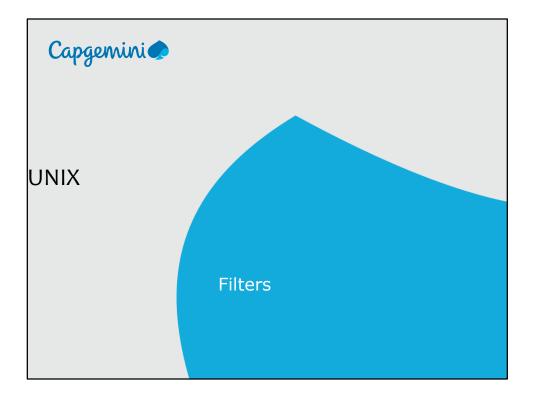
## **Instructor Notes:**



## **Instructor Notes:**

## Lesson Objectives

In this lesson, you will learn:

- Filter commands in UNIX:
  - · Simple Filters
  - Advance Filters



### **Instructor Notes:**

3.1: Simple Filters

### What is a Filter?

Filters are central tools of the UNIX tool kit.

Commands work as follows:

- Accept some data as input.
- Perform some manipulation on the inputted data.
- Produce some output.

Most of them work on set of records, with each field of a record delimited by a suitable delimiter.

When used in combination, they can perform complex tasks too.

### **Simple Filters:**

Filters are central tools of the UNIX tool kit. These commands accept some data as input, perform some manipulation on it and produce some output. Most of them work on set of records, with each field of a record delimited by a suitable delimiter. They serve as useful text manipulators. When used in combination, they can perform complex tasks as well. This lesson discusses some commonly used simple filters.

### **Instructor Notes:**

### head Command

The head command, by default, will display the first 10 lines of a file.

• **Example 1:** To display fist 10 lines from file employee:

\$head employee

• Example 2: To display first 5 lines from file employee:

\$head -5 employee

Single command can be used to display lines from more than one file.

\$ head -1 PuneEmp PKPEmp

### The head Commands:

These are simple horizontal filters.

Using head, it is possible to display beginning of one or more lines form files. By default, the first 10 lines are displayed. Incase a numeric line count argument is specified, the command would display those many lines from the beginning of the file.

To display first 10 lines of the file bigfile, use the following syntax:

```
$ head bigfile
```

Output: cfile1.lst

cfile2.lst errfile file1.txt file2.txt file3.txt mail newdir1 newdir2 newfile.txt

To display first 3 lines of the file bigfile, use the following syntax:

\$ head -3 bigfile

\$ head -1 listing bigfile

Output:

cfile1.lst cfile2.lst errfile

Output:

==> listing <== cfile1.lst ==> bigfile <== cfile1.lst

### **Instructor Notes:**

### tail Command

The tail command is useful to display last few lines or characters of the file.

• Example 1: To display last ten lines from employee:

\$tail employee

Example 2: To display last seven lines:

\$tail -7 employee

• **Example 3:** To display lines from the 10<sup>th</sup> line till end of the file:

\$tail +10 employee

• Example 4: To display last 5 characters of the file:

\$tail -5c employee

### The tail Commands:

### Using the tail command:

Using tail, the end of file can be displayed – default being last 10 lines.

### \$ tail -2 listing

Output: result

testdir2 mail

newdir1

newdir2 newfile.txt

\$ tail +20 bigfile

Output: result

Testdir2

To display last 6 characters from bigfile, use the following syntax:

\$ tail -6c bigfile

le.txt

### **Instructor Notes:**

### cut Command

The cut command retrieves selected fields from a file.

\$ cut [options] <filename>

#### Options:

-c : selects columns specified by list
-f : selects fields specified by list
-d : field delimiter (default is tab)

### cut Command:

You can slice the file vertically with the cut command, and paste laterally with the paste command.

The cut command can be used to retrieve specific column information from a file. In case of fixed record formats, the -c (columns) option can be used to specify column positions. If a delimiter has been used, -f (field) in conjunction with -d (delimiter) options can be used for retrieval. The default delimiter is tab.

