

# EEB 177 Lecture 9

Tuesday Feb 9th, 2017

## Topics

- end of collections...
- for, if, else, while statements

# Office Hours

Tues 11-12

Wednesday 10-11

Terasaki 2149 Today

# Hacking Sessions

Tuesday from 6-8:30 in LS 3209

# Homework #4

Due Wednesday by 5:00 PM

# Projects Deadline #2

Due Friday by 5:00 PM. To complete this assignment:

- Add a README to your Git repository
- Add two files of pseudocode for project. One should describe file input-output. The other should describe data manipulation over a loop.
- due Friday Week 5: Feb 10th  
(4%)

# Pseudocode

A way to plan code without worrying about syntax

Example: Write a program that asks the user for a temperature in Fahrenheit and prints out the same temperature in Celsius.

Pseudocode

```
xx = Get user input  
yy = Convert xx to Celsius  
Output message displaying Celsius temperature
```

python

```
xx = input("What is the temperature in Fahrenheit?")  
yy = (float(xx) - 32) * 5 / 9  
print("The temperature in Celsius is {}".format(yy))
```

Here is another example showing a loop.

Write a program in which a password is set and the program will keep prompting the user to guess it, until they get the word right.

Pseudocode

```
set password
ask user to try to guess password
    if the password is wrong, say access denied
    if the password is right, grant access
```

python

```
password = ("sesame")
hello = input("Enter the password: ")
while hello != password:
    print ("Access denied")
    hello = input("Enter the password: ")
if hello == password:
    print ("Login successful!")
```



# Preliminaries

- Create a new jupyter notebook and save the file "classwork-Tuesday-2-7.ipynb" to your class-assignments directory
- push this to your remote repository
- you can write answers to today's exercises in this file.

# Challenge

Do this:

- Define a list `a = [1, 1, 2, 3, 5, 8]`.
- Extract `[5, 8]` in two different ways.
- Add the element 13 at the end of the list.
- Reverse the list.
- Define a dictionary `m = {"a": ".-", "b": "-...-", "c": '-.-.'}`.
- Add the element `"d": "-.."`
- Update the value `"b": "-..."`

# Dictionaries

Dictionaries are collections of unordered lists where elements (*values*) are accessed by *keys*. We separate *key-value* pairs using a comma. Dictionaries are defined using `{}`

```
# create an empty dictionary
my_dict = {}
# strings, floats or even lists can be values in a dictionary
my_dict = {"a": "test", "b": 3.14, "c": [1, 2, 3, 4]}
GenomeSize = {"Homo sapiens": 3200.0, "Escherichia coli": 4.6}
# A dictionary has no natural order (i.e., the order of keys is not guaranteed)
GenomeSize

{'Arabidopsis thaliana': 157.0,
 'Escherichia coli': 4.6,
 'Homo sapiens': 3200.0}
```

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

```
In [3]: GenomeSize["Arabidopsis thaliana"]  
Out[3]: 157.0
```

You can also add entries by supplying a new key-value pair.

```
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}
```

## copy

```
In [13]: GS = GenomeSize.copy()
```

```
In [14]: GS
```

```
Out[14]:
```

```
{'Arabidopsis thaliana': 157.0, 'Escherichia coli': 4.6,  
'Homo sapiens': 3201.1, 'Saccharomyces cerevisiae': 12.1}
```

# clear

removes all elements

```
In [15]: GenomeSize.clear()  
In [16]: GenomeSize  
Out[16]: {}
```

# get

gets a value from key

```
In [67]: GenomeSize.get("Homo sapiens")  
Out[67]: 3200.0
```

this function is very useful for initializing the dictionary, or to return a special value when the key is not present.

```
In [68]: GenomeSize.get("Mus musculus", 10)  
Out[68]: 10
```



# items

returns key value pairs. Can be used to print contents of a dictionary.

```
for k,v in GS.items():  
    print(k, v)  
( 'Homo sapiens', 3200.0)  
( 'Escherichia coli', 4.6)  
( 'Arabidopsis thaliana', 157.0)
```

# keys, values

These functions return lists of the keys or values of the dictionary.

```
In [72]: GS.keys()
```

```
Out[72]: ['Homo sapiens', 'Escherichia coli', 'Arabidopsis']
```

```
In [74]: GS.values()
```

```
Out[74]: [3200.0, 4.6, 157.0]
```

# Creating dictionaries

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

```
enzymes = {}  
enzymes['EcoRI'] = r'GAATTC' # r before the string  
#tells python to automatically escape every character  
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 z1 + z2
```

```
TypeError: unsupported operand type(s) for +: "dict" and "
```

