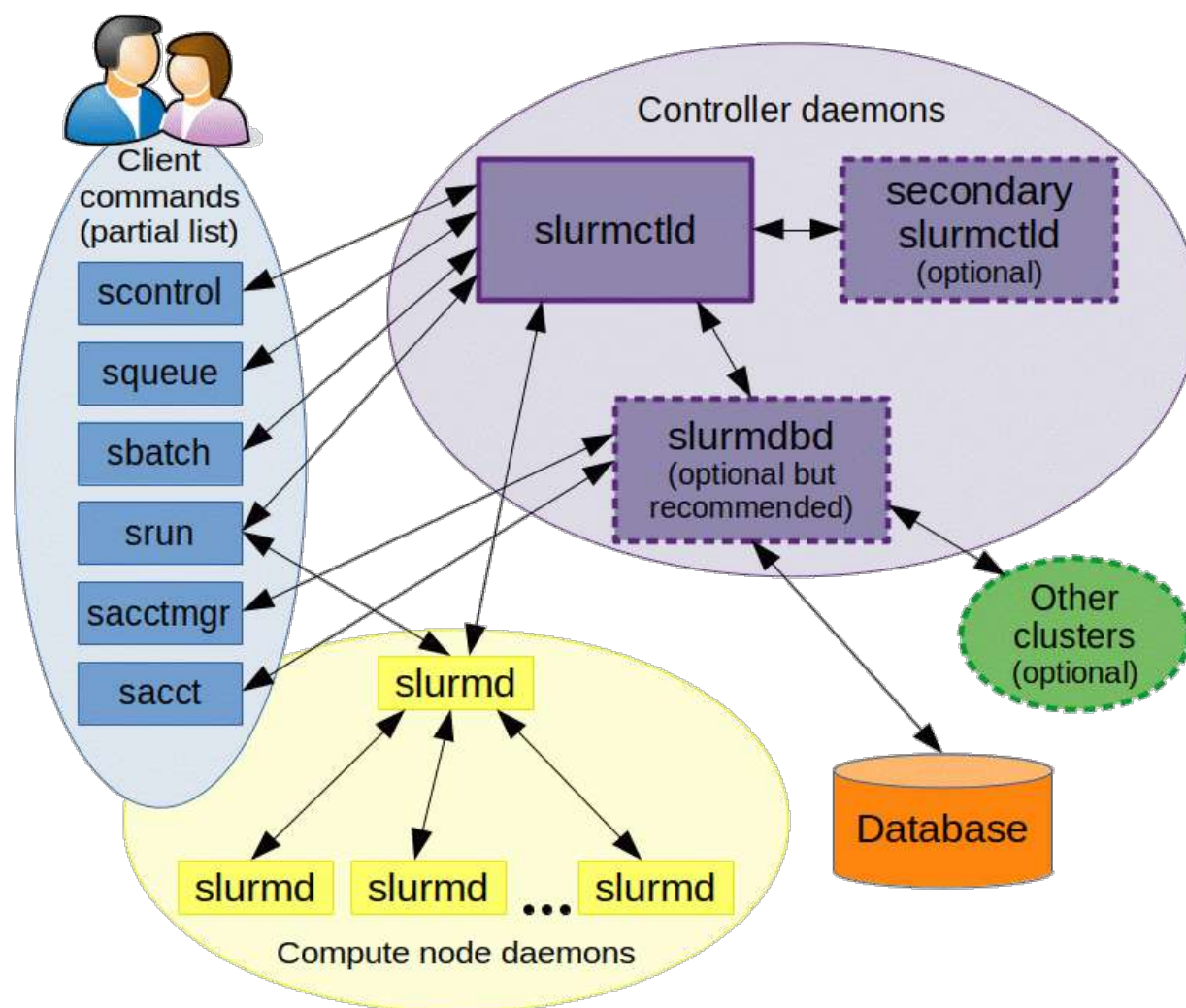


Simple Linux Utility for Resource Management



slurm

workload manager





sinfo

reports the state of partitions and nodes managed by **Slurm**. It has a wide variety of filtering, sorting, and formatting options

```
PARTITION AVAIL  TIMELIMIT  NODES  STATE NODELIST
main*      up        infinite    1    alloc caius
main*      up        infinite    5    idle
augustus,caesar,claudius,nero,tiberius
head       up        infinite    1    idle claudius
test       up        infinite    3    idle test[1-3]
tbg        up        infinite    3    idle test[1-3]
gpu        up        infinite    1    idle jupiter
```



srun

is used to submit a job for execution in real time.

```
$ srun -N 2 -n 48 -t 30 -A xz0123 ./my_small_test_job
```




sbatch

is used to submit a job script for later execution. The script will typically contain one or more **srun** commands to launch parallel tasks.

```
$ sbatch example.sh  
Submitted batch job 119405
```



example.sh

```
#!/bin/bash
#
#SBATCH -p generic          # Partición (cola)
#SBATCH -N 1                # Numero de nodos
#SBATCH -n 1                # Numero de cores (CPUs)
#SBATCH -mem 100            # Bloque de memoria para todos los nodos
#SBATCH -t 0-02:00          # Duración (D-HH:MM)
#SBATCH -o slurm.%N, %j.out  #STDOUT
#SBATCH -e slurm.%N, %j.err  #STDERR
#SBATCH --mail-type=END,FAIL # Notificación cuando el trabajo termina o
falla
#
#SBATCH -- mail-user=micorreo@ehu.eus # Enviar correo a la dirección
```



scancel

is used to cancel a pending or running job or job step. It can also be used to send an arbitrary signal to all processes associated with a running job or job step.

```
$ scancel 119405
```