EEB 177 Lecture 4

Topics

- Advanced shell commands
- Permissions
- Intro to Python

Office Hours

Tues, Wednesday 11-12

Hershey Courtyard (inside Hershey Hall)

Terasaki 2149 on rainy days

Hacking Sessions

This Thursday from 6-8 in LS 3209

Homework #2

Due this Friday by 5:00 PM

Preliminaries

- Start gedit: \$ gedit and save the file "classwork-Tuesday-1-24.txt" to your class-assignments directory
- push this to your remote repository
- you can write answers to today's exercises in this file.

wget

wget retrieves files from a web address. Lets try it! change to the /sandbox directory in CSB/unix then...

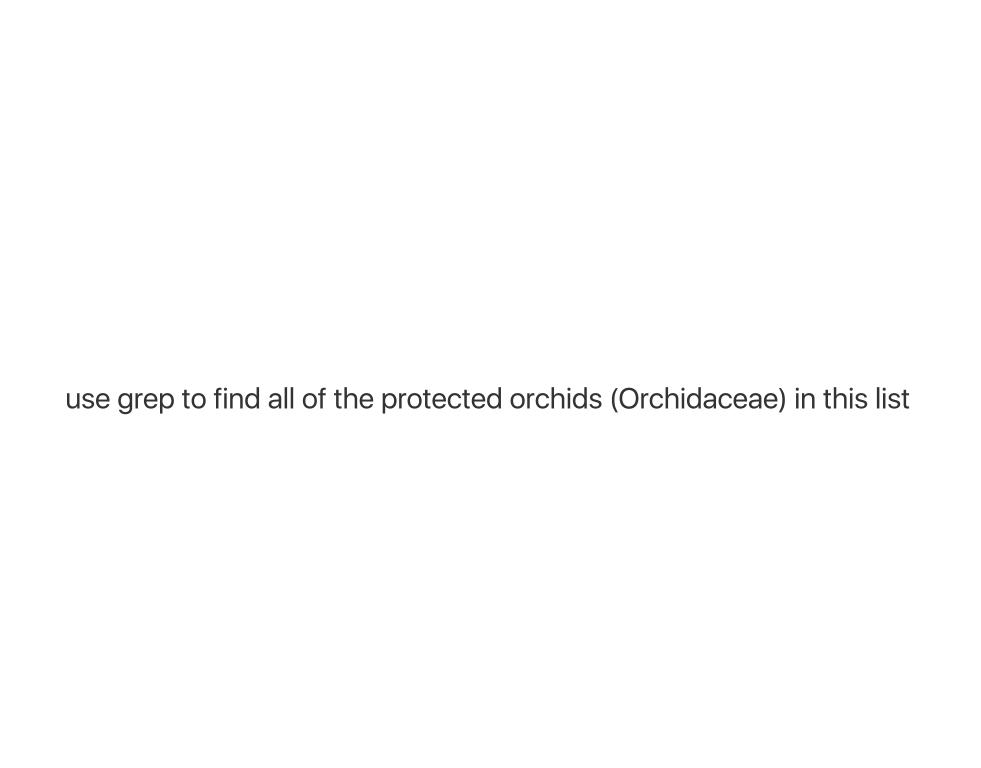
wget http://tinyurl.com/cep-taxa -0 cep-taxa.txt this retrieves the remote file and renames it.

grep review

. Grep is a powerful pattern matching command that can be combined with the regular expressions you used in lab.

Useful grep options:

- -c to count lines
- –w to match words
- -i to make case insensitive
- -n to show line number of match.



use grep to find all of the protected orchids (Orchidaceae) in this list

grep Orchidaceae cep-taxa.txt

how could you count these lines?

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how could you count these lines?

grep -c Orchidaceae cep-taxa.txt

use grep to find all of the falcons in this list.

try searching for falcon. Then try searching for Falcon. What is going on? How can you find occurrences regardless of case?

use grep to find all of the falcons in this list.

try searching for falcon. Then try searching for Falcon. What is going on? How can you find occurrences regardless of case?

grep -i falcon cep-taxa.txt

Macaws are in the genus Ara. Find all of the macaws in this list regardless of case. What is the problem now?

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How can you solve it?

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How can you solve it?

grep −i −w ara cep−taxa.txt

Other grep options

- -B X finds x lines before
- -A X finds X lines after

find the 3 lines before all occurrences of Cebus.

