title: The Unix Shell subtitle: Pipes and Filters

minutes: 20

The Unix Shell: Pipes and Filters

Learning Objectives

- Redirect a command's output to a file.
- Process a file instead of keyboard input using redirection.
- Construct command pipelines with two or more stages.
- Explain what usually happens if a program or pipeline isn't given any input to process.
- Explain Unix's "small pieces, loosely joined" philosophy.

Now that we know a few basic commands, we can finally look at the shell's most powerful feature: the ease with which it lets us combine existing programs in new ways.

Counting

We'll start with a directory called data/articles that contains text files of news articles. They are organized by the region they cover. So africa1.txt is the first article about Africa, africa2.txt the second and so on.

```
$ ls ~/programming-fundamentals/data/articles
```

Wow, that's a lot of files!

Let's go into that directory with cd and run the command wc africa*.txt. wc is the "word count" command: it counts the number of lines, words, and characters in files. Remember that the * in africa*.txt matches zero or more characters, so the shell turns africa*.txt into a complete list of .txt files that start with africa:

```
$ cd ~/programming-fundamentals/data/articles
$ wc africa*.txt
```

```
1
       101
               608 africa95.txt
              9632 africa96.txt
 1
      1576
              3218 africa97.txt
 1
       515
              4177 africa98.txt
 1
       653
       597
 1
              3748 africa99.txt
136
     70982 442767 total
```

If we run wc -1 instead of just wc, the output shows only the number of lines per file:

```
$ wc -l africa*.txt
```

```
1 africa94.txt
1 africa95.txt
1 africa96.txt
1 africa97.txt
1 africa98.txt
1 africa98.txt
1 africa99.txt
```

So we see that each article has only 1 lines. How can this be, if the number of words vary so widely? Open a file and try to guess why.

We can also use -w to get only the number of words, or -c to get only the number of characters.

Which of these files is shortest? It's an easy question to answer when there are only a few files, but what if there were 6000?

Redirecting and Printing

Our first step toward a solution is to run the command:

```
$ wc -w africa*.txt > lengths
```

The > tells the shell to **redirect** the command's output to a file instead of printing it to the screen. The shell will create the file if it doesn't exist, or overwrite the contents of that file if it does.

(This is why there is no screen output: everything that wc would have printed has gone into the file lengths instead.) ls lengths confirms that the file exists:

```
$ ls lengths

lengths
```

We can now send the content of lengths to the screen using cat lengths. cat stands for "concatenate": it prints the contents of files one after another. There's only one file in this case, so cat just shows us what it contains:

```
$ cat lengths
```

```
101 africa95.txt
1576 africa96.txt
515 africa97.txt
653 africa98.txt
597 africa99.txt
591 africa9.txt
```

Sorting

Now let's use the sort command to sort its contents. We will also use the -n flag to specify that the sort is numerical instead of alphabetical. This does *not* change the file; instead, it sends the sorted result to the screen:

```
$ sort -n lengths
```

```
1143 africa134.txt
1155 africa43.txt
1227 africa25.txt
1329 africa24.txt
1576 africa96.txt
```

We can put the sorted list of lines in another temporary file called sorted-lengths by putting sorted-lengths after the command, just as we used lengths to put the output of wc into lengths. Once we've done that,

we can run another command called head to get the first few lines in sorted-lengths:

```
$ sort -n lengths > sorted-lengths
$ head -1 sorted-lengths
```

```
70 africa49.txt
```

Using the parameter -1 with head tells it that we only want the first line of the file; -20 would get the first 20, and so on. Since sorted-lengths contains the lengths of our files ordered from least to greatest, the output of head must be the file with the fewest lines.

Pipes

If you think this is confusing, you're in good company: even once you understand what wc, sort, and head do, all those intermediate files make it hard to follow what's going on. We can make it easier to understand by running sort and head together:

```
$ sort -n lengths | head -1
70 africa49.txt
```

The vertical bar between the two commands is called a **pipe**. It tells the shell that we want to use the output of the command on the left as the input to the command on the right. The computer might create a temporary file if it needs to, or copy data from one program to the other in memory, or something else entirely; we don't have to know or care.

We can use another pipe to send the output of wc directly to sort, which then sends its output to head:

```
$ wc -w africa*.txt | sort -n | head -1

70 africa49.txt
```

This is exactly like a mathematician nesting functions like log(3x) and saying "the log of three times x". In our case, the calculation is "head of sort of word count of africa*.txt".

We can use this logic in many different combinations. For instance, to see how many files are in this directory, we can command:

```
$ ls -1 | wc -l 958
```

This uses wc to do a count of the number of lines (-1) in the output of ls -1.

