



# Introduction to Linux and the Command Line with SimpleVM

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# Hands-On Session

- **Part 1: SimpleVM Introduction**
- Part 2: Linux Command Line



- SimpleVM is a multi-cloud application that eases the access to computational resources of the federated de.NBI Cloud.
- It takes care of any necessary network or volume setup.
- Automatically installed software
- No background knowledge in cloud computing necessary
- Additional Project Modes:
  - Workshop Mode
  - SimpleVM Cluster
- New SimpleVM Portal since October 2023

# de.NBI Cloud - A Solution for (almost) Every Use Case

## Infrastructure-, Platform- and Workflows- as-a-Service



- ▶ High configurability, infrastructure virtualization
- ▶ API access, e.g. for use with Terraform or Ansible
- ▶ Any software of the cloud ecosystem



- ▶ High configurability, service and container orchestration
- ▶ API access, e.g. for use with kubectl
- ▶ Any containerized software / service, Helm charts



- ▶ Beginner-friendly, preconfigured Research Envs
- ▶ ELIXIR guarded interactive browser sessions
- ▶ “One-click” solution for setting up a workshop or cluster

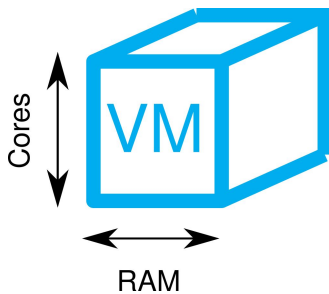


- ▶ Fully established bioinformatics tools and workflows, maintained by the community
- ▶ Point-and-click GUI for composition of bioinformatics workflows
- ▶ Interactive tours and comprehensive training library

# Cloud Components

- Main building block of the cloud is a **virtual machine**.
- A virtual machine **instance** is a running instance of the virtual machine image with resource parameters (Cores and RAM) assigned to it.

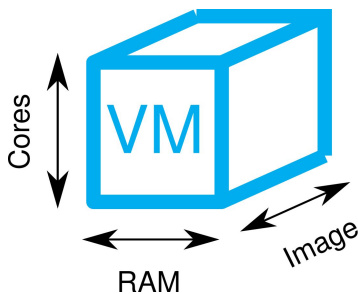
## 1. Create your blueprint



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- A virtual machine **image** is a virtual disk that has a bootable operating system installed on it.

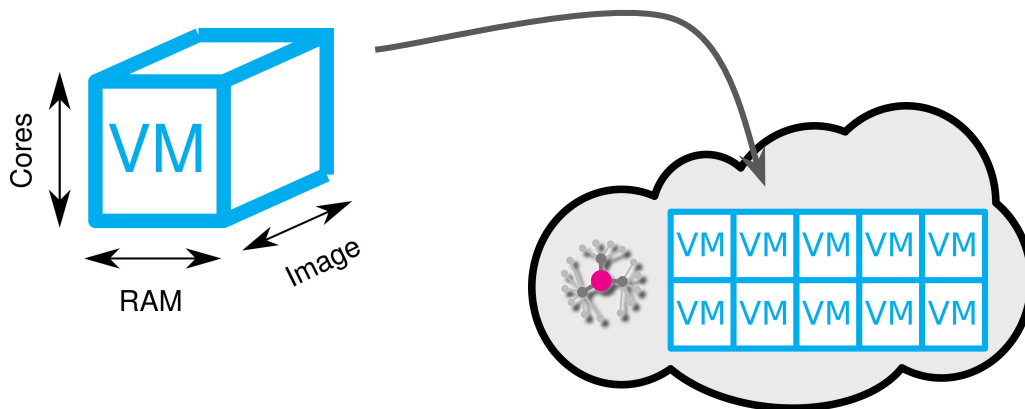
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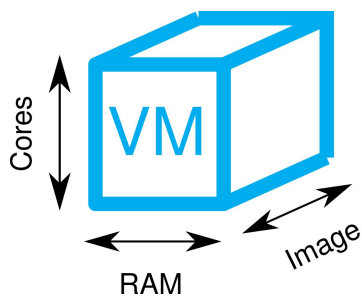
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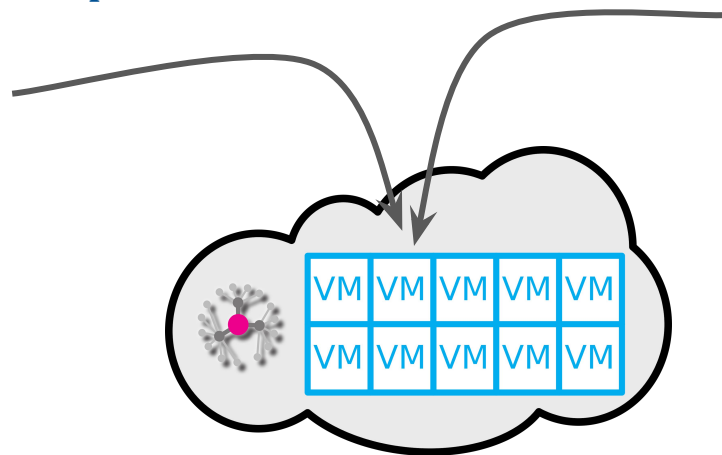
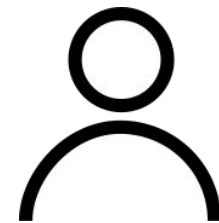
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## 1. Create your blueprint