Other grep options

-n shows the line number of the match.

what are the line numbers of all iguanas in the file?

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-n shows the line number of the match.

what are the line numbers of all iguanas in the file?

grep −i −n −w iguana cep−taxa.txt

finding all lines that do not match

-v returns everything that does not match the grep pattern

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How many lines do not match ara?

what are the line numbers of those lines?

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How many lines do not match ara?

```
grep −i −v −c ara cep−taxa.txt
```

what are the line numbers of those lines?

```
grep -i -v -n ara cep-taxa.txt
```

#finding files with find

find allows you to search for files with specified attributes.

use the wildcard ** to find everything in your sandbox directory.

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find .

```
./temp.txt
./cep-taxa.txt
./.gitignore
./junk
```

if you pass find a path it will give all files and folders in that directory

```
find ../data/
```

```
../data
../data/toremove.txt
../data/Gesquiere2011_data.csv
../data/Saavedra2013_about.txt
...
```

if you pass find a path it will give all files and folders in that directory

```
find ../data/
```

```
../data
../data/toremove.txt
../data/Gesquiere2011_data.csv
../data/Saavedra2013_about.txt
...
```

count all of the files and folders within ../data/

find options

You can search for a specific file with -name

```
find ../data/ -name "n30.txt"
```

This can be helpful when you don't know exactly where you left a file.

```
find /home/eeb-177-student/Desktop/ -name "classwork-
Tues-1-17.txt"
```

/home/eeb-177-student/Desktop/eeb-177/class-assignments/c

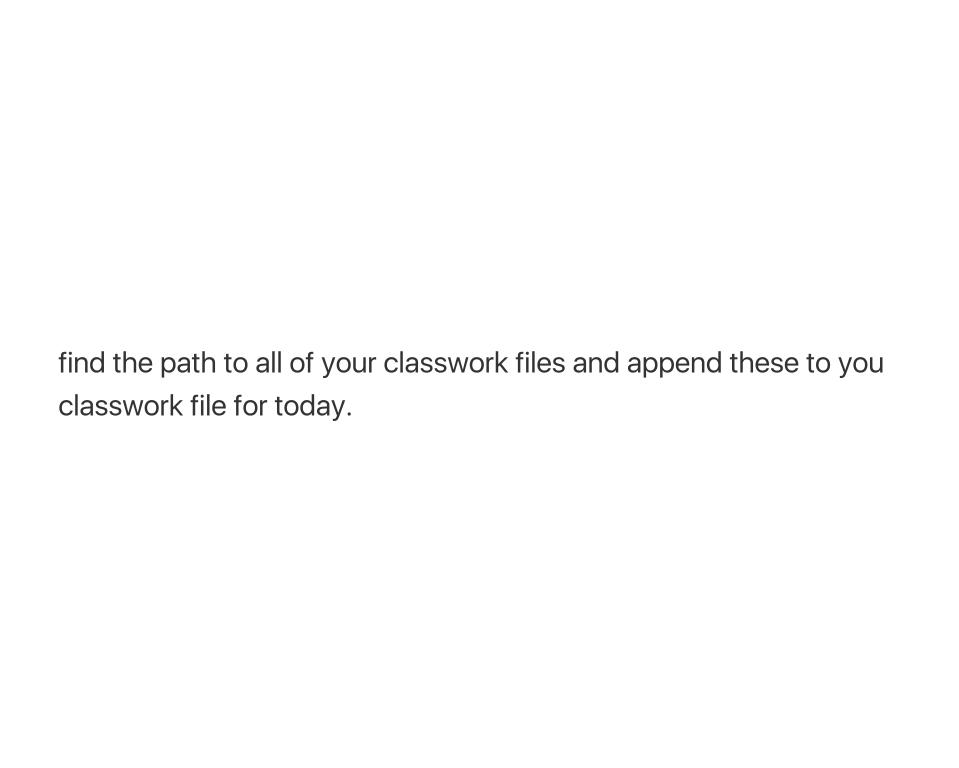
find gets even more powerful with wildcards.

for example, to find all of the files with about in the data directory....

```
find /home/eeb-177-student/Desktop/eeb-177/CSB/unix/ -
iname "*about*"
```

```
/home/eeb-177-student/Desktop/eeb-177/CSB/unix/data/Saaved/home/eeb-177-student/Desktop/eeb-177/CSB/unix/data/Marra/home/eeb-177-student/Desktop/eeb-177/CSB/unix/data/Pacif
```

note that -iname ignores the case in the file names



find the path to all of your classwork files and append these to you classwork file for today.

```
find /home/eeb-177-student/Desktop/ -iname "*class**" >>
/home/eeb-177-student/Desktop/eeb-177/class-
assignments/classwork-Tues-1-24.txt
```

specifying the depth of the search

to restrict the depth in the folder hierarchy of the search, use the maxdepth N option.

