



# H3ABioNet

Pan African Bioinformatics Network for H3Africa

## Introduction to Bioinformatics online course: IBT

## Linux

## Introduction to Linux and Unix and the command line



# Learning Objectives

- ① Understand the Unix file structure
- ② Understand full vs relative paths: when and how to use them
- ③ Learn how to create directories and navigate through the file structure
- ④ Learn some useful shortcuts

# Learning Outcomes

- ① Be able to create a file structure
- ② Be able to navigate through the file structure
- ③ Be able to create text files and view their content
- ④ Be able to use simple shortcuts

# Part 1

## Introduction to Linux and UNIX

# What is Linux?

- UNIX is an **Operating System** (OS) initially developed in the 1960.
- There are many different versions of UNIX, that share common similarities.
- The most popular varieties of UNIX are Solaris, **Linux** and MacOS.
- UNIX systems have a graphical user interface (GUI) making them easier to use.

# Linux vs Unix

- Linux is a “clone” of the original Unix but doesn’t contain its code
- Linux is free and open source, the original Unix is not (but some of its derivatives are)
- All command lines work the same on both

# Why Linux?

- Linux is free and the most popular distributions are Ubuntu, Fedora/Red Hat, Mandriva, etc.
- Low cost and very stable system
- Most secure OS
- Best multi-user and multi tasking OS
- The world's fastest super computers run Linux
- Fast developing OS (many developers)



