ALX LESSON 0x02. Shell, IO Redirections and filters

Shell - Bash

TABLE OF CONTENTS

01

Overview topics

03

Quiz questions 02

Learning Objectives

04

hands on lab practice

01 OVERVIEW topics

Ubuntu Shell Commands

Topics

how to redirect standard output to a file

how to get standard input from a file instead of keyboard

how to send output from one program to the input of another program

how to combine commands and filters with redirections

what are spectial characters

understand what do the white spaces, single qoutes, double qoutes, backslash, comment, pipe, command separator, tilde and how and when to use them

Ubuntu Shell Commands

Topics

how to display a line of text

how to concatenate files and print on the standard output

how to reverse a string

how to remove sections from each line of lines

Ubuntu Shell Commands

Topics

echo cat WC head tail sort uniq tr grep cut sed awk tee find

Slides On Telegram

https://t.me/alx_2023

Ubuntu Shell Commands

Topics



02

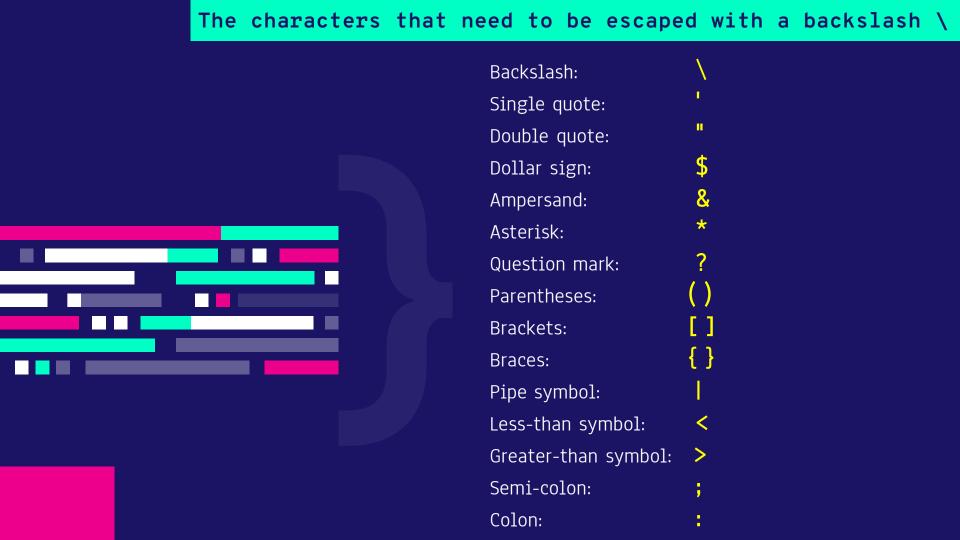
Learning Objectives

echo

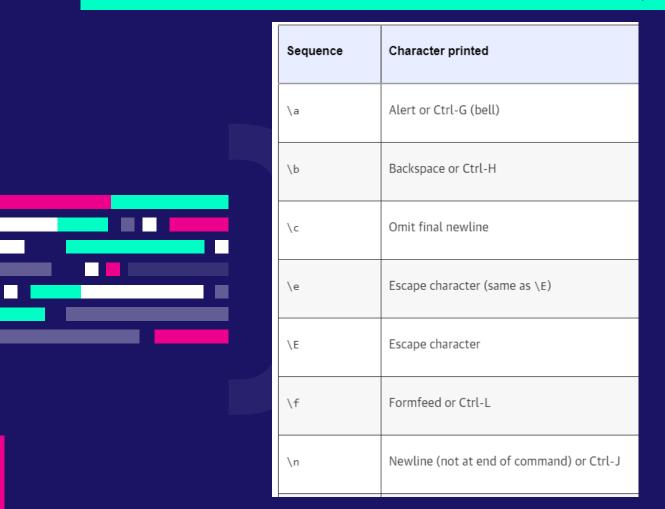
	Parameter	Description	Example
	`-n`	Suppresses the trailing newline character	`echo -n "Hello world"` prints "Hello world" without a newline at the end.
	`-e`	Enables the interpretation of escape sequences	`echo -e "Hello\tworld"` prints "Hello" followed by a tab character and then "world".
	`-E`	Disables the interpretation of escape sequences	`echo -E "Hello\tworld"` prints "Hello\tworld" without interpreting the tab character.
	`-\\`	Escapes the next character and treats it as a literal character	`echo "Hello \\world"` prints "Hello \world".
	1.1	Treats all subsequent arguments as positional parameters	`echo - "Hello world"` prints "Hello world" and any subsequent arguments as separate positional parameters.
	``	Indicates the end of options	`echoHello` prints "-Hello" and treats it as a string, rather than an option.
	`-help`	Displays help information for the `echo` command	`echohelp` displays help information for the `echo` command.