What will this line do?

```
$ find ../data -maxdepth 1 -name "*.txt" | wc -l
```

How many text files are there in ../data if you do not restrict the depth?

You can exclude certain files with not

find ../data/ -not -name "*about*" | wc -l

Permissions

In Unix, each file and directory has an attribute that determines who can read (r), write (w), execute (x), or do nothing (-) to a file. There are three categories of file users

- owner
- group (set of users)
- everyone else

you can see permissions with ls -l

permissions structure

```
Indicates a directory (d) or a file (–)

Group (set of users)

drwxrwxrwx

Other users (everyone else)

User (owner) permissions
```

chmod and chown

These commands change permissions and ownwership (u , g , or o for user, group or other).

```
touch permissions.txt
ls -l
```

```
-rw-rw-r-- 1 eeb-177-student eeb-177-student 0 Jan 24 07:
```

notice that the user may now execute this file.

you can also add and remove permissions for a user with + and -.

```
chmod g+w,u+x permissions.txt
ls -l permissions.txt
-rwxrw-rw-
```

Add write permissions for all users.

Remove read, write, and execute permission from others

text editor that will execute a series of shell commands that you have already learned.

open up gedit and type the following lines:

```
#! /bin/bash
ls -la
echo "Above are the directory listings for this folder"
pwd
echo "right now it is:
date
```

save this file as dir.sh

Paths

there are two standard locations for programs-- /usr/bin and /bin

use ls to see what is in them

Creating a scripts directory and adding it to the path

we want a single working copy of each program on our machines so we need to make sure the shell searches for our programs....

- go to your home directory
- create a directory called scripts
- to add the scripts directory to the path, open the
 bash_profile file in gedit
- add this line (exactly) export PATH="\$PATH:\$HOME/scripts"
- exit and save

Now we have created a program we would like to run and created a path to the scripts directory. What else do we need to do?

hint: where is dir.sh right now?

hint: what permissions do we need to execute a script?

the shebang (#!)

#! is called the shebang--it means that all following contents of script will be sent to the program following the shebang

#! /bin/bash sends it all to bash

remember, new scripts are not executable w/o changing permissions

checking permissions

- cd ~/scripts
- check permission with Is -I
- add permission to execute with chmod u+x

try running your program from different directories. Does it work? Why?	

exercise

Add your scripts directory to your remote repository. You will need to

- git init in your scripts directory
- add your script
- commit your script
- create a remote repo on github
- copy and paste the command lines from the remote repo after you create it.

```
git remote add origin https://github.com/michaelalfaro/eelgit push —u origin master
```

you will be using your scripts directory throughout the quarter, so make sure this repo is working

Programming languages

- There are over 2000
- There is no perfect language for all tasks
- You are already learning several: regular expressions, python, R
- This class does not cover fast, compiled languages like C. Useful for heavy computational tasks

Why Python?

- easy to teach
- readable
- powerful string manipulation, web scraping, and other capabilities
- helps enforce good programming practices

Getting started with python

Python should already be installed on your VM

```
$ ipython
IPython 2.2.0 -- An enhanced Interactive Python.
? -> Introduction and overview of IPython's featu
%quickref -> Quick reference.
help -> Python's own help system.
object? -> Details about 'object', use 'object??' for eximal import scipy
```

to exit ipython: CTRL-D CTRL-D to exit

IPython interactive mode

You can use python and IPython from the command prompt.

```
In [1]: 2 + 2
Out[1]: 4

In [6]: 2 > 3
Out[6]: False

In [11]: "I'm fine, " + "thank you"
Out[11]: "I'm fine, thank you"
```

try it out....

