



Lesson Objectives

In this lesson, you will learn:

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



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.**

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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.

head Command

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

- Example 1: To display first 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
```

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The head Commands:

These are simple horizontal filters.

Using head, it is possible to display beginning of one or more lines from files. By default, the first 10 lines are displayed. In case 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
```

Output:

```
cfile1.lst
cfile2.lst
errfile
```

```
$ head -1 listing bigfile
```

Output:

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

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 10th line till end of the file:

```
$tail +10 employee
```

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

```
$tail -5c employee
```

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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
```

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)

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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
```

cut Command

- **Example 1:** To display 2nd and 3rd field from file bookDetails.lst:

```
$ cut -d'|' -f2,3 bookDetails.lst
```

- **Example 2:** To display characters from 1st to 4th and 31st to 35th from file bookDetails.lst :

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

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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 |

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'|' enum.lst ename.lst
```

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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
```

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

```
$ paste -d'|' enum.lst ename.lst
```

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

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
```

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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`

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
```

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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:

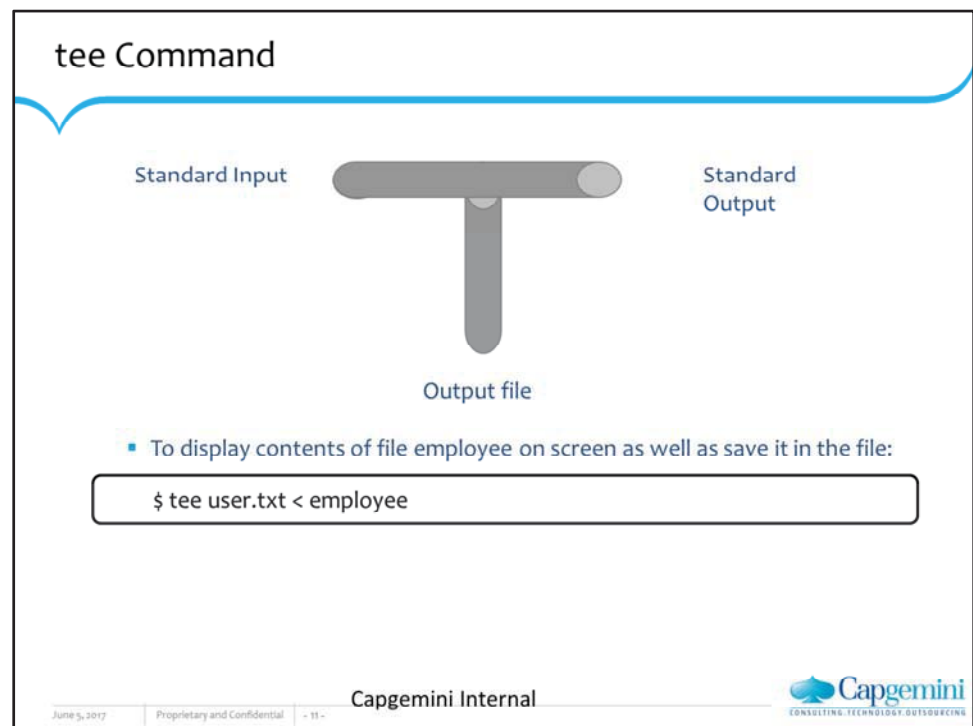
```
$ cat duplist.lst
```

Output: 1
34
30
34
1
23
23
4

Example 2:

```
$ sort -n duplist.lst | uniq
```

Output: 1
4
23
30
34

**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
```

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
```

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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**.


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
```

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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.

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

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
```

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 [^0-9] . |
| . (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. |

grep Command

Examples of Regular Expression:

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

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grep Command: Some more examples:

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

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.

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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.

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`

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fgrep Command:

Example using fgrep:

```
$ cat stud.lst
```

Output:

| | |
|------------------------|-----|
| R001 Pratik Sharma | 425 |
| R002 Pallavi V. | 398 |
| R003 Pratibha Aggarwal | 400 |
| R004 Preeti Agrawal | 390 |
| R005 Prerana Agarwal | 421 |
| R006 Pranita aggarwal | 380 |

```
$cat mypattern
```


Output: Pratik
Pratibha

```
$ fgrep -f mypattern stud.lst
```

Output:

| | |
|------------------------|-----|
| R001 Pratik Sharma | 425 |
| R003 Pratibha Aggarwal | 400 |

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
```

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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 exp2 | Match with expressions exp1 or exp2 |
| (a1 a2)a3 | Match with expression a1a3 or a2a3 |

Some examples of using **egrep** are given:

```
$ cat stud.lst
```

```
Output:  Roo1|Pratik Sharma      |425
         Roo2|Pallavi V.        |398
         Roo3|Pratibha Aggarwal |400
         Roo4|Preeti Agrawal    |390
         Roo5|Prerana Agarwal   |421
         Roo6|Pranita aggarwal  |380
```

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

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

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.



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?

