Lab 1: basic UNIX tools

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This lab is available as a web page² or pdf document³.

Preamble: how to do the labs for this course

This course has a biweekly 1.5-hour lab session where you will work through the instructions and questions either from the web page or the PDF version³, which you can print ahead if you want.

You should work **with a partner**, sharing a single keyboard and screen. This will help you meet other students and learn more, since your partner may have questions or comments you hadn't thought of, and you'll learn to communicate about course concepts, which is important for your understanding (and for the exam!)

For this to work well, each person should ensure that both they *and* their partner understand what is going on. Pay attention to what your partner is doing and discuss what's going on. That means:

- If you know the answer and your partner doesn't, *don't* just type it in and move on to the next question---explain what you have done and why. This helps your partner learn *and* improves your own understanding. Ask if your partner is satisfied with your explanation. Can you think of other examples to help your partner understand? Explaining technical concepts to other people is an important skill, and this will help you practice it.
- If your partner knows the answer and you don't, don't let them move on until you understand too. Don't let them dictate what you should type if you don't understand why. Do ask questions, and tell your partner if their explanation makes sense or not. Remember, just because they think they know the answer doesn't necessarily mean they are right! If it doesn't make sense to you, they need to work harder at explaining, and maybe discover that they are wrong!
- If neither of you knows the answer, ask one of the demonstrators in the lab, and we can try to help you.
- If one person is much more familiar with Python than the other, try putting the less experienced person at the keyboard or at least switching frequently, so they will get more practice with basic coding skills.

¹http://creativecommons.org/licenses/by-nc/4.0/.

²http://www.inf.ed.ac.uk/teaching/courses/anlp/labs/lab1.html

³http://www.inf.ed.ac.uk/teaching/courses/anlp/labs/lab1.pdf

⁴http://computing.help.inf.ed.ac.uk/new-taught-students

⁵http://computing.help.inf.ed.ac.uk/sites/default/files/IntroLecture201617.pdf

⁶https://web.stanford.edu/class/cs124/kwc-unix-for-poets.pdf

⁷http://www.inf.ed.ac.uk/teaching/courses/anlp/labs/lab2.html

You should be able to work through the main set of questions in 1.5 hours. However, we encourage you to go beyond the basic set of questions to further improve your practical skills and understanding, using the "Going Further" section at the end of each lab.

NOTE: This week's lab has a lot to read. Future labs will have less reading and more thinking/coding, but you need to read through some basics first.

Preliminaries: to do before the first lab session

Please do the following **before** arriving at the first lab session.

- 1. Get a DICE (Distributed Informatics Computing Environment) account. All students on an Informatics degree should already have an account. If you are on a degree in another school, you will automatically get a DICE account once you have registered for this class, so please register as soon as possible. In the meantime, please pair up with another student who already has an account and do the lab together using their account.
- 2. Read the School's Computing Support information page for new taught students⁴ and introduction to DICE⁵. The rest of the lab assumes you have read sections 1-4 and 8 (at least up to "other things to consider") of the introduction to DICE.

Goals and motivation of this lab

This lab aims to:

- familiarize you with basic command line tools in the UNIX operating system.
- give you a first taste of working with text data.

What is UNIX?

This operating system is commonly used by computing researchers and programmers; in Informatics we use a version called Scientific Linux, which is installed on all of the student lab machines and which you will be using for the practical labs in this course. (The UNIX tools described here are also available in Apple's OS X, and in Windows by installing additional software such as Cygwin.)

UNIX commands can be a bit confusing at first but are powerful tools for manipulating files and data, including exploratory data analysis. This lab will introduce some basic UNIX commands for exploring and processing files. You can do a lot of useful stuff with just a few commands, and we hope you'll keep using the skills from this lab for the rest of the semester and in your other courses.

But I already know UNIX!

This course is taken by students with many different backgrounds. If you are already familiar with UNIX, please skim through anyway to make sure you know all the commands we discuss and can do the tasks at the end of the lab. You may also want to take a look at the Going Further section for more practice, and you can also help out your classmates who have less experience than you.

