

# Operating System Lab: CS341

## LAB 1: Linux Basics



**Course Instructor: Dr Satendra Kumar**

# Basic Linux Commands

- **mkdir:** The **mkdir** command in Unix-like operating system is used to create new directories.

## Syntax:

- In order to create a single directory the syntax is:

**mkdir directory\_name**

- In order to create multiple directories the syntax is:

**mkdir directory1\_name directory2\_name directory3\_name**

- In order to create a directory with necessary parent directories, we use -p option.

**mkdir -p parent\_directory/sub\_directory**

- **touch**: The touch command creates an empty file if it does not already exist, or updates the access and modification times if it does.

### **Syntax: touch filename**

- **mv**: The mv command in Linux is used to move or rename files and directories.
  - To move a file to a different directory:

### **Syntax: mv file1.txt /path/to/destination/**

- To rename a file:

### **Syntax: mv oldname.txt newname.txt**

- To move a file to a different directory and rename it at the same time

### **Syntax: mv file1.txt /path/to/destination/newname.txt**

➤ **cp:** The cp command in Linux is used to copy files and directories.

- To copy a file to another location:

**Syntax:** `cp file1.txt /path/to/destination/`

- To copy a file and rename it at the same time:

**Syntax:** `cp file1.txt /path/to/destination/newname.txt`

- To copy multiple files to another directory:

**Syntax:** `cp file1.txt file2.txt /path/to/destination/`

- To copy a directory and its contents, you need to use the -r (recursive) option:

**Syntax:** `cp -r dir1 /path/to/destination/`

➤ **ls:** The ls command in Linux is used to list the contents of a directory.

**Syntax:** `ls -R`

Lists files in all subdirectories recursively.

➤ **read:** The read command in Linux is used to take user input and assign it to a variable. It reads a single line of input from standard input (keyboard) or a file.

**Syntax:** `read name`

The -p option allows you to provide a prompt message before reading the input.

**Syntax:** `read -p "Enter your name: " name`

- To check whether a given path is not a directory the following syntax is used:

**Syntax:** `if [ ! -d "path" ]; then`  
                  `# Commands to execute if the path is not a directory`  
          `fi`

- **echo:** Prints a message to the standard output.

**Syntax:** `echo "Message"`

- **for loop:** Loops through a list of items and executes commands for each item.

**Syntax:** `for item in list; do`  
                  `# Commands to execute for each item`  
          `done`

- To append output to a file.

**Syntax:** `echo "Message" >> filename`

- **rm**: Removes files or directories.

**Syntax:** `rm file_name` or `rm directory_name`

- **if-else**: Conditional statement based on the value of a variable.

**Syntax:** `if $variable; then`

**# Commands to execute if variable is true**

`else`

**# Commands to execute if variable is false**

`fi`

- **exit**: Exits the script with a specified status code.

**Syntax:** `exit status_code`

- To check if the number of arguments passed to the script is not equal to 1.

**Syntax:** `if [ "$#" -ne 1 ]; then`

**# Commands to execute if the number of arguments is not 1**

`fi`

- **ps:** Displays information about active processes.

**Syntax:** `ps [options]`

- **ps aux:** Displays all processes for all users in a user-oriented format.

**Syntax:** `ps aux`

- **sort:** Sorts the output of the ps command by a specified field.

**Syntax:** `ps aux --sort=-%cpu`

➤ **awk**: The awk command is a powerful text-processing tool in Unix and Unix-like operating systems. It is used for pattern scanning and processing. awk reads input files line by line, searches for patterns, and performs actions on lines that match the patterns.

**Syntax:** `awk '{ print $1, $2 }' input_file`

By default, awk splits each line into fields based on whitespace (spaces or tabs). Fields can be accessed using \$1, \$2, etc.

## Fields Specifications:

1. **\$1**: Prints the 11th field of the input line. In the context of ps aux, this is typically the command or process name.
2. **\$2**: Prints the 2nd field of the input line. This is the process ID (PID).
3. **\$3**: Prints the 3rd field of the input line. This is the CPU usage percentage.
4. **\$4**: Prints the 4th field of the input line. This is the memory usage percentage.

- **df:** The df command reports the amount of disk space used and available on file systems.

**Syntax:** `df -h /`

Displays the disk usage of the root directory / in a human-readable format.

- **free:** The free command displays information about the system's memory usage.

**Syntax:** `free -h`

Shows the free, used, and total memory available on the system in a human-readable format.