## 2. Connect through ssh

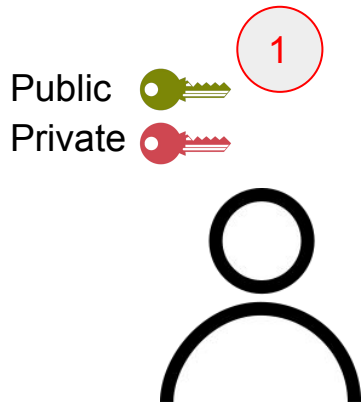




# Cloud Components - Secure Shell Protocol (SSH)

## First Time Usage

1. Generate a key pair locally or let the Portal generate one for you

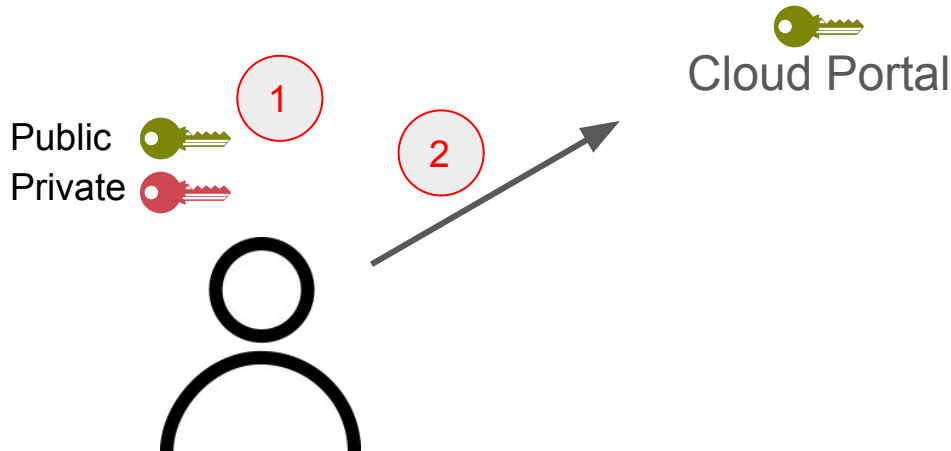


# Cloud Components - Secure Shell Protocol (SSH)

## First Time Usage

1. Generate a key pair locally or let the Portal generate one for you
2. Upload your public key to the de.NBI Cloud Portal/OpenStack

## Connect to your VM

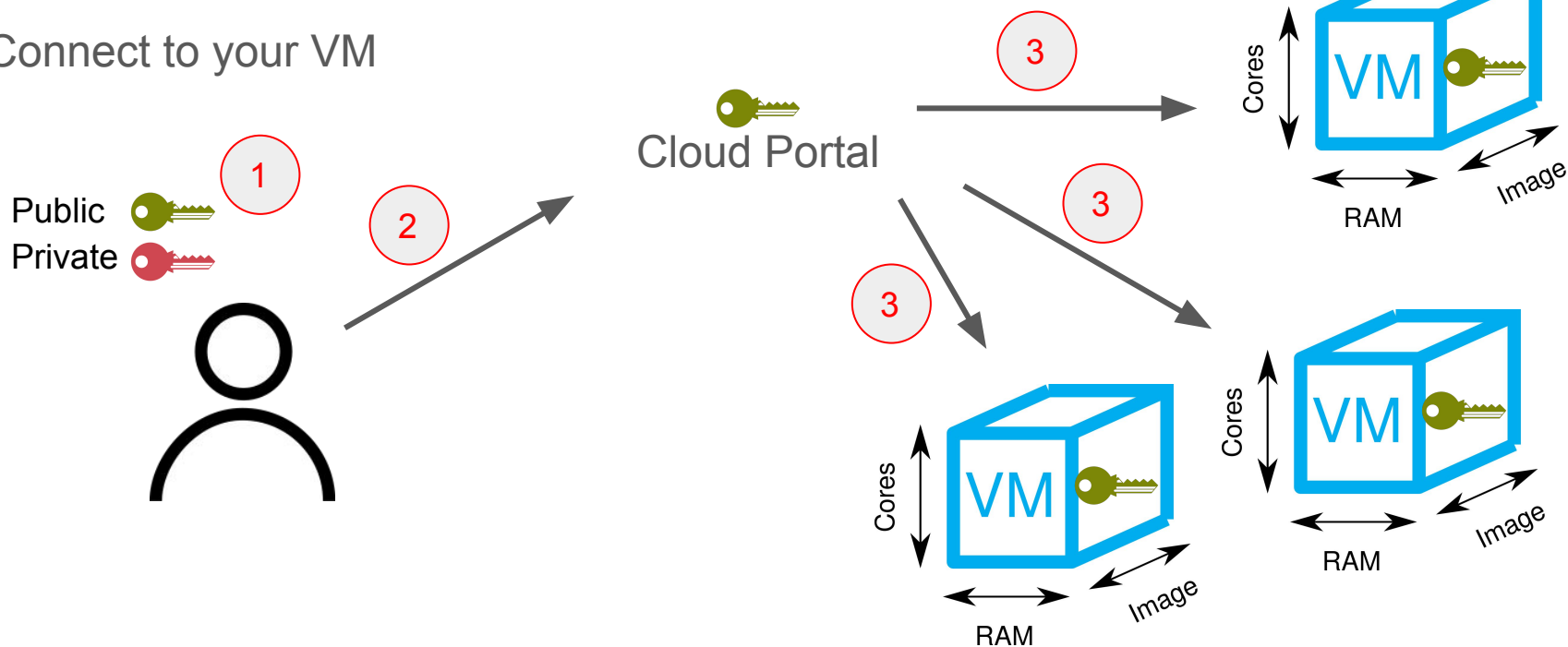


# Cloud Components - Secure Shell Protocol (SSH)

## First Time Usage

1. Generate a key pair locally or let the Portal generate one for you
2. Upload your public key to the de.NBI Cloud Portal/OpenStack
3. The Cloud Portal will set the public key on every new VM

