Imperial College London



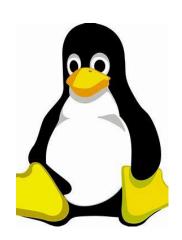
Linux and Command Line

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What is Linux?

- Just like Windows, iOS, and Mac OS, Linux is an operating system. It is free and open-source.
- An operating system is software that manages all of the hardware resources associated with your desktop or laptop.
- Many enterprise-level projects (c/c++/python/java) are deployed to Linux system. One of the most popular platforms, Android, is powered by the Linux operating system.

- Linux has a number of different versions to suit any type of user.
- Distributions (distros): specific group of software and conventions
- Major distributions: Arch, Debian (Linux Mint, Ubuntu, Kali, etc.), Red Hat (CentOS, Fedora, etc.), Slackware





Why Linux?

- For us ... HPC!
- Much of scientific experimentation such as modeling, simulations or data management and mining is made possible by computing -> high performance computing (HPC) cluster.
- HPC environment: Linux operating system & Bash shell
- Available teaching courses https://www.imperial.ac.uk/admin-services/ict/self-service/research-support/rcs/support/getting-started/
- Bioinformatics is highly dependent on Linux-based computers and software.
- 1. With the advent of new technologies (e.g. sequencing), analysis tend to rely more and more on powerful computing clusters.
- 2. Linux systems come with pre-installed versions of most popular programing languages and thus tend to become the natural platform for newly developed software(some programs are exclusively developed for Linux-based systems).

New to Linux

All our systems run Linux, and you'll need some familiarity with working in a terminal. If this is new to you, please read our introductions to the command-line and shell scripting. We also offer regular training courses.

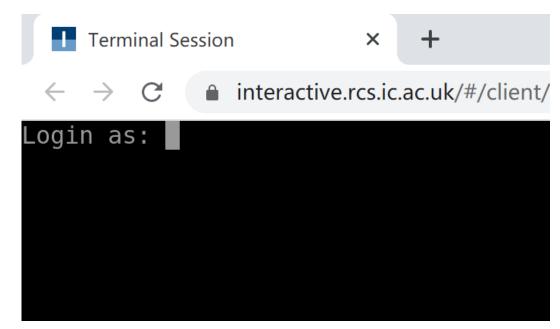


Running your first job

Our resources are batch processing systems. Rather than being run directly from the command line, jobs get submitted to a queue where they are held until compute resources become free. A job is defined by a shell script that contains all of the commands to be run.

Environment

- Terminal
- Mac easier Terminal
- Windows harder "run Linux on Windows" possible: Ubuntu, CentOS, Fedora...
- Login via the browser http://login.rcs.ic.ac.uk
 Without VPN? Try this: https://copy.sh/v86/?profile=linux26



Basic Bash commands

• ls:

Description: lists contents of current directory.

Common options:

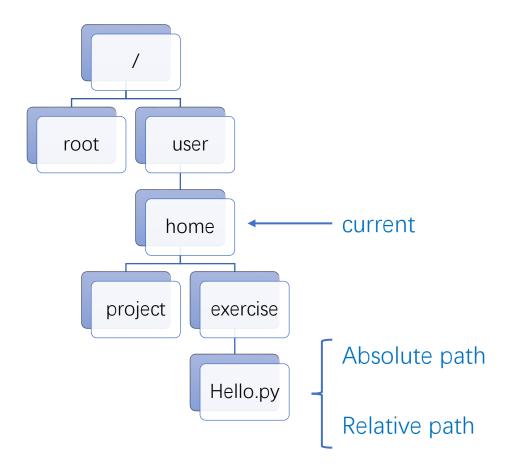
- -a lists all files of the current directory including the hidden files.
- -l lists contents of the current directory with extra details.

Examples: list all the contents in a specific directory.

- man: Displays the manual page of a given command.
- help: Displays information about built-in commands.

• pwd:

<u>Description:</u> prints working directory (the full path), or your location on the filesystem) <u>Absolute path (full path) and relative path</u> Absolute path and relative path



• cd:

<u>Description:</u> change directory <u>Examples:</u>

change directory to

- 1. child directory
- 2. parent directory
- a specific directory using absolute path
- 4. a specific directory using relative path
- 5. your home directory
- 6. the previous directory

Folders

• mkdir:

<u>Description:</u> makes a directory Examples:

- 1. Specify the path to where you want to create the new directory.
- 2. Create multiple directories.
- 3. Create nested directories.

• rmdir:

<u>Description:</u> removes a directory, which should be empty.

Files

cp:

<u>Syntax:</u> cp source_file destination_file <u>Description:</u> copy files and folders from one place to another.

Examples:

- 1. Copy files to a different path.
- 2. Copy multiple files.

• rm:

Description: remove files.

Options:

- -r Remove directories and their contents recursively.
- -d Remove empty directories. Equivalent to using rmdir.

Examples: remove multiple files with similar names.

• mv:

Examples:

- 1. move a file to different directory
- 2. move multiple files and the use of *
- 3. rename a file
- 4. move a directory
- 5. rename a directory

"

cp command = copy and paste

mv command = cut and paste

• cat:

<u>Description:</u> concatenate files and displays the output to the shell.

Common options:

- -b number all the non-empty lines
- -n number all output lines

Examples: Create a new file and check. (> redirects the output from a command.)

Editors: vim, Sublime, Eclipse, etc.

• wget:

<u>Description:</u> a network downloader which is used to download files from the server. It supports HTTP, HTTPS, and FTP protocols, as well as retrieval through HTTP proxies. Common options:

- -c resume a partially downloaded file
- -o save the downloaded file under a different name
- -P save the file to a specific location

- |: redirect the output of the first command as the input to next command.
- more: shows the contents of one or more files, one page at a time.
- less: displays the content of the file or the output of a command.

head:

<u>Description:</u> print the top N numbers of data of the given input. Common options: -n num: prints the first 'num' lines (default: 10).

• tail:

<u>Description:</u> print the last N numbers of data of the given input. Common options: -n num: prints the last 'num' lines (default: 10).

• wc:

<u>Description:</u> count the number of lines, words, characters, and bytes of each given file or standard input and print the result.

Common options:

- -l --lines Print the number of lines.
- -w --words Print the number of words.
- -m --chars Print the number of characters.
- -c --bytes Print the number of bytes.
- -L --max-line-length Print the length of the longest line.