

## **Linux Commands**

Here's a more thorough list of commands found in the file, briefly describing what each one does:

- `ls`: Lists files and directories.
- `ls -l`: Lists files with details in long format.
- `ls -r`: Lists files in reverse order.
- `ls -R`: Recursively lists files in directories.
- `ls -lt`: Lists files sorted by modification time, in long format.
- `ls -la`: Lists all files, including hidden ones, in long format.
- `cd`: Changes the current directory.
- `cd ././`: Changes to the current directory (effectively does nothing).
- `pwd`: Prints the current working directory.
- `cat <file>`: Displays the content of a file.
- `cat > <file>`: Creates a file and allows writing to it.
- `cat >> <file>`: Appends content to a file.
- `touch <file>`: Creates an empty file.
- `mkdir <dir>`: Creates a new directory.
- `mkdir <dir> && cd <dir>`: Creates and enters a new directory.
- `ls *.js`: Lists all .js files in the directory.
- `ls -R *.js`: Recursively lists all .js files in subdirectories.
- `grep`: Searches for patterns in files or outputs.
- `grep <pattern> <file>`: Searches for a pattern in a specific file.
- `ls -lR | grep <pattern>`: Lists files recursively and filters the output based on the given pattern using `grep`.

- `mv <file1> <file2>`: Renames or moves a file or directory.
- `touch <file>`: Creates a new empty file.
- `cp <source> <destination>`: Copies or renames a file. If copying within the same directory with a new name, it acts as a renaming command.
- `cp -r <source> <destination>`: Recursively copies a directory.
- `rm <file>`: Removes a file.
- `rm -r <dir>`: Recursively removes a directory.
- `chmod <mode> <file>`: Changes the file permissions.
- `echo <string>`: Prints a string to the terminal.
- `cat <file>`: Displays the contents of a file.
- `cat >> <file>`: Appends content to a file.
- `head <file>`: Displays the first few lines of a file.
- `tail <file>`: Displays the last few lines of a file.
- `head -n <number> <file>`: Displays the first n lines of a file.
- `tail -n +<number> <file>`: Displays lines starting from the n-th line of a file.
- `wc <file>`: Displays word, line, and character counts of a file.
- `grep <pattern> <file>`: Searches for a pattern in a file.
- `grep <pattern> <file> | wc`: Combines the output of grep with wc to count lines, words, and characters matching the pattern.
- `wc -l`: Displays the line count.
- **CHMOD** -

Here's how the numbering system works:

4 represents read permission (r)

2 represents write permission (w)

1 represents execute permission (x)

0 represents no permissions (-)

These numbers are then added together to create a single digit for each category (owner, group, others). Here's a breakdown:

$7 = 4 + 2 + 1$  (rwx: read, write, and execute)

$6 = 4 + 2$  (rw-: read and write)

$5 = 4 + 1$  (r-x: read and execute)

$4 = 4$  (r--: read only)

$3 = 2 + 1$  (-wx: write and execute)

$2 = 2$  (-w-: write only)

$1 = 1$  (--x: execute only)

$0 = 0$  (---: no permissions)

So, when you see a permission like `chmod 755`, it means:

Owner:  $7 (4+2+1) = \text{rwx}$  (read, write, execute)

Group:  $5 (4+1) = \text{r-x}$  (read, execute)

Others:  $5 (4+1) = \text{r-x}$  (read, execute)

- **`grep "s" -c a2.txt:`**

Counts the number of lines in `a2.txt` that contain the letter "s" and displays the count (2 in this case).

- **`grep "s" -h a2.txt:`**

Searches for lines containing "s" in `a2.txt` but omits the file name in the output, showing only the matching lines.

- **grep "one" -h a2.txt:**

Searches for lines containing "one" in a2.txt and displays them without the file name in the output. Matches exact case.

- **grep "one" -hi a2.txt:**

Searches for lines containing "one" in a2.txt case-insensitively (-i), and displays the matching lines without the file name.

- **grep "one" -hir ./:**

Recursively searches for the word "one" (case-insensitively) in all files and directories starting from the current directory (./), without showing the file names.

- **grep "one" -n a2.txt:**

grep searches for the string "one" in a2.txt.

-n shows line numbers where matches are found.

- **grep "one" -w a2.txt:**

grep searches for the string "one" in a2.txt.

-w restricts the search to whole words only.

- **history** command displays a list of previously executed commands in the terminal, allowing you to view and reuse them. You can use **!number** to re-run a specific command from the history list.

- **bash a3.sh:**

Executes the script a3.sh. It produced an error due to unexpected characters.

- **grep -v "one" a2.txt:**

Displays lines from a2.txt that do not contain the string "one".

- **grep -A 5 "one" a2.txt:**

Shows lines containing "one" and the 5 lines following each match.

- **grep -B 5 "one" a2.txt:**

Shows lines containing "one" and the 5 lines preceding each match.

- **grep -C 5 "one" a2.txt:**

Displays lines containing "one" along with 5 lines before and after each match.

- **sed -i 's/one/two/' a2.txt:**

Replaces the first occurrence of "one" with "two" on each line in the file `a2.txt` and saves the changes directly to the file.

- **sed -ibackup 's/one/two/' a2.txt:**

Replaces the first occurrence of "one" with "two" on each line in `a2.txt` and saves the changes to the file. Additionally, it creates a backup of the original file named `a2.txtbackup`.

- **sed '15,20 s/two/one/' a2.txt:**

Attempts to replace "two" with "one" only in lines 15 through 20 of `a2.txt` but was missing the correct command format. The correct format should be `sed '15,20 s/two/one/' a2.txt`.

- **awk '/two/{print \$0}' a2.txt:**

Prints lines from `a2.txt` that contain the string "two".

- **awk '{gsub(/two/, "three")}{print}' a2.txt:**

Replaces all occurrences of "two" with "three" in each line of `a2.txt` and prints the modified lines.

- **awk 'BEGIN {print "START OF FILE\n-----"} {print} END {print "-----\nEND OF FILE"}' a2.txt:**

Prints "START OF FILE" at the beginning, followed by the contents of `a2.txt`, and then prints "END OF FILE" at the end.

- **awk '{print \$1, \$2}' a2.txt:**

Prints the first and second fields of each line from `a2.txt`. If there are fewer than two fields, only the first field is printed.

- **awk -F "," '{print \$1, \$2}' a2.txt:**

Sets the field separator to a comma and prints the first and second fields of each line from `a2.txt`. This command didn't change the output since `a2.txt` does not use commas as separators.

- **awk '{count[\$1]++} END {print count["two"]}' a2.txt:**

Counts occurrences of each unique value in the first field of `a2.txt` and prints the count of the value "two".