## Connect to your VM



# Hands-On Session

- Go to the first part of the workshop:

<https://github.com/deNBI/simpleVMWorkshop/blob/main/part1.md>

- **Please start with part 1 of the SimpleVM workshop**
- ~ 10 Minutes

Internet:

WiFi Name:

LAN1-JBB

WiFi PWD:

jbb45\_45

# Hands-On Session

- Part 1: SimpleVM Introduction
- **Part 2: Linux Command Line**

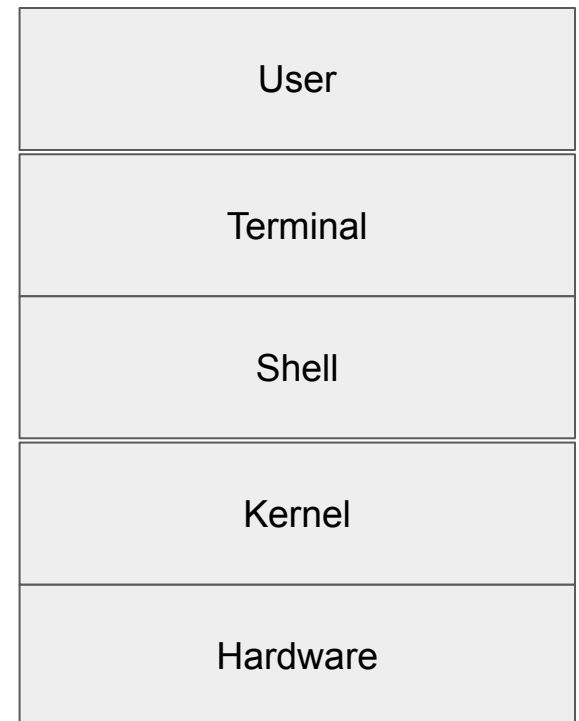
## Part 2: Linux Command Line Introduction

# Linux Command Line Introduction - Unix/Linux?

- Linux is a flavour of **Unix**
- **Unix**: set of programs to interact with the computer
- Main Properties:
  - **Multitasking environment**: Multiple processes can run in parallel
  - **Multiuser system**: Multiple users can use the same Linux system at the same time

# Linux Command Line Introduction - Unix/Linux?

- The heart of a Linux system is the **kernel**
- The kernel allocates system resources
- You, as a user, use a **shell** to interact with the kernel
- The **terminal** is a graphical interface that lets you interact with the shell





# Linux Command Line Introduction - Part 1

- Almost every command has a usage and manual **man** page

Get help and usage message: **--help / -h / -? / --usage**

```
ls --help
```

Read the manual: **man**

```
man ls
```

 one page down

 one page up

 quit



If **man** does not give any result, try:

**info**  
**help**



Don't know what command to use? Try:

**apropos**

# Linux Command Line Introduction - Part 1

- Data in Unix is organized in files
  - Three types of files:
    - ordinary files (e.g. a program or text),
    - directories,
    - special files (e.g. shortcuts)
- Directories organized in hierarchical, tree like structure
  - with “/” as its root.
    - /home/ubuntu/projects/unix-course
    - /home/ubuntu/Downloads/test.txt

# Linux Command Line Introduction - Part 1

- Commands for navigation: **pwd**, **ls**, **cd**
- Places to navigate (current: `/home/ubuntu/dir`, aim: `/home/ubuntu`):
  - Absolute path: **cd** `/home/ubuntu/`
  - Relative path: **cd** `..`
  - Home directory: **cd** `~`
- Commands to manipulate the filesystem tree:
  - **mkdir**, Syntax: `mkdir directory_to_create`
  - **rm -r**, Syntax: `rm -r directory_to_remove`
- View a file:
  - **less** `file_to_view`
- Other:
  - **file**: Returns you the type of the file
  - **df**: Allows you check the usage of your disk. (size of the free and used space)

# Linux Command Line Introduction - Part 1

- Permissions:
  - You can view the permissions of file with “**ls -l**”

```
xterm
ubuntu@sebastianmain-03787:~$ ls -l
total 36
drwxr-xr-x 2 root root 4096 Nov 21 13:11 Desktop
drwxr-xr-x 2 ubuntu ubuntu 4096 Nov 22 12:18 Documents
drwxr-xr-x 2 ubuntu ubuntu 4096 Nov 22 12:18 Downloads
drwxr-xr-x 2 ubuntu ubuntu 4096 Nov 22 12:18 Music
drwxr-xr-x 2 ubuntu ubuntu 4096 Nov 22 12:18 Pictures
drwxr-xr-x 2 ubuntu ubuntu 4096 Nov 22 12:18 Public
drwxr-xr-x 2 ubuntu ubuntu 4096 Nov 22 12:18 Templates
drwxr-xr-x 0 root root 0 Jan 1 1970 thinclient_drives
drwxr-xr-x 3 root root 4096 Apr 8 2019 Trimmomatic-0.39
drwxr-xr-x 2 ubuntu ubuntu 4096 Nov 22 12:18 Videos
ubuntu@sebastianmain-03787:~$
```

permissions	owner	user	group	size (kb)	last modified	file name
drwxr-xr-x	2	root	root	4096	Nov 21 13:11	Desktop
drwxr-xr-x	2	ubuntu	ubuntu	4096	Nov 22 12:18	Documents
drwxr-xr-x	2	ubuntu	ubuntu	4096	Nov 22 12:18	Downloads
drwxr-xr-x	2	ubuntu	ubuntu	4096	Nov 22 12:18	Music
drwxr-xr-x	2	ubuntu	ubuntu	4096	Nov 22 12:18	Pictures
drwxr-xr-x	2	ubuntu	ubuntu	4096	Nov 22 12:18	Public
drwxr-xr-x	2	ubuntu	ubuntu	4096	Nov 22 12:18	Templates
drwxr-xr-x	0	root	root	0	Jan 1 1970	thinclient_drives
drwxr-xr-x	3	root	root	4096	Apr 8 2019	Trimmomatic-0.39
drwxr-xr-x	2	ubuntu	ubuntu	4096	Nov 22 12:18	Videos

