

# Unit II File Management commands Filters (Part I)

# Sorting Files

- To sort the contents of file

`$ sort filename`

- to sort multiple file together

`$ sort f1 f2 f3`

- to save output to a file named “final”

`$ sort -o final filename`

- to sort only unrepeated data

`$ sort -u filename`

# Sorting Files

- to sort data from terminal

```
$ sort
```

- to combine the contents of terminal & a file & then perform sorting

```
$ sort - filename
```

- to merge only sorted files

```
$ sort -m f1 f2
```

- to sort the file in reverse/descending order

```
$ sort -r f1
```

# Information on Files

- To generate one or more of the following statistics:  
newline count, word count, and character count

## **wc filename**

A line is any group of characters not containing a newline

A word is a group of characters not containing a space, tab or newline

A character is the smallest unit of information, and includes a space, tab, and newline

# Information on Files

- to count only lines

```
$ wc -l filename
```

- to count only words

```
$ wc -w filename
```

- to count only characters

```
$ wc -c filename
```

- to display the length of longest line in a file

```
$ wc -L filename
```

# Uniq command

It discards all the successive identical lines except one from the input and writes the output.

- `uniq filename`
- `uniq -c filename -- with count`

The options of `uniq` command are:

- `c` : Count of occurrence of each line.
- `d` : Prints only duplicate lines.
- `D` : Print all duplicate lines
- `f` : Avoid comparing first N fields.
- `i` : Ignore case when comparing.
- `s` : Avoid comparing first N characters.
- `u` : Prints only unique lines.
- `w` : Compare no more than N characters in lines

# What is common?

- To find the common lines in two “SORTED” files

```
$ comm f1 f2
```

- it displays a three-columnar output, first column contains lines unique to f1, second column contains lines unique to f2, while third column contains lines common to both f1 and f2

- to drop any column, use column number as option

```
$ comm -2 f1 f2
```

```
$ comm -12 f1 f2
```

# File Comparison: cmp command

- Used to compare the two files byte by byte
- Helps you to find out whether the two files are identical or not.
- Reports the location of the first mismatch.
- Displays no message and simply returns the prompt if the files compared are identical.

## Syntax

```
cmp [OPTION]... file1 [file2 [skip1 [skip2]]]
```

File1 and file2 are file names

skip1 and skip2 specify the number of bytes to skip at the beginning of each file which is zero by default



```
cmp f1 f2
```

the files are compared byte by byte, if found a mismatch, location is displayed, but if files are identical, no message is displayed & the prompt is returned

```
cmp -i 5:6 file1.txt file2.txt
```

Will skip 5 bytes from first file and 6 bytes from second file

```
cmp -n 30 file1.txt file2.txt
```

Will search 30 bytes from the starting

# diff command

- diff file1 file2
- The first line of the **diff** output will contain:
  - line numbers corresponding to the first file,
  - a letter (**a** for *add*, **c** for *change*, or **d** for *delete*), and
  - line numbers corresponding to the second file.
- Lines preceded by a < are lines from the first file;
- lines preceded by > are lines from the second file.

# pg command

- Displays the contents of text files, one page at a time
- Syntax: `pg [-number] [-p string] [-cefnrs] [+line] [+/pattern/] [file...]`

# Head command

- Print the first N number of data/lines of the given input.
- By default, it prints the first 10 lines of the specified files.
- If more than one file name is provided then data from each file is preceded by its file name.
- **Syntax:**
  - head [OPTION]... [FILE]...

# Viewing Files - head

to view first 5 lines from file.txt

**\$head -n 5 file.txt**

to view first 10 bytes from the beginning of file

\$ head -c 10 filename

\*also counts newline character

Try head without any arguments !!!