What data will we use and what will we do with it?

We will explore the data from a corpus of parent-child interactions, building up to computing the child's *mean length of utterance* (MLU) in one or two files. MLU measures a child's language development: it is just the average number of words (or, sometimes, morphemes) in each of the child's utterances (spoken sentences).

Lab Setup

To set up for the labs, log into your DICE account and open a *shell* (also called a *terminal*) following the instructions in the Introduction to DICE document. You should see a *command prompt* of the form:

[hostname]username:

Now, create a directory (folder) called anlp that you will use for all your ANLP work. You can do this by entering the following command at the prompt. (*Remember*, mkdir stands for 'make directory')

```
mkdir anlp
```

Now, create a subdirectory for your labs, and then another one for this lab. (You cannot create a subdirectory before creating the directory it belongs inside.)

```
mkdir anlp/labs
mkdir anlp/labs/lab1
```

Downloading the data

In this lab we will be working with data from CHILDES, the Child Language Data Exchange System. CHILDES is a collection of many different corpora in many different languages, all contributed by researchers of child language development. Different researchers have different interests, so the corpora contain different types of data (transcriptions, audio, and/or video) and different kinds of annotations (ranging from detailed phonetic transcripts to morphological and syntactic annotations).

We will use the Providence corpus for this lab. To download the corpus, click on the following URL or paste it into your web browser: http://homepages.inf.ed.ac.uk/sgwater/teaching/Providence.zip. You should get a dialog box asking what to do with the file; choose Open with Archive Manager (default). In the next window, click Extract and then navigate to your lab1 directory by double-clicking in the right-hand pane on the anlp folder, then lab1. Then click Extract.

ls

Go back to your terminal window and cd (meaning change directory) into the lab1 directory:

```
cd anlp/labs/lab1
```

If you ever get confused or forget which directory you are in, use the pwd (print working directory) command. Try it now:

pwd

Now let's *list* the contents of the current directory using ls:

ls

You should now see a subdirectory called Providence, which contains the data you just downloaded. What do you see in it? What about in the further subdirectories?

```
ls Providence
ls Providence/Ethan
```

Looking at the data

less

To get an idea of what is in the files you just downloaded, type:

cd Providence/Ethan
less eth01.cha

What information is in the metadata at the top of each file? (Hint: child language researchers use the format y;m.d to indicate a child's age in years;months.days)

(less may seem like a funny name for this command which shows you *more* of the file; it's because there was an earlier similar command called more, so when this newer version was developed the developers decided to give it a cute name. UNIX developers like cute names.)

When you are looking at a file using less, you can scroll up and down using the arrow keys or <PageUp>/<PageDown>. <space> also acts like <PageDown> and <Enter> acts like <down-arrow>.

What do you see in the rest of the file? (Hint: the string of numbers at the end of each line is a code that links to a time point in the audio recording of this data. The audio isn't included here but can be obtained from the CHILDES database.)

One final useful thing you can do with less is search for a particular string in the file by typing / followed by the string. Try it: type

/the

You should see all of the instances of the being highlighted.

To stop viewing the file and go back to the terminal command line, type q.

Actually, the eth01. cha file maybe is not so interesting. Look now at eth50. cha. What are some of the main differences between the data in these two files? Can you think of an explanation for those differences? (If you want to look at both files simultaneously, you could start a new terminal and open one file in each terminal.)

head, tail

Sometimes we just need a quick peek at part of a file. What does this command do?

head eth01.cha

What about this one?

tail eth01.cha

These commands can be useful, for example, if we want to look over the range of ages in the files we have. Try:

head *.cha

What does the * do?

man

We don't really need to look at the first 10 lines of each file to see the ages, we only need the first five lines. If we could print only the first five lines, we could see the information we want more compactly. Most UNIX commands, including head, have many possible options to change their behavior. To see what options are available, look at the *manual* for the command:

man head

You should see:

NAME

head - output the first part of files

SYNOPSIS

```
head [OPTION]... [FILE]...
```

DESCRIPTION

Print the first 10 lines of each FILE to standard output. With more than one FILE, precede each with a header giving the file name. With no FILE, or when FILE is -, read standard input.