Python operators

```
+ Addition
- Subtraction
* Multiplication
/ Division
** Power
% Modulo
// Integer division == Equals
!= Differs
> Greater
>= Greater or equal &, and Logical and |, or Logical or
!, not Logical not
```

Variables

You typically manipulate variables in programming languages.

```
In [12]: x = 5
In [13]: x
Out[13]: 5
In [14]: x + 3
Out[14]: 8
In [15]: y = 8
In [16]: x + y
Out[16]: 13
In [17]: x = "My string"
In [18]: x + " is now longer"
Out[18]: "My string is now longer"
In [19]: x + y
TypeError: cannot concatenate "str" and "int" objects
```

Python recognizes variable types and methods for converting among them

```
In [20]: x
Out[20]: "My string"
In [21]: y
Out[21]: 8
In [22]: x + str(y)
Out[22]: "My string8"
In [23]: z = "88"
In [24]: x + z
Out[24]: "My string88"
In [25]: y + int(z)
Out[25]: 96
```

Collections

Python has variables that are collections of other objects. lists are collections of ordered values and are defined by [].

```
# Anything starting with # is a comment
In [26]: MyList = [3,2.44,"green",True]
In [27]: MyList[1]
Out[27]: 2.44
In [28]: MyList[0] # NOTE: FIRST ELEMENT -> 0
Out[28]: 3
In [29]: MyList[3]
Out[29]: True
In [30]: MyList[4]
IndexError: list index out of range
In [31]: MyList[2] = "blue"
In [32]: MyList
Out[32]: [3, 2.44, "blue", True]
```

note that indexing in python starts at 0.

{.smaller}

append, sort, and count are methods that work on lists

```
In [33]: MyList[0] = "blue"
In [34]: MyList
Out[34]: ["blue", 2.44, "blue", True]
In [35]: MyList.append("a new item")
In [36]: MyList
Out[36]: ["blue", 2.44, "blue", True, "a new item"]
In [37]: MyList.sort()
In [38]: MyList
Out[38]: [True, 2.44, "a new item", "blue", "blue"]
In [39]: MyList.count("blue")
Out[39]: 2
```

More list operations

Try these

```
In [71]: apes = ["Homo sapiens", "Pan troglodytes", "Gori
In [72]: #how long is list?
In [73]: len(apes)
Out[73]: 3
In [74]: #index elements with brackets
In [75]: apes[1] # what element will this be
Out[75]: 'Pan troglodytes'
In [76]: apes.reverse() #lists are ordered
In [77]: apes[0]
Out[77]: 'Gorilla gorilla'
#if you know the element but do not know position, use in
In [78]: apes.index("Homo sapiens")
Out[78]: 2
#get the last element of any list with [-1]
In [79]: apes[-1]
Out[79]: 'Homo sapiens'
```

Subsetting lists

You can make a new list by telling python the elements of the original list to include

```
ranks = ["kingdom","phylum", "class", "order", "family"]
In [81]: ranks
Out[81]: ['kingdom', 'phylum', 'class', 'order', 'family'
lower_ranks = ranks[2:5]
In [83]: lower_ranks
Out[83]: ['class', 'order', 'family']
```

Note how python indexing works:numbers are inclusive at the start and exclusive at the end.

Appending to lists

Try this

```
apes = ["Homo sapiens", "Pan troglodytes", "Gorilla goril
print("There are " + str(len(apes)) + " apes")
apes.append("Pan paniscus")
print("Now there are " + str(len(apes)) + " apes")
```

append() changes the variable in place!

+ and extend

Try this

```
apes = ["Homo sapiens", "Pan troglodytes", "Gorilla goril
monkeys = ["Papio ursinus", "Macaca mulatta"]
primates = apes + monkeys
print(str(len(apes)) + " apes")
print(str(len(monkeys)) + " monkeys")
print(str(len(primates)) + " primates")

platyrrhines = ["Cebus", "Sapajus", "Aotus"]
monkeys.extend(platyrrhines)
print monkeys
```

{.smaller}

Dictionaries are collections of key-value pairs. Very useful for data without a natural order. Defined by \{\}.

```
In [1]: GenomeSize={"Homo sapiens": 3200.0,
    "Escherichia coli": 4.6, "Arabidopsis thaliana": 157.0}
In [2]: GenomeSize
Out[2]:
{"Arabidopsis thaliana": 157.0,
    "Escherichia coli": 4.6,
    "Homo sapiens": 3200.0}
```