echo

```
echo "hello world!"
echo "my bank balance is: $500"
echo "my bank balance is: \$500"
echo $PWD
echo "The current date and time is $(date)"
echo number_{1..5}
echo front-{A,B,C}-Back
echo front-{A..C}-Back
echo {A..Z}
echo *
echo "Hello\tworld\nWelcome to\n\nUnix/Linux"
echo -e "Hello\tworld\nWelcome to\n\nUnix/Linux"
```



The characters that need to be interpreter -e



	\r	Return (Enter) or Ctrl-M
	\t	Tab or Ctrl-I
	\v	Vertical Tab or Ctrl-K
H	\n <i>nnn</i>	The eight-bit character whose value is the octal (base-8) value nnn where nnn is 1 to 3 digits
	\0 nnn	The eight-bit character whose value is the octal (base-8) value <i>nnn</i> where <i>nnn</i> is 0 to 3 digits
	∖×HH	The eight-bit character whose value is the hexadecimal (base-16) value HH (one or two digits)
	\\	Single backslash

shell redirection operators

Operator	Description	Example
`<`	Input Redirection: read input from a file or command	`command < input.txt`
`>`	Output Redirection: write output to a file or device	`command > output.txt`
`>>`	Append Output Redirection: append output to a file	`command >> output.txt`
`2>`	Error Redirection: write error output to a file	`command 2> error.txt`
`&<`	Input/Output Redirection: read input and write output	`command &< input_output.txt`
`&>`	Input/Output Redirection: write output and error output	`command &> output_error.txt`
` \ `	Input/Output Redirection: read input and write output	`command <> input_output.txt`
`<<`	Here Document: redirect multiple lines of input to a command	`command << END`
`<<<`	Here String: redirect a string as input to a command	`command <<< "hello"`

shell redirection operators

File	File Descriptor	
Standard Input STDIN	0	
Standard Output STDOUT	1	
Standard Error STDERR	2	

echo "hello, world!" > outputfile.txt
echo "again, hello" >> outputfile.txt

wc << outputfile.txt
cat <<< outputfile.txt</pre>

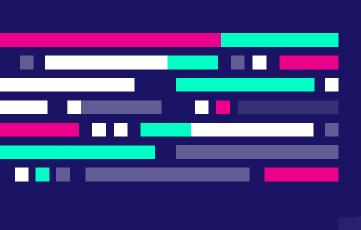
wc < outputfile.txt

2>&1

ls /path/to/missing/directory 2> error.txt
ls /path/to/nonexistent/directory &> output_and_error.txt
ls {.,/path/to/nonexistent/directory} &> output_and_error.txt
ls {.,/path/to/nonexistent/directory} > output_and_error.txt

shell redirection operators

`&`	Runs a command in the background
`;`	Separates multiple commands on a single command line, runs them sequentially
`&&`	Separates multiple commands on a single command line, runs them conditionally (only runs the next command if the previous command succeeds)



The I operator connects the output of one command to the input of another command, allowing for powerful command pipelines to be constructed.

ls -l | grep ".txt"
ls -l ; echo "done"
ls -l && echo "okay"
ls fredkorhan && echo "not okay"
echo Don't work if the command before it
works first

cat

The cat command is a utility command in Linux. One of its most common usages is to print the content of a file onto the standard output stream. Other than that, the cat command also allows us to write some texts into a file.

Display the contents of a file: cat filename.txt

Concatenate two or more files and display the output:

cat file1.txt file2.txt

Append the contents of one file to another file: cat file1.txt >> file2.txt



Create a new file and add content to it: cat > newfile.txt

This will create a new file called newfile.txt and allow you to type in content. Press Ctrl-D to save and exit.



The wc command in Linux is a command-line utility used to count the number of lines, words, and characters in a file or input from the standard input.

Count the number of lines, words, and characters in a file:

wc filename.txt

Count the number of lines, words, and characters in multiple files:

wc file1.txt file2.txt

Count the number of lines, words, and characters in the output of another command:

ls I wc

WC

Options:

-c, --bytes print the byte counts

-m, --chars print the character counts

-l, --lines print the newline counts

-L, --max-line-length print the maximum display width

-w, --words print the word counts

wc -m filename.txt

wc -l filename.txt

wc -w filename.txt

wc -L filename.txt

head

The head command in Linux is a command-line utility used to display the first few lines of a file or input from the standard input.