# Linux Command Line Introduction - Part 1

- Permissions:
  - You can view the permissions of file with “**ls -l**”
  - Modify permissions by using **chmod**
    - **-rw-r-- --** = (**user**, **group**, **others**)
    - r = readable, w = writable, x = executable
    - Example: **chmod g+w file** (make file writable for the group)
- Execute a program:
  - You can execute a local program by providing the path of the file:  
“/path/to/my\_program\_to\_execute”
  - You can cancel any command with “Ctrl+c”

# Linux Command Line Introduction - Part 1

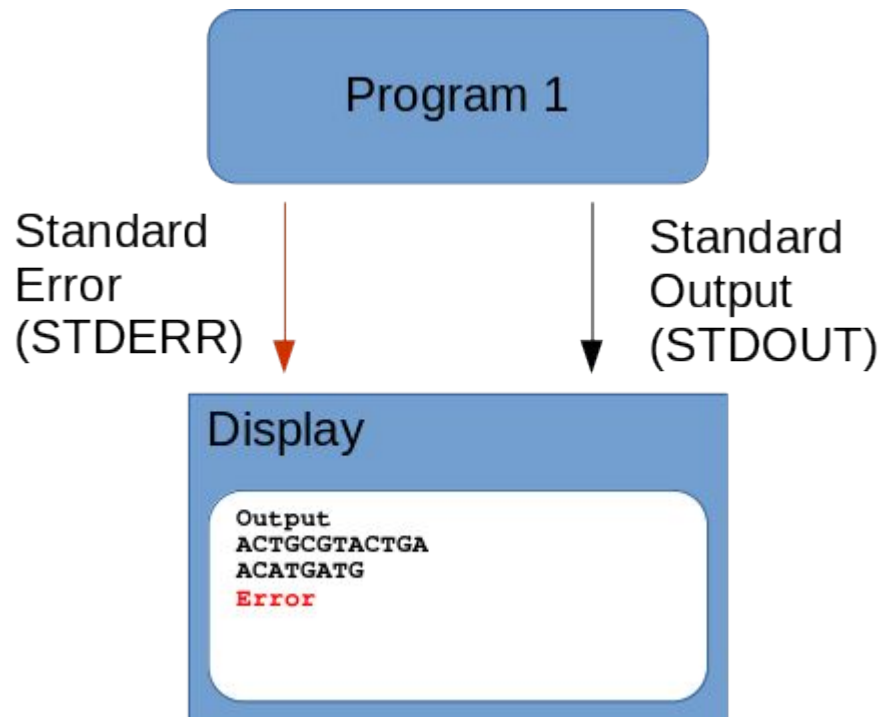
- Go to the second part of the SimpleVM workshop:

<https://github.com/deNBI/simpleVMWorkshop>

- Once you are done, continue with the actual Linux Course:
- Your task is to **identify the correct commands** by using the list of commands and execute them. Feel free to experiment. Take a look at the solution **if absolutely necessary**.
- Tutorial: <https://github.com/deNBI/unix-course>
- Please just do part 1 of the tutorial!
- ~20 Minutes