### \$ cat bookDetails.lst

Output: 1001|Unix for You |375 1002|Learning Unix |250 1003|Unix Shell Programming |450 1004|Unix Device Drivers |375 1005|Advanced Unix Concepts |450

The following command will display first 4 characters followed by 31st to 35th characters:

### \$ cut -c1-4,31-35 bookDetails.lst

Output: 1001|375 1002|250 1003|450 1004|375 1005|450

### **Instructor Notes:**

## cut Command

• **Example 1:** To display 2<sup>nd</sup> and 3<sup>rd</sup> field from file bookDetails.lst:

\$ cut -d"|" -f2,3 bookDetails.lst

• **Example 2:** To display characters from 1<sup>st</sup> to 4<sup>th</sup> and 31<sup>st</sup> to 35<sup>th</sup> from file bookDetails.lst:

\$ cut -c1-4,31-35 bookDetails.lst

### cut Commands:

To display 2nd and 3rd field from file bookDetails.lst: In the given command on the above slide, the –d option specifies field delimiter is |. Hence it will consider that in bookDetails file there are 3 fields separated by '|' character. The –f option will specify to display 2nd and 3rd field.

\$ cut -d"|" -f2,3 bookDetails.lst

Output: Unix for You | 375

Learning Unix | 250
Unix Shell Programming | 450
Unix Device Drivers | 375
Advanced Unix Concepts | 450

### **Instructor Notes:**

## paste Command

The paste command is used for horizontal merging of files.

\$paste <file1><file2><Enter>

- Options : -d (Field delimiter)
- Example 1: To paste enum.lst and ename.lst files:

\$ paste enum.lst ename.lst

 Example 2: To paste enum.lst and ename.lst files with '|' character as delimiter:

\$ paste -d'l' enum.lst ename.lst

### paste Command:

Several files can be pasted laterally, with specific delimiters, with the paste command.

### \$ cat enum.lst

Output: 1010

2021

3718

4135

5765

### \$ cat ename.lst

Output: Abc

Zxc

Qwe

Jkl

Uio

### \$ paste enum.lst ename.lst

### \$ paste -d"|" enum.lst ename.lst

Output: 1010	Abc		Output:	1010 Abc
	2021	Zxc		2021 Zxc
	3718	Qwe		3718 Qwe
	4135	Jkl		4135 Jkl
	5765	Uio		5765 Uio

### **Instructor Notes:**

## sort Command

The sort command is useful to sort file in ascending order.

\$sort <filename>

- Options are:
  - · -r : Reverse order
  - · -n : Numeric sort
  - -f : Omit the difference between Upper and lower case alphabets
  - · -t : Specify delimiter
  - · -k : to specify fields as primary or secondary key
- Example:

```
$ sort -t"|" +1 bookDetails.lst
$sort -k3,3 -k2,2 employee
```

#### sort Command:

### Sorting a file with the sort command:

The sort command sorts a file (which may or may not contain fixed length records) on line by line basis. Default sorting is in the ascending ASCII order, which can be reversed by using the –r option.

Sorting can be done on one or more fields by specifying the delimiter using –t option. It is also possible to specify character positions within fields.

Using the -m option, it is also possible to merge any number of sorted files.

Since the sorting is done on the basis of ASCII collating sequence, incase of sorting of numbers, -n option needs to be used.

### \$ sort -t"|" +1 bookDetails.lst

Output: 1005|Advanced Unix Concepts | 450 1002|Learning Unix | 250 1004|Unix Device Drivers | 375 1003|Unix Shell Programming | 450 1001|Unix for You | 375

To sort file employee on 3rd field as primary key and 2nd field as secondary key, use the following syntax:

```
$ sort -t"|" -k3,3 -k2,2 Employee
```

To consider only 3rd and 4th character from 2nd field for sorting employee file, use the following syntax:

```
$sort -t"|" -k2.3,2.4 employee
```

### **Instructor Notes:**

## uniq Command

The uniq command fetches only one copy of redundant records and writes the same to standard output.

- u option: It selects only non-repeated lines.
- -d option: It selects only one copy of repeated line.
- -c option: It gives a count of occurrences.

To find unique values, the file has to be sorted on that field.

• Example: To find unique values from file duplist.lst

\$ uniq duplist.lst

### uniq Command:

The uniq command requires a sorted file as input. It fetches only one copy of redundant records and writes the same to standard output.

The **–u** option can be used to select only non-repeated lines, while the **–d** option can be used to select only one copy of repeated line. It is also possible to get a count of occurrences with the -c option.

### Example 1:

### \$ cat duplist.lst

Output: 1

34 23

23

30

#### Example 2:

### \$ sort -n duplist.lst | uniq

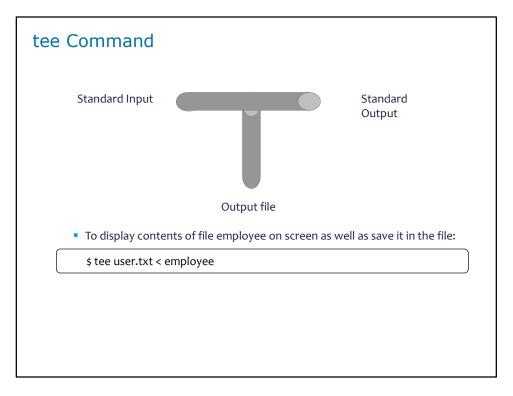
# Output: 1

23

30

34

### **Instructor Notes:**



#### tee Command:

The tee command copies the standard input to the standard output and also to the specified file.