This will display the first 10 lines of filename.txt by default:

head filename.txt

Display a specified number of lines from a file: head -n 15 filename.txt

This will display the first 10 lines of file1.txt and file2.txt by default.

head file1.txt file2.txt

To display the first 50 bytes of a file, you would use the following command: head -c 50 filename.txt

head

To display the file name before each output line head -v filename.txt



tail

The tail command is used to display the last few lines of a file. Here are some examples of how to use tail: tail file.txt

This command will display the last 5 lines of the file file.txt
tail -n 5 file.txt

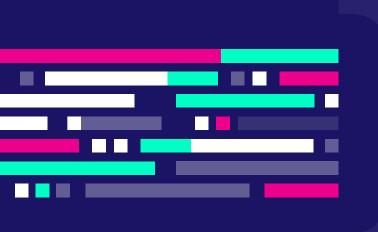
This command will display the last 10 lines of the file log.txt, and then keep the file open and display any new lines that are added to the file in real time tail -f log.txt

This command will display the last 1000 bytes of the file file.txt tail -c 1000 file.txt

tail

This command will display all lines in file.txt to the fifth line.

tail -n +5 file.txt



sort

The sort command is used to sort lines of text files in alphabetical or numerical order. By default, it sorts lines in ascending order.

This command will sort the lines in file.txt in alphabetical order and print the sorted lines to the terminal.

sort file.txt

This command will sort the lines in file.txt in reverse alphabetical order and print the sorted lines to the terminal.

sort -r file.txt

This command will sort the lines in numbers.txt in numerical order and print the sorted lines to the terminal.

sort -n numbers.txt

sort

This command will sort the lines in numbers.txt in reverse numerical order and print the sorted lines to the terminal.

sort -nr numbers.txt

This command will sort the lines in file.txt in alphabetical order and remove any duplicates, printing only the unique lines to the terminal.

sort -u file.txt

uniq

The uniq command is used to identify and remove duplicates from a sorted text file or input stream. By default, it compares adjacent lines in the input and removes any duplicates.

This command will read file.txt and remove any duplicate lines, printing only the unique lines to the terminal. uniq file.txt

This command will read file.txt and count the number of occurrences of each unique line, printing the count and the line to the terminal.

uniq -c file.txt

This command will read file.txt and print only the lines that appear more than once (i.e., the duplicate lines). uniq -d file.txt

uniq

This command will read file.txt and print only the lines that appear exactly once (i.e., the unique lines).

uniq -u file.txt

sort file.txt | uniq



The tr command in Linux is used to translate or delete characters. It can be used to replace or remove characters from a text file or input stream.

This command will replace all occurrences of the letter "o" with the number "O" in the input string "hello world", and print the result to the terminal:

echo "hello world" | tr "o" "0"

This command will read the contents of file.txt and convert all lowercase letters to uppercase letters using the [:lower:] and [:upper:] character classes, and then print the result to the terminal.

cat file.txt | tr "[:lower:]" "[:upper:]"

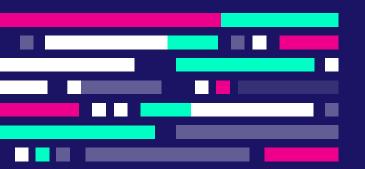


This command will read the contents of file.txt and remove all punctuation marks (such as periods, commas, and semicolons) using the [:punct:] character class, and then print the result to the terminal.

cat file.txt | tr -d "[:punct:]"

This command will replace all consecutive spaces in the input string "hello world" with a single space using the -s option, and print the result to the terminal:

echo "hello world" | tr -s " "



This command will search the file file.txt for any lines that contain the word "hello", and print those lines to the terminal.

grep "hello" file.txt

This command will search the file file.txt for any lines that contain the word "hello", regardless of the case (since we're using the -i option), and print those lines to the terminal. grep -i "hello" file.txt

This command will search recursively through the directory directory/ for any files that contain the word "hello", and print the names of those files to the terminal.

grep -r "hello" directory/

This command will search the file file.txt for any lines that begin with the word "hello" (since we're using the ^ character to anchor the search to the beginning of the line), and print those lines to the terminal.

grep "hello" file.txt

This command will list all running processes using the ps command, and then filter the results to show only the ones that contain the word "chrome" (in this case, we're using grep to search the output of the ps command). ps -ef | grep "chrome"

This command will print all lines in file.txt that do not contain the word "hello".

grep -v "hello" file.txt

This command will search all files in the current directory for the word "hello", and print the names of the files that contain it.

grep -l "hello" *

This command will print the number of lines in file.txt that contain the word "hello".

grep -c "hello" file.txt

This command will print all lines in file.txt that contain the whole word "hello", but not lines that contain partial matches like "hello there" or "helloworld".

grep -w "hello" file.txt

-E This option allows you to use extended regular expressions:

grep -E "hellolworld" file.txt

This command will print all lines in file.txt that contain either the word "hello" or the word "world", using the I character to separate the two options.