Processes, Inputs, Outputs, and Filters

Here's what actually happens behind the scenes when we create a pipe. When a computer runs a program --- any program --- it creates a **process** in memory to hold the program's software and its current state. Every process has an input channel called **standard input**. (By this point, you may be surprised that the name is so memorable, but don't worry: most Unix programmers call it "stdin". Every process also has a default output channel called **standard output** (or "stdout").

The shell is actually just another program. Under normal circumstances, whatever we type on the keyboard is sent to the shell on its standard input, and whatever it produces on standard output is displayed on our screen. When we tell the shell to run a program, it creates a new process and temporarily sends whatever we type on our keyboard to that process's standard input, and whatever the process sends to standard output to the screen.

Here's what happens when we run wc -w africa*.txt > lengths. The shell starts by telling the computer to create a new process to run the wc program. Since we've provided some filenames as parameters, wc reads from them instead of from standard input. And since we've used > to redirect output to a file, the shell connects the process's standard output to that file.

If we run wc -w africa*.txt | sort -n instead, the shell creates two processes (one for each process in the pipe) so that wc and sort run simultaneously. The standard output of wc is fed directly to the standard input of sort; since there's no redirection with >, sort's output goes to the screen. And if we run wc -w africa*.txt | sort -n | head -1, we get three processes with data flowing from the files, through wc to sort, and from sort through head to the screen.

This simple idea is why Unix has been so successful. Instead of creating enormous programs that try to do many different things, Unix programmers focus on creating lots of simple tools that each do one job well, and that work well with each other. This programming model is called "pipes and filters". We've already seen pipes; a **filter** is a program like wc or sort that transforms a stream of input into a stream of output. Almost all of the standard Unix tools can work this way: unless told to do otherwise, they read from standard input, do something with what they've read, and write to standard output.

The key is that any program that reads lines of text from standard input and writes lines of text to standard output can be combined with every other program that behaves this way as well. You can *and should* write your programs this way so that you and other people can put those programs into pipes to multiply their power.

Redirecting Input

As well as using > to redirect a program's output, we can use < to redirect its input, i.e., to read from a file instead of from standard input. For example, instead of writing wc africa1.txt, we could write wc < africa1.txt. In the first case, wc gets a command line parameter telling it what file to open. In the second, wc doesn't have any command line parameters, so it reads from standard input, but we have told the shell to send the contents of africa1.txt to wc's standard input.

Rochelle's Pipeline: Concatenating Files.

Rochelle has her bulk text downloads in the new-york-times/2015-01-01 directory described earlier.

As a quick sanity check, she types:

```
$ cd ~/programming-fundamentals/data/new-york-times/2015-01-01
$ wc -l *.TXT
```

```
63661 human-rights-2000.TXT
56035 human-rights-2001.TXT
60045 human-rights-2002.TXT
46873 human-rights-2003.TXT
62611 human-rights-2004.TXT
55557 human-rights-2005.TXT
62905 human-rights-2006.TXT
44866 human-rights-2007.TXT
54580 human-rights-2008.TXT
66540 human-rights-2009.TXT
573673 total
```

Now she wants to concatenate (or merge) all of these text files into one big text file that she can later use to parse into a CSV.

```
$ cat *.TXT > all.TXT
$ wc -l *.TXT
```

```
573673 all.TXT
63661 human-rights-2000.TXT
56035 human-rights-2001.TXT
60045 human-rights-2002.TXT
46873 human-rights-2003.TXT
62611 human-rights-2004.TXT
55557 human-rights-2005.TXT
62905 human-rights-2006.TXT
44866 human-rights-2007.TXT
54580 human-rights-2008.TXT
66540 human-rights-2009.TXT
```

Notice that all.TXT is the sum of all the lines of the other *.TXT files.

Exercises

Challenge 1

Create a file under my_files called "sorts.txt" that contains these data



Sort it numerically and place the output in a file called "nsorts.txt". Then, sort it alphabetically and place the output in a file called "asorts.txt". Are these files the same? Why or why not?

Challenge 2

How would you count all of the words contained in articles about asia found in data/articles/? Store this number under my_files/ in a file called "asia_count"

Challenge 3

The command uniq removes adjacent duplicated lines from its input. For example, if a file salmon.txt contains:

coho
coho
steelhead
coho
steelhead
steelhead

then uniq salmon.txt produces:

coho steelhead coho steelhead

You have a file called data/animals.txt contains the following data:

deer
rabbit
raccoon
rabbit
deer
fox
rabbit
bear

Process this list such that there is only one value for each animal, and save it in a file under my_files/ called "unique_animals.txt"

When you feel you have met these challenges successfully, cd into test/ and type

. 1-3_test.sh

Adapted from: <u>Software Carpentry (http://software-carpentry.org/v5/novice/shell/03-pipefilter.html)</u>