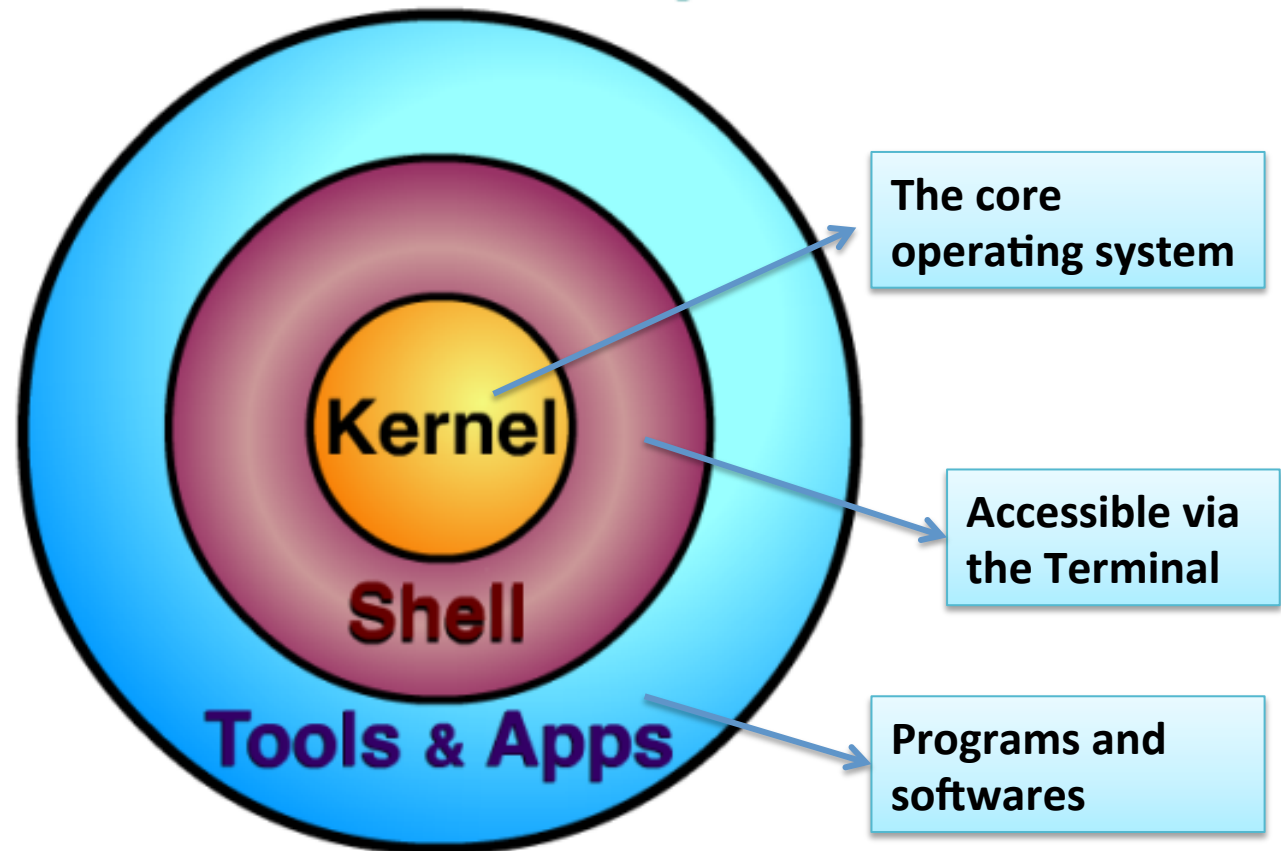
**Very popular as servers OS**

# Linux distributions

- Different Linux distributions are available  
<http://distrowatch.com/>
- Ubuntu distribution is easy and convenient to use for beginners
- A simple guide to install Ubuntu in your machine:  
<http://www.ubuntu.com/download/desktop/install-ubuntu-desktop>



# Parts of the UNIX System



Adapted from: [www.usna.edu](http://www.usna.edu)

- A **terminal** refers to a wrapper program which runs a shell
- There are many different Unix shells, the most popular shell for interactive use include **Bash**: the default on most Linux installations

