EEB 174 Homework week 2

Preliminaries

Create a new repo in your <u>eeb-177</u> Desktop folder called <u>homework</u>. Follow the <u>instructions</u> from week 0 for creating a new git repo.

Create a file called week-2-hw-eeb174.txt in your homework directory.

Use this file for your homework answers.

Question 1.

Lets revisit the Pacifici2013_data.csv file from class today (in CSB/unix/data. Show the Order of the first 5 species in the data set.

(hint: you will need cut, I, tail, and head)

Question 2.

How many unique families are in this data set? Append the line "There are X unique families:" (fill in the value for x) to your exercise file. Then append the list of unique families to your hw file.

hint: you will need the uniq command. Also, you will to sort your data before you apply uniq as uniq only collapses non-unique values when they are contiguous.

Question 3.

Work through the Allesina example 1.7.3 that introduces the tr command and shows how the commands you have learned can be used to subset the Pacifici2013_data.csv. Append your command history for 1.7.3 only to your hw file.

Question 4.

What is the 10th largest species in the database? What is the 10th smallest?

Question 5.

Work through section 1.7.4 in the Allesina text on grep. Append your command history for 1.7.3 only to your hw file.

Question 6.

Do exercise 1.11.1 in Allesina.

hint 1: you will need to use the squeeze option (-s) to deal with repeat spaces for #5.

hint 2: #7 requires numeric sorting on the number of reads. How could you get a column of numreads only and then how could you cut the number from the text "numread"?

Question 7

Read the software carpentry section on loops in the shell here. Then do the following.

- create a directory called temp in CSB/unix/sandbox
- cd to the Saavedra2013 directory in CSB/unix/data

• write a loop that will copy all of the text files in this directory to your newly created temp directory. The copied files should all be renamed to include "edited-" in front of the original file name. Append this loop to your HW file.

hint: you will need to give the cp command the relative (or absolute) path to the target directory

hint use the echo command to see the new file name in the loop before you copy as in the *Nelle's Pipeline: Processing Files* section of the loops page.

Question 8

Work through the software carpentry exercise on finding things <u>here</u>. Answer the questions in the orange boxes at the end of this section in your HW file.

∕ Using grep	
find Pipeline Reading Comprehension	
✓ Matching and Subtracting	
Little Women 🖸	
Finding Files With Different Properties	

Commit all of your work as you go.

Due Dates

Homework must be committed by Monday, January 23rd at 5:00 PM.