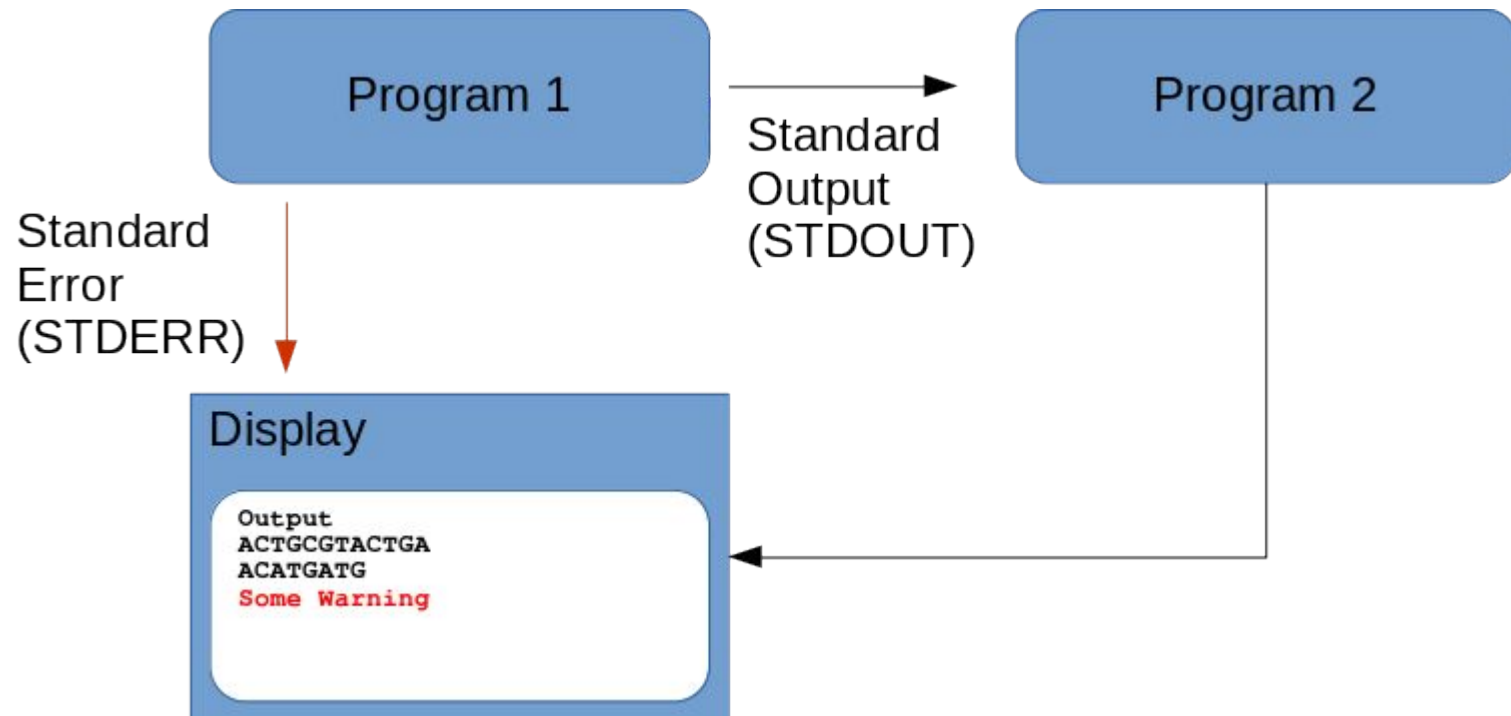
# Linux Command Line Introduction - Part 2

- Many unix commands support streams (files) as input and/or output



# Linux Command Line Introduction - Part 2

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- Streams can be **redirected**

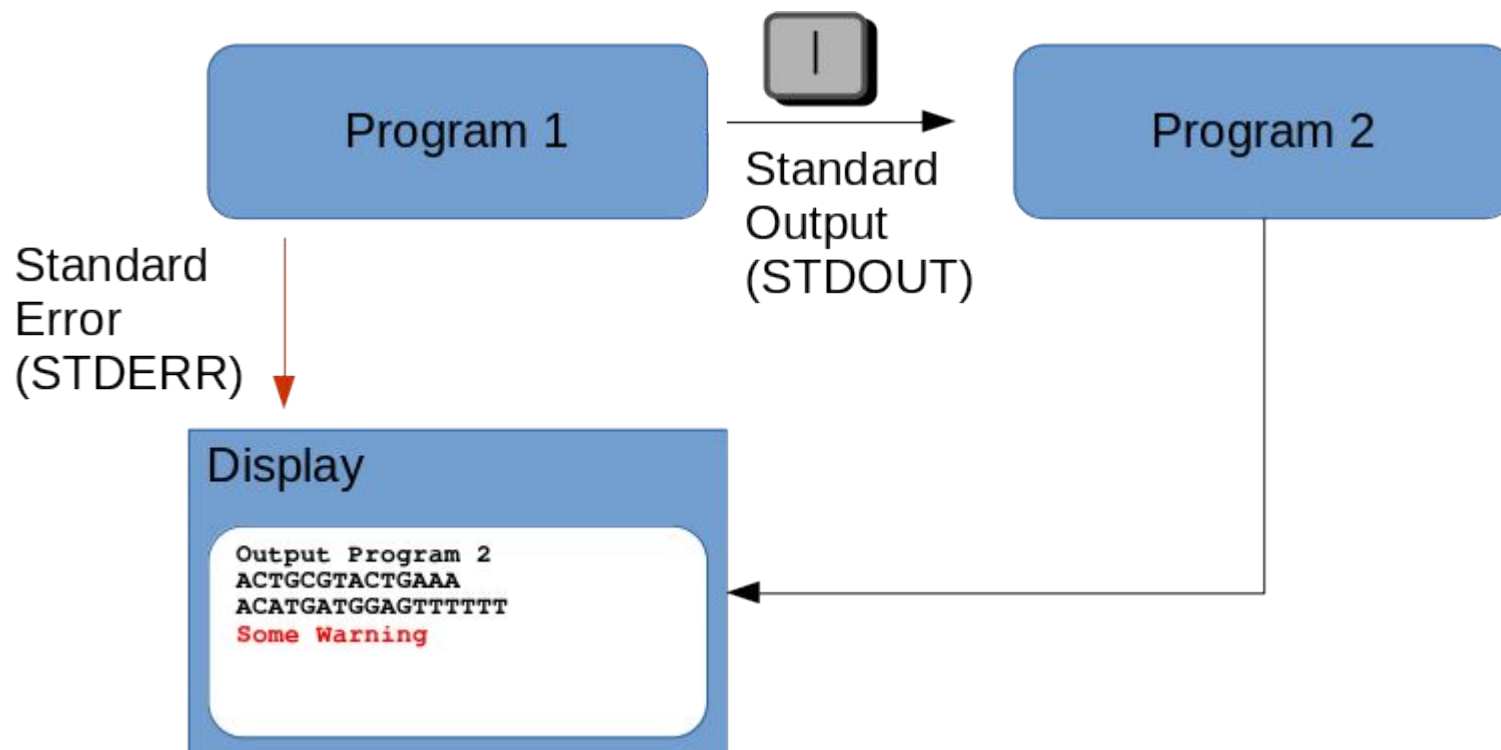


How?