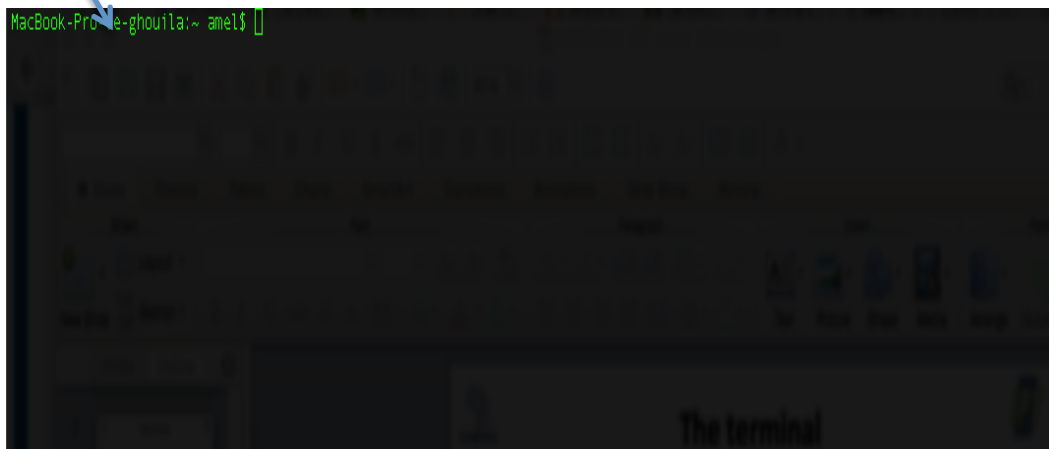
# The Terminal

**Shell prompt**

**User name**

**Machine name**

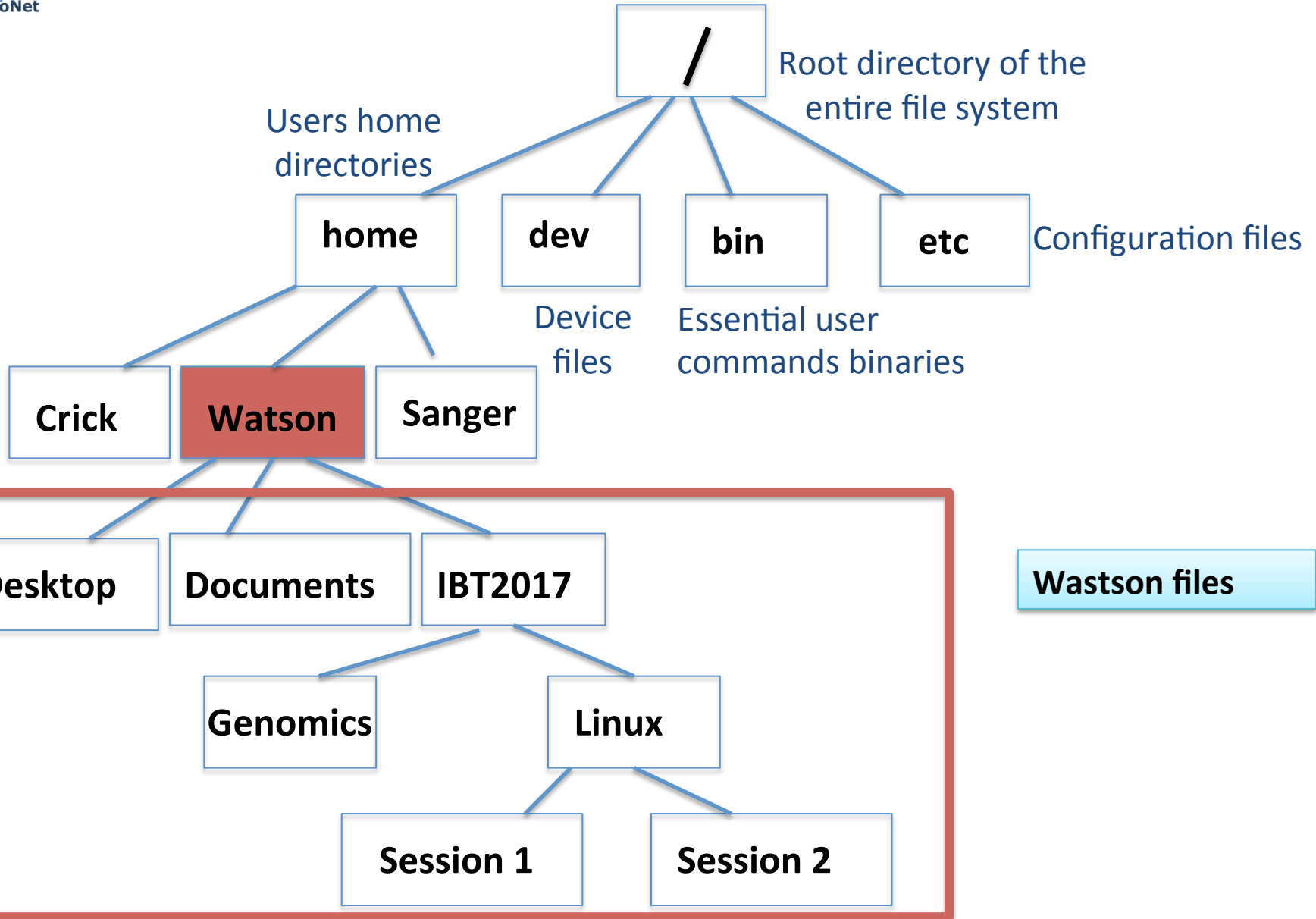
Even though it is a command line interface, the mouse is still handy (scroll, copy, paste, etc.)