If it is required to see the output on screen as well as to save output to a file, the **tee** command can be used. The **tee** command uses both standard input and standard output.

To display list of users and it's count both on screen, use the following:

Who | tee /dev/tty | wc -l

If we give command as **who|wc -l**, it will display only number of users on screen. However, in the above command **tee** will save the o/p to /dev/tty file which is terminal device file. Hence the o/p of **who** will be displayed on screen and also will be transferred as i/p to **wc** command.

In the following command, **sort** will sort the file and o/p will be transferred to **tee** command. It will display o/p on screen as well as store it in file **sorted\_file.txt**. The o/p will be given to **uniq** command which will give count of lines in the file. The head will find first 12 lines and store it in **top12.txt** file.

\$sort somefile.txt | tee sorted\_file.txt | uniq -c | head 12 > top12.txt

### **Instructor Notes:**

3.2: Advanced Filters

### find Command

The find command locates files.

find <path list> <selection criteria> <action>

 Example 1: To locate the file named .profile starting at the root directory in the system -print specify the action:

\$ find / -name .profile -print

 Example 2: To locate the file named myfile starting at the root directory in the system

find / -type f -name "myfile" -print

### find Command (locating files with find):

The **find** command is used to find files matching a certain set of selection criteria. The **find** command searches recursively in a hierarchy, and also for each pathname in the pathname-list (a list of one or more pathnames specified for searching).

The syntax of the find command is given in the following format:

\$ find <path list> <selection criteria> <action>

The find command first looks at all the files in the directories specified in the path list. Subsequently, it matches the files for one or more selection criteria. Finally it takes action on those selected files.

\$ find / -name .profile -print

The above command will locate the **.profile** files in the system.

\$find . -name \*stat

The above command will locate all file names ending with stat.

### **Instructor Notes:**

## grep Command

The syntax for grep command is as follows:

grep <options> <pattern> <filename(s)>

• **Example:** The following example will search for the string Unix in the file **books.lst**. The lines which match the pattern will be displayed.

grep 'Unix' books.lst

### grep Command:

The **grep** command is used to locate a pattern / expression in a file / set of files. There are many options that are available for obtaining different types of outputs.

The syntax for the grep command is as follows:

grep <options> <pattern> <filename(s)>

The grep command scans the file(s) specified for the required pattern, and outputs the lines containing the pattern. Depending on the options used, appropriate output is printed. The grep command compulsorily requires a pattern to be specified, and the rest of the arguments are considered as file names in which the pattern has to be searched.

### **Instructor Notes:**

## grep Command

### Options of grep:

- c : It displays count of lines which match the pattern.
- n : It displays lines with the number of the line in the text file which match the pattern.
- v : It displays all lines which do not match pattern.
- i : It ignores case while matching pattern.
- -w: It forces grep to select only those lines containing matches that form whole words

### **Instructor Notes:**

## grep Command

• **Example 1:** To print all lines containing "rose" regardless of case:

\$grep -i rose flower.txt

• Example 2: To print all lines containing "rose" as a word:

\$grep -w rose flower.txt

• Example 3: To print all lines not containing "rose":

\$grep -v rose flower.txt

## **Instructor Notes:**

## grep Command

## Regular Expression:

Expression	Description
^ (Caret)	match expression at the start of a line, as in ^A.
\$ (Question)	match expression at the end of a line, as in A\$.
\ (Back Slash)	turn off the special meaning of the next character, as in \^.
[](Brackets)	match any one of the enclosed characters, as in [aeiou]. Use Hyphen "-" for a range, as in [0-9].
[^]	match any one character except those enclosed in [], as in [^o-9].
. (Period)	match a single character of any value, except end of line.
* (Asterisk)	match zero or more of the preceding character or expression.
\{x,y\}	match x to y occurrences of the preceding.
\{x\}	match exactly x occurrences of the preceding.
\{x,\}	match x or more occurrences of the preceding.