# Linux Command Line Introduction - Part 2

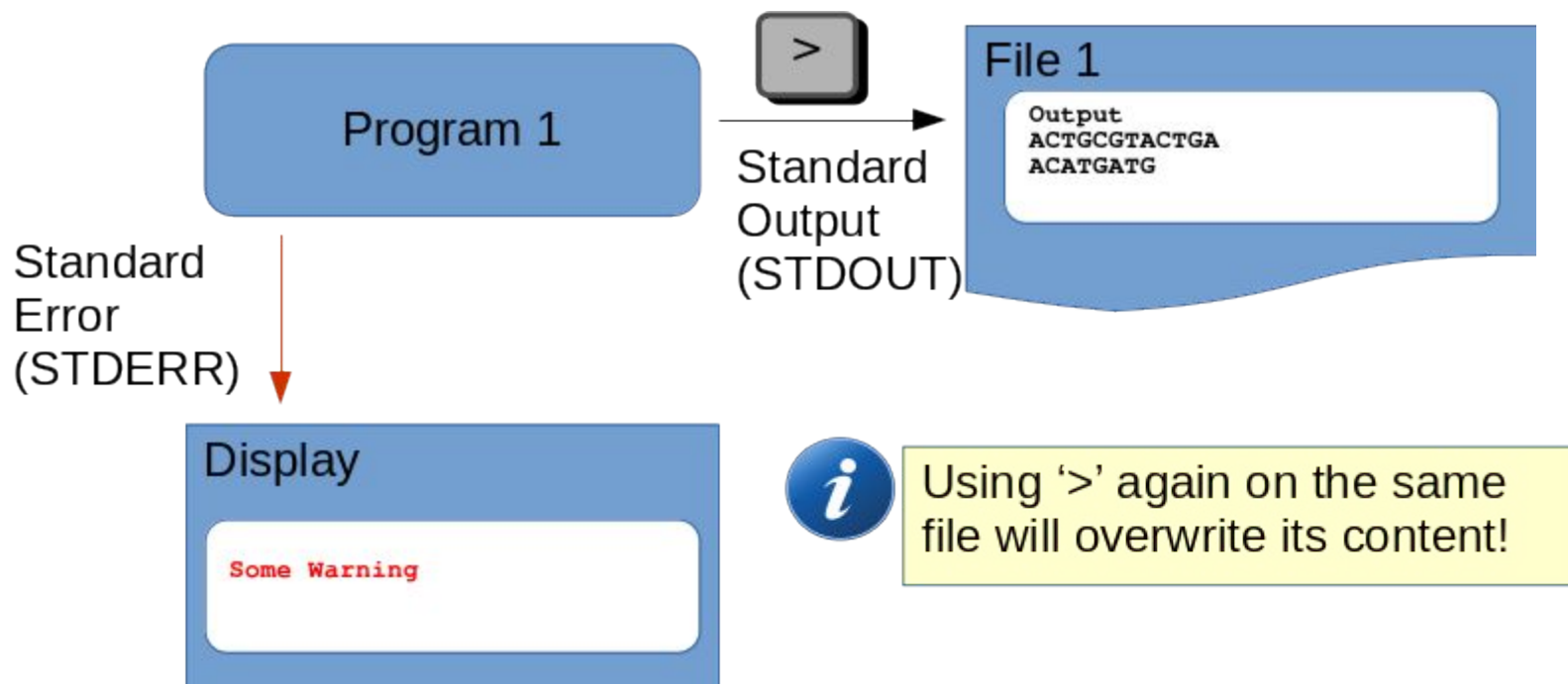
- Many unix commands support streams (files) as input and/or output
- Streams can be **redirected**