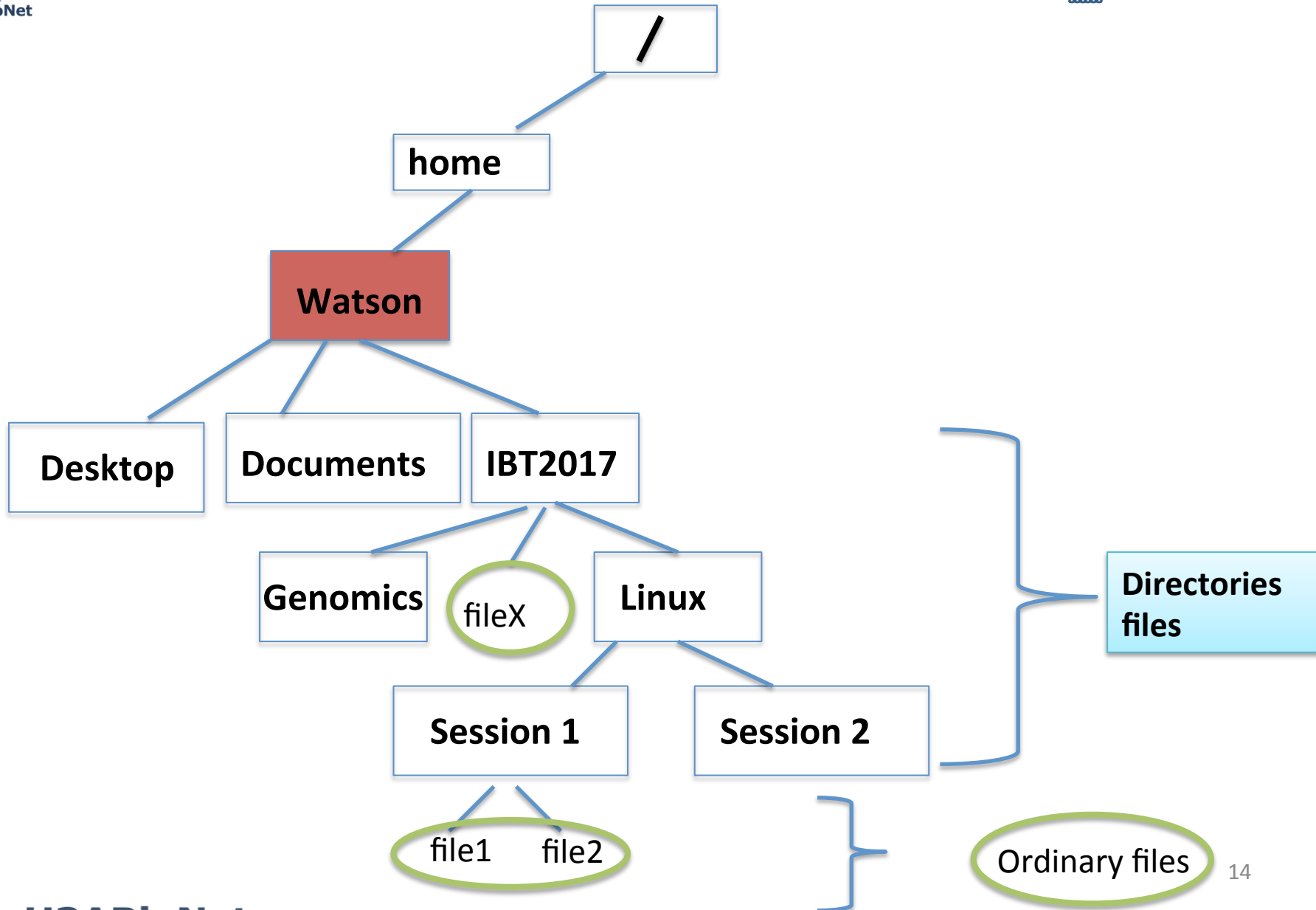
## Part 2

# File-system under UNIX

# Linux files structure



Watson files



# Home directory and working directory

- When you first log in on a UNIX system, the working directory is your **home** directory.
- While working you will be associated to one directory called the **working directory** or the **current directory**
- An abbreviation of the working directory is displayed as part of the prompt on your terminal
- The command **pwd** gives the absolute path of the working directory

# What is a path or a pathname?

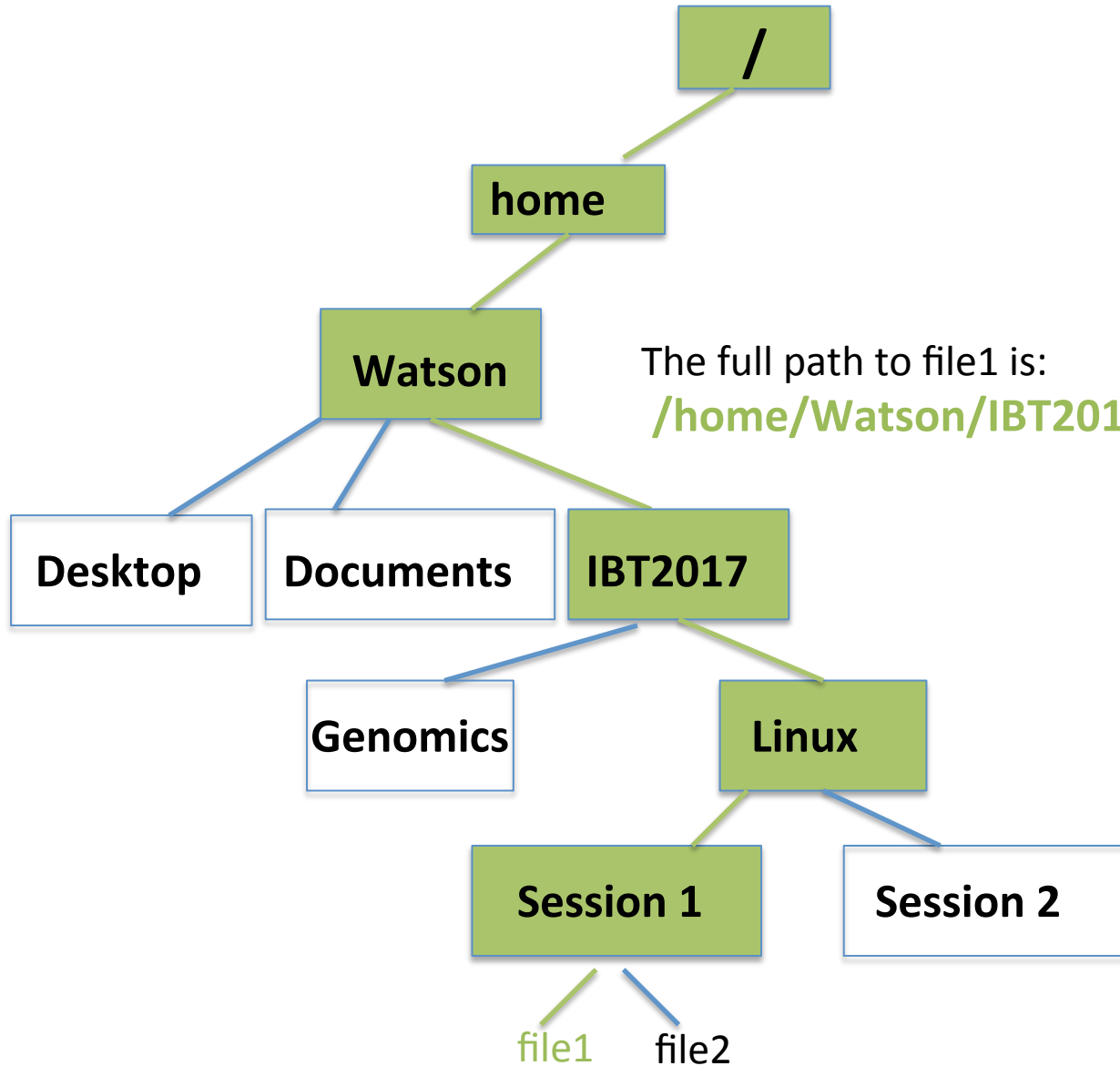
- A path locates a given file in the system hierarchy
- An **absolute path** in the file system hierarchy for a given file or folder describes the parents all the way up to the root
- A **relative path** describes the path to the file starting from the **current working directory**