You can access the value of a dictionary by suppling the key for that pair.

```
In [3]: GenomeSize["Arabidopsis thaliana"]
Out[3]: 157.0
In [4]: GenomeSize["Saccharomyces cerevisiae"] = 12.1
In [5]: GenomeSize
Out[5]:
{"Arabidopsis thaliana": 157.0,
   "Escherichia coli": 4.6,
   "Homo sapiens": 3200.0, "Saccharomyces cerevisiae": 12.1
```

Adding keys and changing values in dictionaries

```
# ALREADY IN DICTIONARY!
In [6]: GenomeSize["Escherichia coli"] = 4.6
In [7]: GenomeSize
Out[7]:
{"Arabidopsis thaliana": 157.0,
 "Escherichia coli": 4.6,
 "Homo sapiens": 3200.0,
 "Saccharomyces cerevisiae": 12.1}
In [8]: GenomeSize["Homo sapiens"] = 3201.1
In [9]: GenomeSize
Out [9]:
{"Arabidopsis thaliana": 157.0,
 "Escherichia coli": 4.6,
 "Homo sapiens": 3201.1,
 "Saccharomyces cerevisiae": 12.1}
```

Creating dictionaries

You will often create a dictionary and then populate it. Try this!

```
enzymes = {}
enzymes['EcoRI'] = r'GAATTC' # r before the string tells |
enzymes['AvaII'] = r'GG(A|T)CC'
enzymes['BisI'] = r'GC[ATGC]GC'
enzymes.keys()
enzymes.values()
```

You can use zip() to turn two lists into a dictionary

```
keys = ('name', 'age', 'food')
values = ('Monty', 42, 'spam')
zip(keys, values) #makes a list of tuples
my_new_dict = dict(zip(keys, values))
my_new_dict
```

tuples contain sequences that are immutable and are defined by ().

```
In [12]: FoodWeb=[("a","b"),("a","c"),("b","c"),("c","c")
In [13]: FoodWeb[0]
Out[13]: ("a", "b")
In [14]: FoodWeb[0][0]
Out[14]: "a"
# Note that tuples are "immutable"
In [15]: FoodWeb[0][0] = "bbb"
TypeError: "tuple" object does not support item assignment in [16]: FoodWeb[0] = ("bbb","ccc")
In [17]: FoodWeb[0]
Out[17]: ("bbb", "ccc")
```

Use tuples when order matters.

sets are lists without duplicate elements

```
In [1]: a = [5,6,7,7,7,8,9,9]
In [2]: b = set(a)
In [3]: b
Out[3]: set([8, 9, 5, 6, 7])
In [4]: c=set([3,4,5,6])
In [5]: b & c
Out[5]: set([5, 6])
In [6]: b | c
Out[6]: set([3, 4, 5, 6, 7, 8, 9])
In [7]: b ^ c
Out[7]: set([3, 4, 7, 8, 9])
```

The operations are: Union | (or); Intersection & (and); symmetric difference (elements in set b but not in c and in c but not in b), ^; and so forth.

You can concatenate similar collection elements with +

```
In [1]: a = [1, 2, 3]
In [2]: b = [4, 5]
In [3]: a + b
Out[3]: [1, 2, 3, 4, 5]
In [4]: a = (1, 2)
In [5]: b = (4, 6)
In [6]: a + b
Out[6]: (1, 2, 4, 6)
In [7]: z1 = \{1: "AAA", 2: "BBB"\}
In [8]: z2 = {3: "CCC", 4: "DDD"}
In [9]: z1 + z2
     TypeError Traceback (most recent call last)
     ----> 1 71 + 72
TypeError: unsupported operand type(s) for +: "dict" and
```

Python programming best practices

We will try to instill as many standard practices as we can at the beginning.

- Wrap lines so that they are less than 80 characters long. You can use parentheses () or signal that the line continues using a "backslash".
- Use 4 spaces for indentation, no tabs.
- Separate functions using a blank line.
- When possible, write comments on separate lines.
- Use docstrings to document how to use the code, and comments to explain why and how the code works.

best practices continued

- Naming conventions:
- _internal_global_variable
- a_variable
- SOME_CONSTANT
- a_function
- Nevercallavariablel 1 or O or o
- use two or more letters for variables: xx better than x
- Use spaces around operators and after commas:
 a = func(x, y) + other(3, 4).