Example: `ls -la | less`

# Linux Command Line Introduction - Part 2

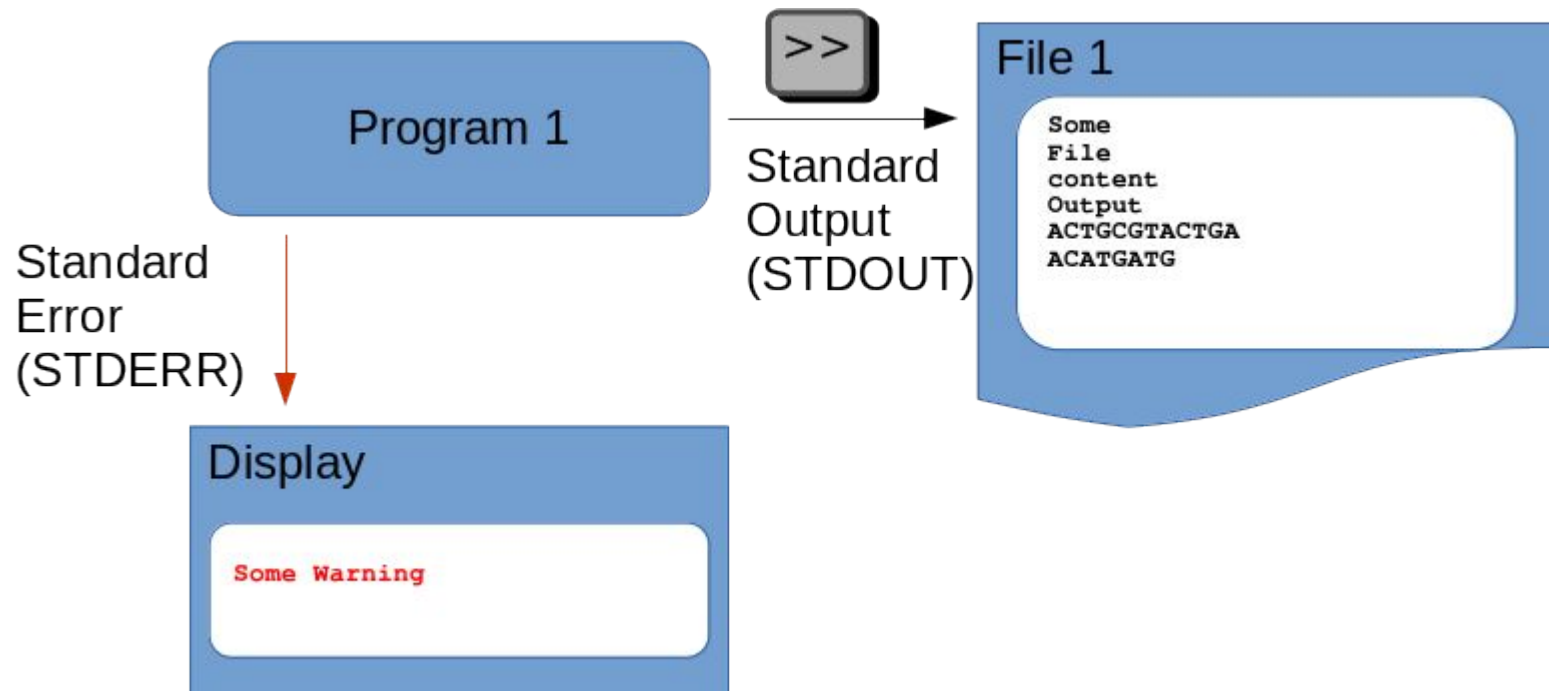
- Many unix commands support streams (files) as input and/or output
- Streams can be **redirected into files**



Example: `ls -la ~ > myHomeDir.txt`

# Linux Command Line Introduction - Part 2

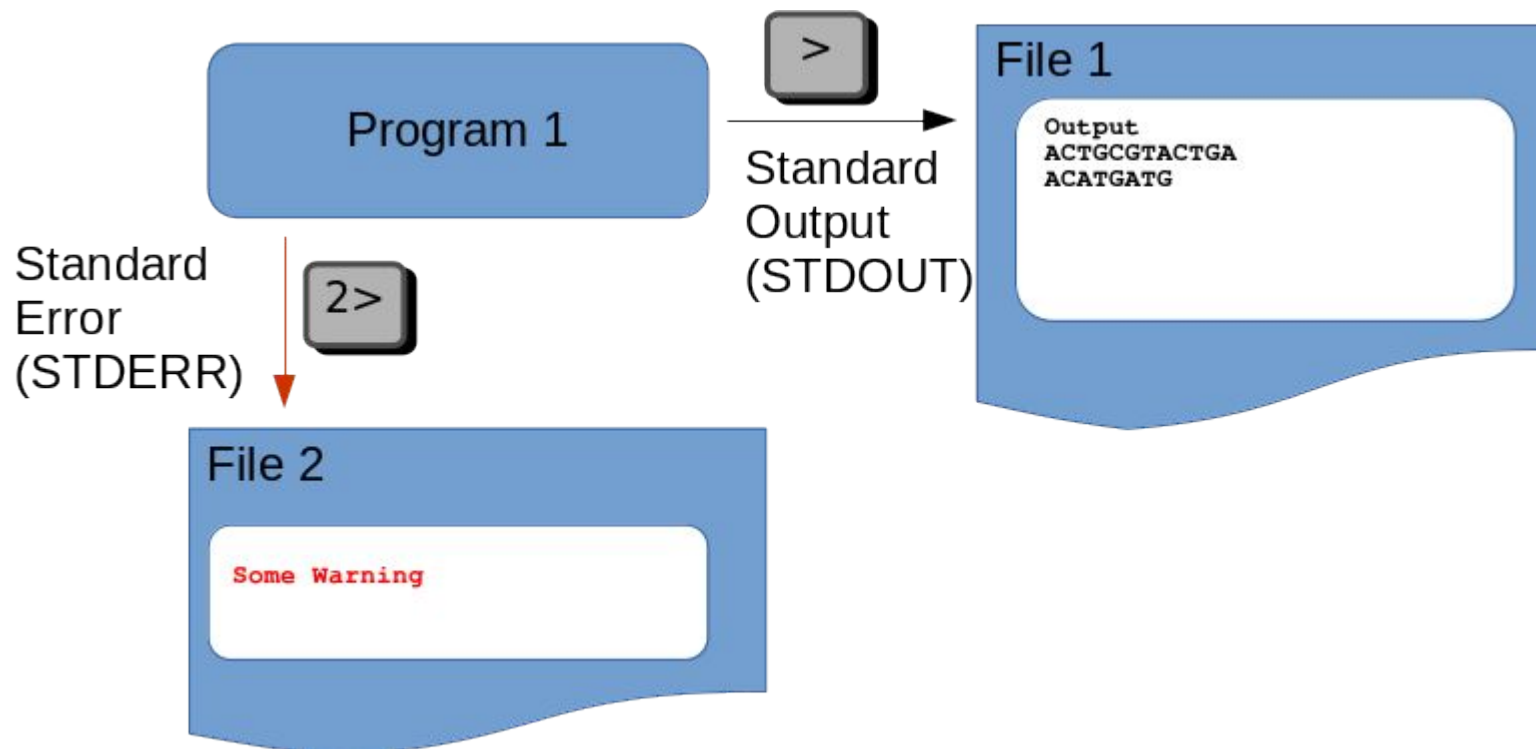
- Many unix commands support streams (files) as input and/or output
- Streams can be **appended to files**