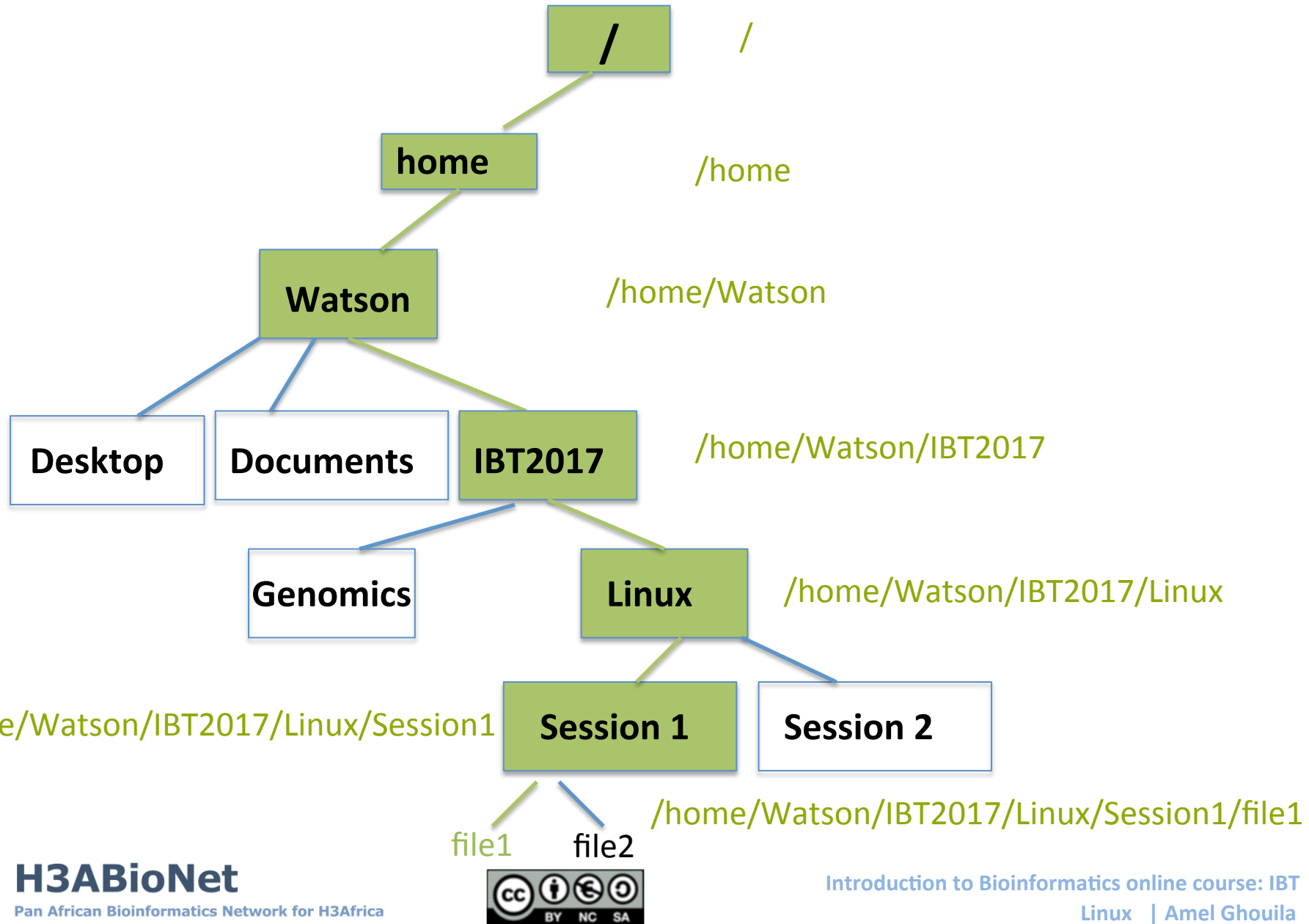
## ~ (your home directory)