## **Instructor Notes:**

## grep Command

## Examples of Regular Expression:

Example	Description
grep "smile" files	search files for lines with 'smile'
grep '^smile' files	'smile' at the start of a line
grep 'smile\$' files	'smile' at the end of a line
grep '^smile\$' files	lines containing only 'smile'
grep '\^s' files	lines starting with '^s', "\" escapes the ^
grep '[Ss]mile' files	search for 'Smile' or 'smile'
grep 'B[oO][bB]' files	search for BOB, Bob, BOb or BoB
grep '^\$' files	search for blank lines
grep '[0-9][0-9]' file	search for pairs of numeric digits

# grep Command: Some more examples:

Example	Description	
grep '^From: ' /usr/mail/\$USER	list your mail	
grep '[a-zA-Z]'	any line with at least one letter	
grep '[^a-zA-Z0-9]	anything not a letter or number	
grep '[0-9]\{3\}-[0-9]\{4\}'	999-9999, like phone numbers	
grep '^.\$'	lines with exactly one character	
grep "smug"	'smug' within double quotes	
grep ""*smug"*'	'smug', with or without quotes	
grep '^\.'	any line that starts with a Period "."	
grep '^\.[a-z][a-z]'	line start with "." followed by 2 lowercase letters	

### **Instructor Notes:**

## fgrep Command

The fgrep command is similar to grep command. Syntax:

\$fgrep [-e pattern list][-f pattern-file][pattern][Search file]

The fgrep command is useful to search files for one or more patterns, which cannot be combined together.

It does not use regular expressions. Instead, it does direct string comparison to find matching lines of text in the input.

### fgrep Command:

The **fgrep** command can also accept multiple patterns from command line as well as a file. However, it does not accept regular expressions – only fixed strings can be specified. The **fgrep** command is faster than **grep** and **egrep**, and should be used while using fixed strings.

The **egrep** and **fgrep** commands to some extent overcome the limitations of **grep**. However, the principal disadvantage of the grep family of filters is that there are no options available to identify fields. Also it is very difficult to search for an expression in a field. This is where the **awk** command is very useful.

### **Instructor Notes:**

## fgrep Command

### Options of fgrep command:

- -e pattern\_list :
  - It searches for a string in pattern-list.
- -f pattern-file :
  - It takes the list of patterns from pattern-file.
- pattern
  - It specifies a pattern to be used during the search for input.
  - It is same as grep command.
- E.g To search employee file for all patterns stored in mypattern file
   \$ fgrep -f mypattern employee.lst

### fgrep Command:

Example using fgrep:

\$ cat stud.lst

Output:Roo1|Pratik Sharma|425Roo2|Pallavi V.|398Roo3|Pratibha Aggarwal|400

Roo4|Preeti Agrawal |390 Roo5|Prerana Agarwal |421 Roo6|Pranita aggarwal |380

\$cat mypattern

Output: Pratik

Pratibha

\$ fgrep –f mypattern stud.lst

Output: Roo1|Pratik Sharma | 425

Roo3|Pratibha Aggarwal |400

### **Instructor Notes:**

## egrep Command

The egrep command works in a similar way. However, it uses extended regular expression matching.

Syntax:

```
egrep [ -e pattern_list ] [-f file ] [ strings ] [ file]
```

 Example: To find all lines with name "aggrawal" even though it is spelled differently:

```
$ egrep '[aA]gg?[ar]+wal' stud.lst
```

### egrep Command (extending grep):

The egrep command offers all the options of the grep command. In addition, it is possible to specify alternative patterns. The table given below gives the extended

Expression	Significance
ch+	Match with 1 or more occurrences of character ch
ch?	Match with 0 or more occurrences of character ch
exp1 e xp2	Match with expressions exp1 or exp2
(a1 a2) a3	Match with expression a1a3 or a2a3

Some examples of using egrep are given:

Output: Roo3|Pratibha Aggarwal |400 Roo4|Preeti Agrawal |390 Roo5|Prerana Agarwal |421 Roo6|Pranita aggarwal |380

### **Instructor Notes:**

## **Summary**

In this lesson, you have learnt:

- The head and tail filter commands filter the file horizontally.
- The cut and paste commands filter the file vertically.
- -m option of sort command is used to merge two sorted files.
- The tee command helps us to send o/p to standard o/p as well as to file.
- grep, fgrep, and egrep commands use to search files for some pattern.



**Filters** UNIX

## **Instructor Notes:**

## **Review Questions**

Question 1: \_\_\_\_ command to display directory listing on screen as well as store it in dirlist.lst.

Question 2: \_\_\_\_ filter commands filter file vertically?

Question 3: \_\_\_\_ filter commands filter file horizontally?