Mandatory arguments to long options are mandatory for short options too.

-n, --lines=[-]K
 print the first K lines instead of the first 10; with the leading '-', print all but the last K lines of each file

-v, --verbose
 always print headers giving file names

Note: you should be able to move up and down in the man pages using the same keys you used for less [I hope; the default for the student account might not allow you to move upwards...], and you can also return to the command line using q.

The man page starts with the name of the command and a brief synopsis of how to use it. In this case, it says that the head command can be followed by zero or more *options* (sometimes called *flags*) and then zero or more *files* as arguments (things to act on). The square brackets indicate that these options and files need not be included at all, and the . . . indicate that you can include more than one of each.

The description after the synopsis says what will happen if you include zero or multiple files. (*Standard input* refers to the text you input in the terminal. Try entering head with no filename to see how this works. You will need to enter some more text after that! To quit, type Ctrl-c.)

The next part of the description tells you what the possible options are and if they require arguments themselves. Many (in this case all) options have both a short and a long form, which are equivalent. For example, to print just the first 5 lines of each file, we could either use:

```
head -n5 eth01.cha
```

or:

```
head --lines=5 eth01.cha
```

(To try this yourself, you will need to either use another terminal or quit the man page first by typing q.)

The K specified in the description is a variable indicating a required argument to the -n option. Here we use the value 5 for K. Notice that this option also has a non-required argument [-] so if we wanted to print all but the last 5 lines we could type:

```
head -n-5 eth01.cha
```

As noted above, you can specify multiple options at once. Compare the output of the following commands:

```
head -n5 *.cha
head -n5 -g *.cha
```

Now that you know how to read a man page, you may want to look at the man pages for some of the other commands we've seen to see what options are available for them.

There is also a *lot* of help about these commands on the Internet. You can find more examples or maybe a better explanation than the man page has. If you know the name of the command you want to use, like grep, try searching for UNIX command grep. If you know what you want to do but not the name of the command, searching is trickier, but try searches like UNIX search regular expression or even UNIX find string in file. Get in the habit of looking for help--you will start to recognize some of the main help forums and which ones are most useful to you.

Being lazy

As you may have noticed, most UNIX commands are very short to avoid lots of typing. You can avoid even more typing by using *tab completion* and *history*.

Tab completion

Type the following into your terminal. Instead of pressing <enter> at the end, press the <tab> key:

```
head e
```

You should find that the filename has been automatically extended to eth. If you type 01 and then <tab> again, you will have the complete filename. In this case the tabbing didn't save many keystrokes, but for long filenames it can save a lot. You can use tab completion for nearly anything in UNIX (commands, filenames, directories). The operating system will complete as much as it can but will stop where there are multiple possible extensions (like here, where all files in the current directory start with eth but we then had to type in 01 to disambiguate before tabbing again.) In DICE, tab completion is set up so that if you hit <tab> twice when there are multiple completions, the system will print out all possible completions for you to see.

History

Your terminal keeps track of all the commands you have entered. To access the previous command, just type the up-arrow key. If you hit <up> twice, you will get the command before that, and so forth. This can save a lot of typing.

Finding and counting things in files

wc

Try this command:

```
wc eth01.cha
```

What are the numbers that you got back? What do you think WC stands for? (Hint: look at the man page!)

Using a single command, can you figure out which of the first nine Ethan files has the most lines in it?

```
wc -I eth0*.cha
```

grep

Sometimes we want some more specific information. The number of lines in a file roughly indicates how much language data is in there, but what if we want to know how many utterances are spoken by a particular person? Try this command:

说话

```
grep 'MOT' eth01.cha
```

What did this command do?

Using the man page for grep, figure out what option you can use with grep to compute the number of utterances spoken by the mother in this file.