- ~ refers to the home directory in a given file system
- The tilde ~ character can be used to specify paths starting at your home directory

# Absoulte path?



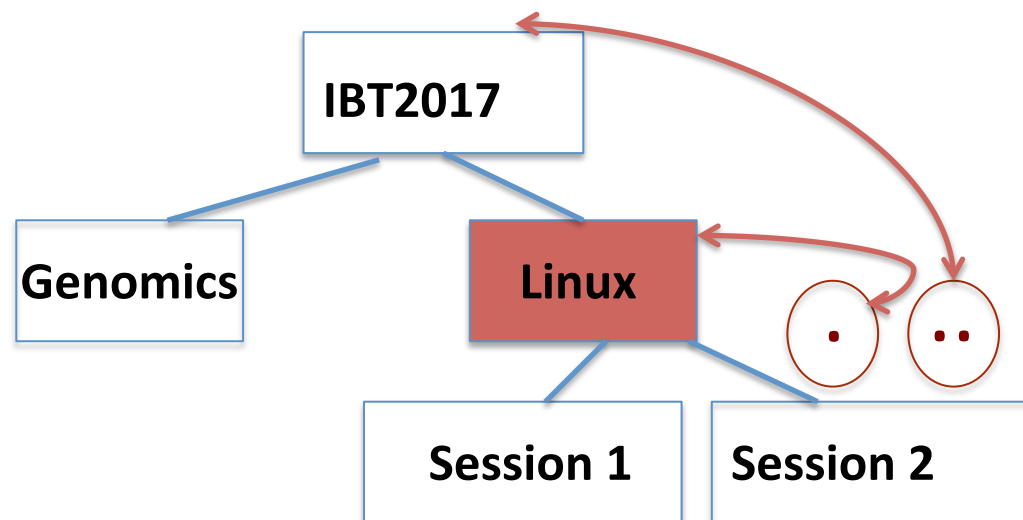
# Absoulte path?



