Viewing files

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1. Outcomes

By the end of this document, you should be able to:

- 1. Open any kind of file using the command line, using cat, less, head, and tail
- 2. Know what a text editor is, and how to use one, for example using nano and vim

2. Introduction

At some point, you will want to view the files you have created. Luckily, many file types can be viewed directly in the terminal. This is useful, as it allows you to quickly check the contents of the file, and modify accordingly. Other file formats are not easily readable, and require specialised software to view and edit them - for example binary or compressed files.

3. Using cat

cat is a command that reads data from a file and outputs it to the terminal. It is a very simple command, and is often used to view the contents of a file.

Text redirection

The > symbol is used to take whatever is on the left and redirect the output. There's a whole tutorial on this, so don't be worried if you haven't come across it yet.

Shell 1: Create a file called hello.txt, and view the contents of the file using cat

```
# Create a file called hello.txt
echo "Hello, world!" > hello.txt
# View the contents of hello.txt
cat hello.txt
```

cat can be used to view the contents of multiple files at the same time. For example:

Shell 2: Create two files called hello.txt and world.txt, and view the contents of both files using cat

```
# Create a file called hello.txt
echo "Hello, world!" > hello.txt
# Create a file called world.txt
echo "World, hello!" > world.txt
# View the contents of hello.txt and world.txt
cat hello.txt world.txt
```

If you have very large files though, cat will dump the entire contents of the file to the terminal. This is not always desirable, as it can be difficult to read. Instead, you can use less, head, or tail.

4. Using less

less is a command that allows you to view the contents of a file, one page at a time. This is useful for viewing large files, as you can scroll through the contents of the file, and search for specific text.

The 'seq' command

seq is an in built BASH command which print out sequences of numbers. In it's default form it will print 0 to whatever number you specify, on separate lines.

Shell 3: Create a file called large.txt, and view the contents of the file using less

```
# Create a file called large.txt
seq 1000 > large.txt
# View the contents of large.txt using less
less large.txt
```

You can scroll through the file using the arrow keys, and search for specific text by typing $\ /$ followed by the text you want to search for Press n to go to the next occurrence of the text. Press q to exit less.

5. Using head and tail

head and tail are commands that allow you to view the first or last few lines of a file, respectively. This is useful for quickly checking the contents of a file, without having to view the entire file. or example, to view the first 10 lines of a file:

Shell 4: Create a file called numbers.txt, and view the first 10 lines of the file using head

```
# Create a file called numbers.txt
seq 100 > numbers.txt
# View the first 10 lines of numbers.txt using head with the -n flag
head -n 10 numbers.txt
```

Similarly, you can view the last 10 lines of a file using tail:

Shell 5: View the last 10 lines of the file numbers.txt using tail

```
# View the last 10 lines of numbers.txt using tail with the -n flag
tail -n 10 numbers.txt
```

6. Using a text editor

It is possible to edit files using the terminal, using a text editor. There are many text editors available, but two of the most commonly used are nano and vim. nano is a simple text editor that is easy to use, whereas vim is a more complex text editor that has a steeper learning curve.

To edit a file using nano, simply type nano followed by the name of the file you want to edit:

Shell 6: Edit the file hello.txt using nano

```
# Edit the file hello.txt using nano
nano hello.txt
```

Now you can edit the contents of the file. To save the changes and exit nano, press Ctrl + X, then Y, and finally Enter.

The very basics of vim are as follows. To enter vim, type vim followed by the name of the file you want to edit (or the name of a file if it exists already):

Shell 7: Edit the file hello.txt using vim

```
# Edit the file hello.txt using vim
vim hello.txt
```

In <code>vim</code>, you can move the cursor using the arrow keys. To enter insert mode, press <code>i</code>. Now you can edit the contents of the file. To save the changes and exit insert mode, press <code>Esc</code>, then <code>:wq</code> (i.e. write and then quit), and finally <code>Enter</code>. If you want to exit <code>vim</code> without saving the changes, press <code>Esc</code>, then <code>:q!</code>, and finally <code>Enter</code>. This is only the start, <code>vim</code> has many so-called 'key bindings' that allow you to do things more quickly than you would be able to in <code>nano</code>. But these are beyond the scope of this document.