In case you are wondering, <code>grep</code> stands for *Globally search for a Regular Expression and Print*. We won't discuss regular expressions (REs) in detail here, but they are covered in Section 2.1 of the Jurafsky and Martin textbook, and also in the Computer Programming for Speech and Language Processing course. (The <code>grep</code> man page also explains the syntax that <code>grep</code> uses for REs.) REs are a powerful way to match patterns in text and we hope you will learn to love them. For now, we will just show you a few simple REs to show some of what you can do.

First, look at this command but don't run it yet:

```
grep 'MOT.*the' eth01.cha
```

What do you think it will do? (Hint: . matches any character, and * means "zero or more repetitions of the previous character")

To see if you were right, run the command and look at the first few lines of output. Even better, let's only print out the first few lines of output. We can do that by combining two commands we already know, grep and head:

```
grep 'MOT.*the' eth01.cha | head
```

pipes and redirection

What just happened?

The | (vertical bar on the keyboard, called *pipe* in UNIX-world) tells the operating system to use the *output* of the first command as the *input* to the second command. (Another way to think of this is that it turns the output of the first command into "standard input", which as you will recall, is what is used as input to head if no filename is given.

Can you think of a way to use pipe to combine two commands we already know to count the number of lines that include the string MOT without using the - C option for grep? grep 'MOT' eth01.cha | wc - | 648

Instead of sending the output of one command to the input of another, we can also just send the output of a command into a file. This can be useful if we want to save the output for later inspection, and we do it using the > (output redirect) character:

```
grep 'MOT.*the' eth01.cha > my_output.txt
less my_output.txt
```

grep again

Coming back to regular expressions, let's make sure you understand what the RE 'MOT.*the' is matching in the grep command. What is the difference between the following two commands?

```
grep 'MOT.*the' eth01.cha
grep 'MOT.*\bthe\b' eth01.cha
```

You may want to look at just the start of the output by piping it to head. Or try using the --color=auto option for grep (unfortunately the color is lost if you pipe the output).

In general, backslashes are used in regular expressions to indicate something special (like a word boundary or any whitespace character) and are also used to make characters that are *usually* special (like *) be interpreted literally. So, for example, if we want to match only the lines spoken by the mother, we should use '^*MOT' rather than just 'MOT', because the first version only matches the exact string *MOT at the beginning of a line (indicated by the ^). The second version also matches lines that are not spoken by the mother but happen to contain the string MOT. (Are there any?)

As you can start to see, regular expressions can be very powerful but proper use takes time and practice, and we will not go any further here.

Computing MLU

Believe it or not, we now have all the commands we need to be able to compute the child's mean length of utterance in any given file. Remember, <u>MLU</u> is the average number of words spoken by the child in each of the utterances in the file. What is Ethan's MLU in the file eth50.cha? Assume for this question that a word is any whitespace-delimited string of characters (including punctuation) in the transcription. So, for example, you should count fill him with pom+poms xxx . as six words.

Hint: you can't get the answer entirely automatically. Use the commands you know to get a few key numbers and then perform some basic arithmetic on them yourself. This works fine if we only care about the MLU for one or two files. If we wanted to compute MLU for all the files, we would probably want to write a short program to do it.

Going Further

If you want, try your hand at one of the following, in whatever order you like:

- 1. We showed how to get all the lines from a file that contain the word 'the'. But what if we actually want to count the number of occurrences of 'the'? Can you figure out how to do this using grep, possibly in combination with other commands we've seen? (*Hint*: look at other command line options to grep. If you aren't sure whether the output is what you want, make sure you test it! Creat a small file of your own where you know the right answer.)
- 2. Suppose you are only allowed to use the same options to grep that we used earlier in the lab. By reading the man page or searching the Web, figure out how the tr command works, and use it together with commands we've seen to count the number of occurrences of 'the' in a file.
- 3. We've covered a few useful commands here, but there are quite a few more that can come in handy for text processing and other NLP-related tasks. To learn more about those, you may want to go through the Unix for Poets⁶ tutorial.
- 4. Explore the data a bit more using Python, by going through Lab 2 from 2014⁷. (We won't use that lab this year because we have a tutorial instead, but that doesn't mean it isn't useful!)