Example: `ls -la ~ >> myHomeDir.txt`

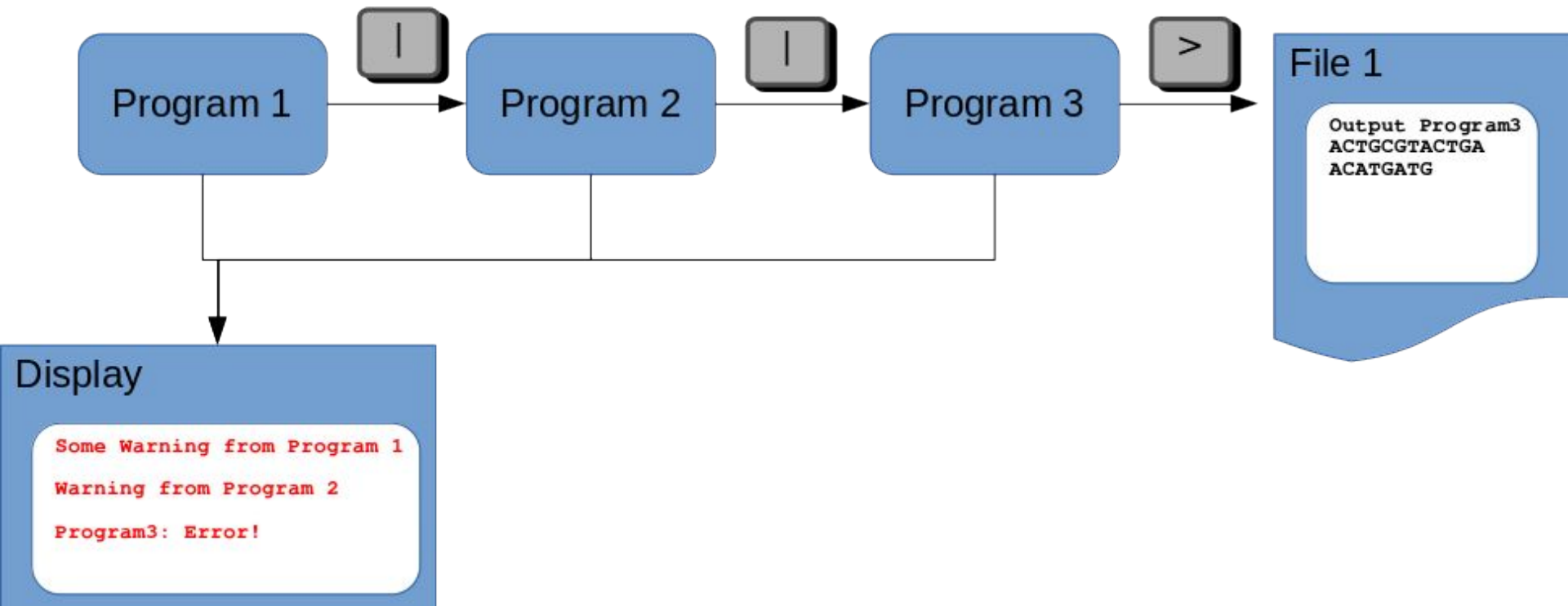
# Linux Command Line Introduction - Part 2

- Many unix commands support streams (files) as input and/or output
- You can also **redirect STDERR**



# Linux Command Line Introduction - Part 2

- Many unix commands support streams (files) as input and/or output
- You can build small **pipelines** by 'connecting' tools



Example: `ls ~ | grep .txt | sort`

# Linux Command Line Introduction - Part 2

- Many unix commands support streams (files) as input and/or output
- Streams can be **redirected**
- **Into files**
  - Stdout: Text that is produced by the program can be redirected by using “>”.
    - Example: `./myProgram > output.txt`
  - Stderr: Error messages are send through stderr and can be redirected using “2>”.
    - Example: `./myProgram 2> error.txt`
  - Stdin: The stdin input stream accepts text as input that can be provided by using Linux pipes
    - Example: `./myProgram1 | ./myProgram2`
- Linux processes
  - You can list all running programs using “**ps**”
  - You can terminate processes by using “**kill** id\_of\_process”.

# Linux Command Line Introduction - Part 2

- **Search for patterns:**
  - **grep:** Search for specific patterns
    - Example: **grep "error" output.log**
    - Search for the term "error" in output.log and displays every line that contains this term
    - Special characters in the pattern:
      - **"^"**: denotes the beginning of the line
      - **"\$"**: denotes the end of a line
- **Search and replace:**
  - **sed:** search and replaces a pattern in a file (stream)
    - syntax: **sed 's/pattern/replacement/'**

# Linux Command Line Introduction - Part 2

- **Wildcards** can be used perform actions on multiple files at the same time
- Common wildcards:

<code>*</code>	Zero or more of any characters
<code>?</code>	Exactly one character – any
<code>[ACGT]</code>	One of the specified characters
<code>[A-Z][0-9]</code>	... works also with alphanumerical ranges
<code>{10..13}</code>	... but for numerical ranges you need 'brace expansion'

- Example:

<code>*.fasta</code>	Matches all files ending with .fasta
<code>sequence_?.fasta</code>	Matches all files beginning with sequence_, ending with .fasta and one character in between
<code>sequences.fast[aq]</code>	Matches sequences.fasta and sequences.fastq



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# Acknowledgements

Cloud Governance Team and Cloud Portal Team:

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