grep -A 1 "lazy" file.txt

the -A 1 option tells grep to print one line after each match of the pattern "lazy"



cut

The cut command in Linux is used to extract specific columns or fields from a file or input stream. It can be used to split a file into parts based on a delimiter or a fixed width.

cut is used to extract the first five characters from each line of the file.txt.

cut -c 1-5 file.txt

cut is used to extract the first and third fields from each line of the file.csv, which are separated by commas. cut -d , -f 1,3 file.csv

sed

stands for "stream editor" and is a powerful tool for text manipulation in Linux and Unix-based systems. It can be used to perform a wide variety of operations on text files and streams, such as searching, replacing, inserting, and deleting text.

Here, s is the sed command to search and replace text, and test and sample are the search string and replacement string, respectively.

sed 's/test/sample/' example.txt

Here, 2d is the sed command to delete the second line. sed '2d' example.txt

sed

sed '1a This is a new line.' example.txt

Here, 1a is the sed command to append text after the first line, and the backslash and newline are used to separate the command from the text to insert.



awk

Awk is a powerful text processing tool that is commonly used in Unix and Linux environments. It allows you to search for and manipulate patterns in text data, making it a useful tool for a wide range of tasks.

awk options 'pattern { action }' filename

awk -F',' '{ print \$1 }' data.txt

Here, the -F option specifies the field separator as a comma. The '{ print \$1 }' action tells awk to print the first field (i.e., the fruit names) for each line of input.

awk -F'[.,]' '{print \$1}' prices.txt

This command uses a regular expression to split each line of the prices.txt file into fields using either a . or , as the field separator.



awk

awk -F':' '{print \$1,\$7}' /etc/passwd

This command prints the first and seventh fields (username anda shell) of each line in the /etc/passwd file, using : as the field separator and a space as the output field separator.



tee

The tee command is used to read the standard input and then write it to both the standard output and one or more files.

ls -l | tee file.txt

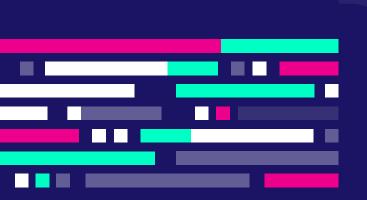
This command will write the output of the ls -l command to the file file.txt and also display it on the terminal.

echo "New line" | tee -a file.txt

This command will append the string "New line" to the end of the file file.txt.

echo "Hello world" | tee file1.txt file2.txt

This command will write the string "Hello world" to both file1.txt and file2.txt.



tee



cat bigfile.txt | tee -i output.txt

This command will read the contents of bigfile.txt and write them to output.txt, ignoring any interrupt signals that may be sent to the tee command.

-i or --ignore-interrupts: This option tells tee to ignore interrupt signals like CTRL+C, allowing it to continue running even if the user tries to stop it.

The find command is a powerful tool for searching for files in a directory hierarchy.

Find all files in the current directory and its subdirectories: find .

Find all directories in the current directory and its subdirectories:
find . -type d

that have the extension ".txt":

find . -name "*.txt"

find . -iname "FILE.txt" Similar to -name, but performs

Find all files in the current directory and its subdirectories

find . -iname "FILE.txt" Similar to -name, but performs a case-insensitive search.

Find all files in the current directory and its subdirectories that were modified more than 30 days ago:
find . -mtime +30

Find all files in the current directory and its subdirectories that are larger than 1 megabyte:

find . -size +1M

Find all files in the current directory and its subdirectories that have not been accessed in the last 7 days:

find . -atime +7

Find all files in the current directory and its subdirectories that are owned by the user "john":

find . -user john

Find all files in the current directory and its subdirectories that are not owned by the current user:

find . ! -user \$(whoami)

Find all files in the current directory and its subdirectories that are executable by the owner:

find . -perm /u+x

Find all files in the current directory and its subdirectories that are writable by anyone:

find . -perm /o+w

find . -mindepth 1 -type f

This will list all regular files (not directories or symbolic links) in the current directory and its subdirectories.

find . -mindepth 2 -type f

This will list all regular files **not** in the current directory but **in** its subdirectories.

find . -maxdepth 1 -type f

This will list all regular files (not directories or symbolic links) in the current directory and its subdirectories.

find . -maxdepth 2 -type f

This will list all regular files in the current directory and its subdirectories.

find . -name "*.txt" -exec cat {} \;

This command will find all files in the directory that end with .txt and execute the cat command on each of them.

find . -type f -name "*.txt" -print

This command will find all files in the directory that end with .txt and print their names to the terminal.



04

Hands on lab Practice





Subscribe

To stay updated with latest videos

Share

To let the others know more

hanks