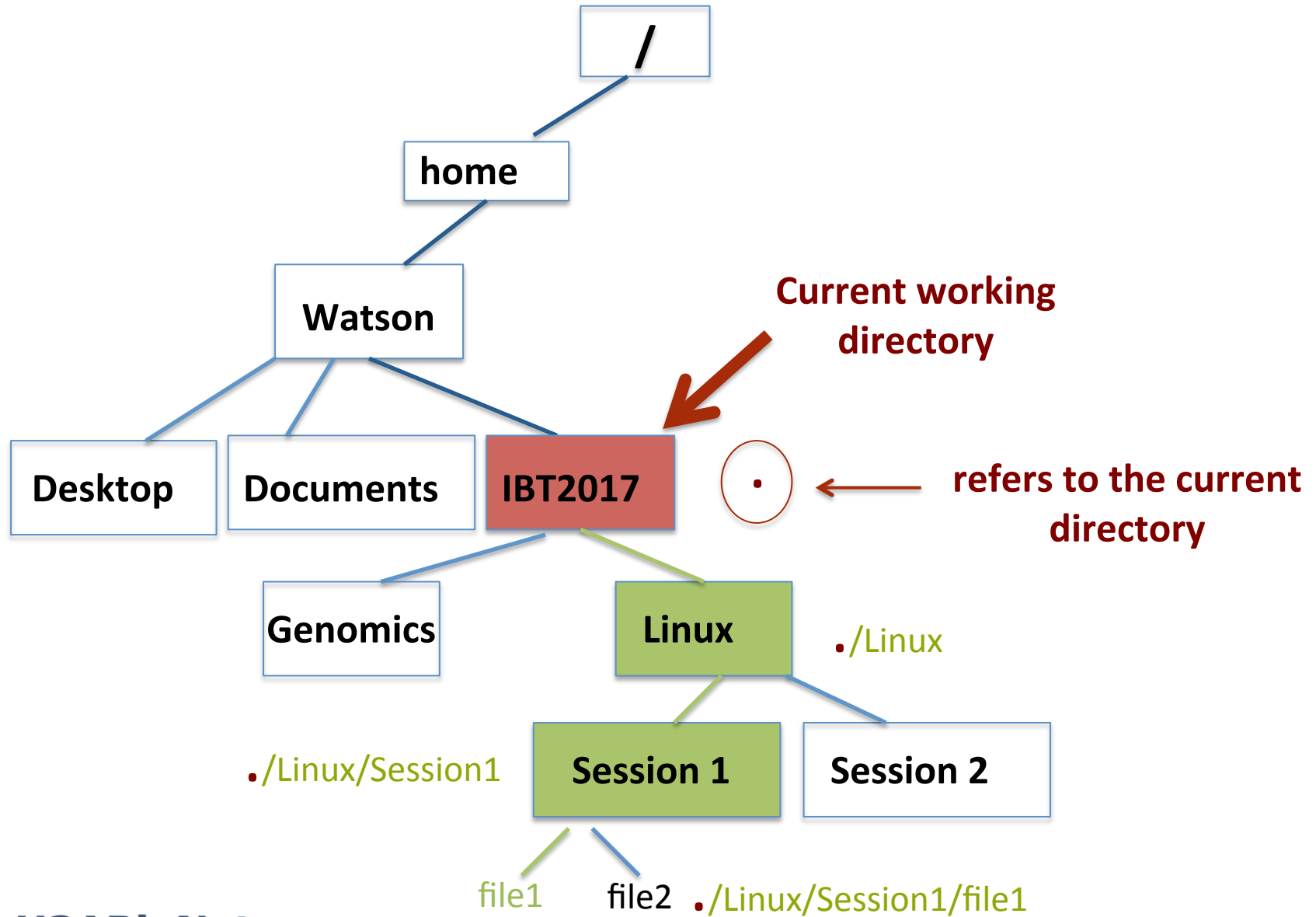
# Refer to the parent and current directories

Every directory has two special sub-directories:

- ◆ . (dot): the current directory
- ◆ .. (dot-dot): the parent directory



# Relative path?



# First test of the terminal

- Open the Terminal on your system
- The shell prompt will appear

Ok, let's try some typing !

## Part 3

# Creating directories and navigating through the file structure

# Commands for manipulating directories

mkdir	Make directory: creates a new directory
rmdir	Removes a directory
pwd	Displays the absolute path of the current working directory
cd	Change directory: allows moving from one directory to another
ls	Lists a directory content



# pwd command

- pwd: **print working directory**
- Displays the **absolute path** of your current location in the file system
- Try **pwd** on your terminal
- You should see: /home/YourUsername

# ls command

- ls lists the content of the current directory by default
- Command structure **ls** [OPTION] [dirname]
- Some useful options:
  - -l: shows sizes, modified date and time, file or folder name and owner of file and permissions
  - -a: List all files including hidden file starting with '.'
  - -lh: shows sizes in easier readable format
  - -R: recursively lists sub-directories
  - -ls: sorting by file sizes

# Create a directory

- mkdir: **makes a directory**
- Command structure: **mkdir** **dirname** [path]
- mkdir **dirname**: would create a directory with the specified **dirname**
- The new created directory will be created in your current working directory
- If you want to create it elsewhere, you have to specify the path: mkdir **dirname** **path**

# Commands basic structure

**command** **[–options]** **[arguments]**

Example:

**ls –lh** /home/Watson/IBT2016

**pwd**

**mkdir** Test1

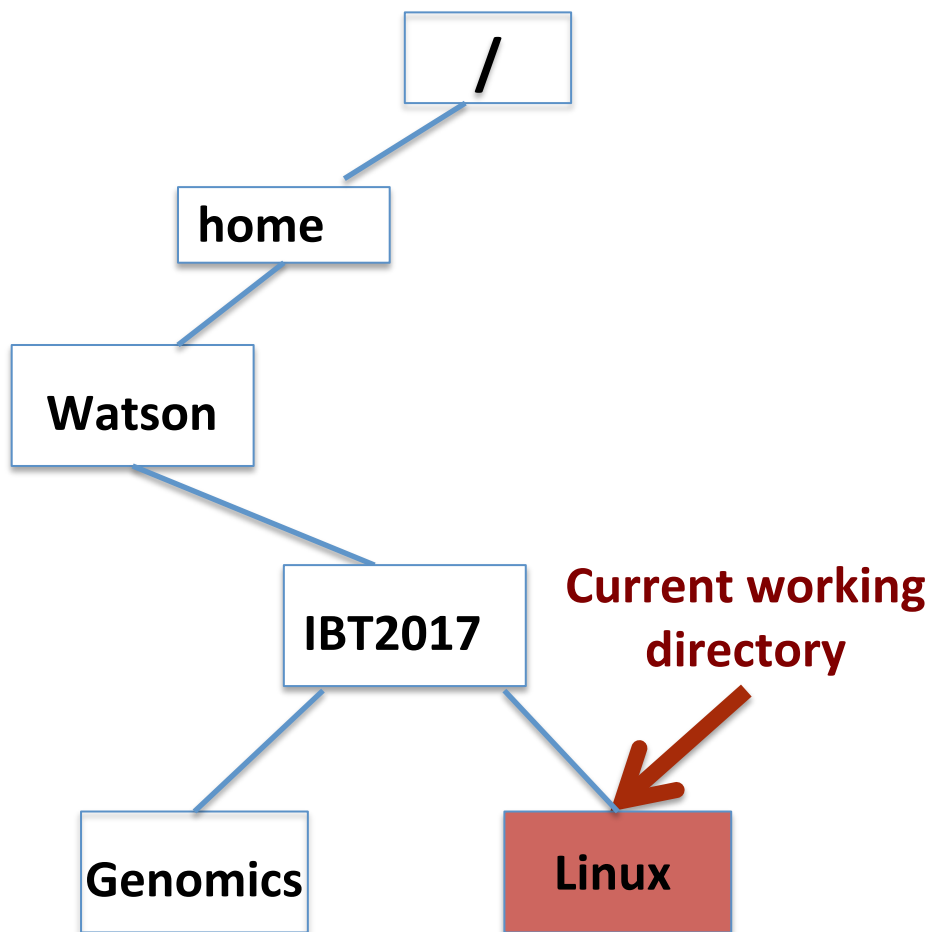
# What you should know about file names in Linux

- No real distinction between the names of ordinary files and the names of directory files.
- No two files in the same directory can have the same name.
- Files in different directories can have the same name.
- Linux is case-sensitive: **Sanger**, **sanger** and **SANGER** are different and would represent three distinct files.
- In most cases, file extensions are optional (.txt, .exe, etc.)

# Move in the files system

- **cd**: change the working directory
- Command structure: **cd** <path>
- The path name of the directory you want to move to should be specified
- You can specify either the absolute path or the relative path

# Move in the files system: example



- Move to Watson directory

1. `cd /home/Watson`
2. `cd ../..`
3. `cd .. + cd ..`

- Move to Genomics directory

1. `cd /home/Watson/IBT2017/Genomics`
2. `cd ../Genomics`
3. `cd .. + cd Genomics`

# Remove a directory

- `rmdir`: **removes a directory**
- Command structure: `rmdir` **dirname** **[path]**
- It would remove the **dirname** directory
- The directory should be in your current working directory
- If you want to remove it from elsewhere, you have to specify the path: `rmdir` **dirname** **path to the directory**
- **`rmdir`** works if there is no contents in the directory



# Remove a directory

- **rm** works if there is no contents in the directory
- If the directory contains files or sub-directories, an error message will appear:  
**“Directory not empty”**
- There is an option to remove **-r**, which stands for **recursive**, that will **recursively** remove a directory and its contents

# How to get help for a command from the terminal?

- **man** `commandname` displays

## Some useful shortcuts and Links

# Useful Links

- **cd**: cd followed by nothing will change the working directory to your home directory
- **cd** ~user\_name: moves to the specified user home directory

# Linux, some useful CLI key combinations

- **Ctrl+A**: move the cursor to the beginning of the command line
- **Ctrl+C**: end a running program and return to the prompt
- **Ctrl+D**: logout from the current shell session: equivalent to exit
- **Tab**: autocomplete a file name
- **Tab Tab**: displays command completion possibilities
- **Ctrl+L**: clear the terminal

# Thanks

- Shaun Aron & Sumir Panji