-Waits for user input

# Viewing Files - head

- to view name of the file along with the contents

```
$ head -v filename
```

- to view first few lines of more than 1 file

```
$ head f1 f2 f3
```

- to suppress filenames when multiple files are being used

```
$ head -q f1 f2 f3
```

# Tail command

- Print the last N number of data of the given input.
- By default it prints the last 10 lines of the specified files.
- If more than one file name is provided then data from each file is preceded by its file name
- **Syntax:**
  - `tail [OPTION]... [FILE]...`

# Viewing Files - Tail

- To display last few lines of the file

`$ tail filename`      {by default shows 10 lines}

- to view last n lines

`$ tail -n filename`

- to view desired “n” no. of bytes from the ending of file

`$ tail -c n filename`      {also counts newline character & n is mandatory}

- Try tail without any arguments !!!



# Viewing Files - Tail

- to view name of the file along with the contents

```
$ tail -v filename
```

- to view first few lines of more than 1 file

```
$ tail f1 f2 f3
```

- to suppress filenames when multiple files are being used

```
$ tail -q f1 f2 f3
```

# Viewing Files-more

- To view files having content not fitting in the screen at a time

\$ more filename

- to view “n” no. of lines

\$ more -n filename

- to prompt the user with the message "[Press space to continue, 'q' to quit.]" and will display "[Press 'h' for instructions.]" when an illegal key is pressed

\$ more -d filename

- to squeeze multiple blank lines into one

\$ more -s filename

# Viewing Files

- press “space bar” or “f” to move one page forward
- Press [Enter] to move one line forward
- Press “2f” to move 2 pages forward
- Press “b” to move one page backwards
- Press “3b” to move 3 pages backward
- Press “q” to quit
- Press “h” to get help
- !cmd to execute UNIX command cmd

# less command

- *Less* is a command similar to *more*, but which allows backward movement in the file as well as forward movement.
- *less* does not have to read the entire input file before starting, so with large input files it starts up faster than text editors like *vi*

# cut command

- To select a list of columns or fields from one or more files.
- Specify either the -c option to cut by column or -f to cut by fields.
- Fields are separated by tabs unless you specify a different field separator with -d
- **Based on column (-c)**
  - cut -c 4 f1 only 4<sup>th</sup> column
  - cut -c -3 f1 beginning of line to 3<sup>rd</sup> column
  - cut -c 2-3,6-8 f1 2<sup>nd</sup> to 3<sup>rd</sup> & 6<sup>th</sup> to 8<sup>th</sup> column
  - cut -c 6-7,9- f1 6<sup>th</sup> to 7<sup>th</sup> & 9<sup>th</sup> to end of the line

# Based on fields (-f)

- `cut -f 1,4 f1` (default field separator is tab)
- `cut -d ":" -f 2,5-8 f1` give respective fields, for all the lines containing ":" or not
- `cut -s -d ":" -f 2,4 f1` give respective fields, only for the lines containing delimiter
- `cut - -complement -f1 filename` gives all fields except first field
- `cut -d ":" -f1,4 - -output-delimiter="$"*" filename` changes the output delimiter to \* rather than ":" by default

# Paste command

- Merges the lines from multiple files
- paste [options] files-list
  - Options
    - -d : Specify of a list of delimiters.
    - -s : Paste one file at a time instead of in parallel.
- paste file1 file2
- paste -d"|" file1 file2
- paste -s file1 file2
  - The paste command reads each file in sequentially. It reads all the lines from a single file and merges all these lines into a single line.
- Specifying multiple delimiters.
  - paste -d"|," file1 file2 file3

# Pipe

- Shells' sequence of interpretation of a command
- Used to connect two commands together so that the output from one program becomes the input of the next program.
- To make a pipe, put a vertical bar (|) on the command line between two commands.



```
echo hi | wc -c
```

```
echo "hi" | echo -e "\n welcome" | wc -l
```

```
ls -l | sort -r
```

```
ls -l a* | wc -l
```

```
echo -n "welcome" ; ls a* | wc -l
```

- Assignment 3.1