# collections recap

```
# round brackets --> tuple
In [1]: type((1, 2))
Out[1]: tuple
# square brackets --> list
In [2]: type([1, 2])
Out[2]: list
# curly brackets, sequence of values --> set
In [3]: type({1, 2})
Out[3]: set
# curly brackets, key-value pairs --> dictionary
In [4]: type({1: "a", 2: "b"})
Out[4]: dict
```



# for

for loops

imagine you wanted to print out each element of this list

```
apes = ["Homo", "Pan", "Gorilla"]
```

you could print these out seperately

```
print(apes[0] + " is an ape")  
print(apes[1] + " is an ape")  
print(apes[2] + " is an ape")
```

A better way is to use the for loop syntax

```
apes = ["Homo sapiens", "Pan troglodytes", "Gorilla gorilla"]  
for ape in apes:  
    print(ape + " is an ape")
```

for loops have the following structure:

```
for x in y:  
    do something
```

y is a list (or list-like object)

x is a variable names

the colon sets off an indented block of code (the body of the loop)

## a more complex loop

```
apes = ["Homo", "Pan", "Gorilla"]
for ape in apes:
    name_length = len(ape)
    first_letter = ape[0]
    print(ape + " is an ape. Its name starts with " + first_letter)
    print("Its name has " + str(name_length) + " letters")
```

# the range function

```
for number in range(6):  
    print(number)
```

## more on range

With two numbers, range will count up from the 1st number (inclusive) to the second (exclusive):

```
for number in range(3, 8):  
    print(number)
```

## range and step

With three numbers, range will count up from the 1st to the second with the step size given by the third:

```
for number in range(2, 14, 4):  
    print(number)
```

# while loops

a while loop runs until some condition is met.

```
count = 0
while count < 10:
    print(count)
    count = count + 1
```

## program flow in python

In its simplest form, a program is just a series of instructions (statements) that the computer executes one after the other. In Python, each statement occupies one line (i.e., it is terminated by a newline character). Other programming languages use special characters to terminate statements (e.g., ; is used in C).

Lets demonstrate statements that control flow in python with a simple program.



## conditional tests

A condition is simply a bit of code that can produce a true or false answer.

```
print(3 == 5)
print(3 > 5)
print(3 <= 5)
print(len("ATGC") > 5)
print("GAATTC".count("T") > 1)
print("ATGCTT".startswith("ATG"))
print("ATGCTT".endswith("TTT"))
print("ATGCTT".isupper())
print("ATGCTT".islower())
print("V" in ["V", "W", "L"])
```

we use conditional tests to control the flow of our program

```
expression_level = 125
if expression_level > 100:
    print("gene is highly expressed")
```

## if, elif, else

if and else create branching points in your program resulting in the execution of different blocks of code depending on a condition.

```
x=4
if x % 2 == 0:
    print("Divisible by 2") #body of loop
```

note indentation to designate loop body

We can use else to specify an action when a condition is not met.

```
x=4
if x % 2 == 0:
    print("Divisible by 2")
else:
    print("Not divisible by 2")
```

```
expression_level = 125
if expression_level > 100:
    print("gene is highly expressed")
else:
    print("gene is lowly expressed")
```

If we have multiple cases that need to be evaluated, elif is useful...

```
x = 17
if x % 2 == 0:
    print("Divisible by 2")
elif x % 3 == 0:
    print("Divisible by 3")
elif x % 5 == 0:
    print("Divisible by 5")
elif x % 7 == 0:
    print("Divisible by 7")
else:
    print("Not divisible by 2, 3, 5, 7")
```

## Challenge

change the program below so that it reports the temperature in Fahrenheit or Celsius depending on user input.

```
xx = input("What is the temperature in Fahrenheit?")  
yy = (float(xx) - 32) * 5 / 9  
print("The temperature in Celsius is {}".format(yy))
```

## if, elif, and else part II

we can handle complex flow with these statements.

```
file1 = open("one.txt", "w")
file2 = open("two.txt", "w")
accs = ['ab56', 'bh84', 'hv76', 'ay93', 'ap97', 'bd72']
for accession in accs:
    if accession.startswith('a'):
        file1.write(accession + "\n")
    else:
        file2.write(accession + "\n")
```



use and, or and not to specify complex conditionals

```
accs = ['ab56', 'bh84', 'hv76', 'ay93', 'ap97', 'bd72']
for accession in accs:
    if accession.startswith('a') and accession.endswith('6'):
        print(accession)
```

```
accs = ['ab56', 'bh84', 'hv76', 'ay93', 'ap97', 'bd72']
for accession in accs:
    if accession.startswith('a') or accession.startswith('b'):
        print(accession)
```

```
accs = ['ab56', 'bh84', 'hv76', 'ay93', 'ap97', 'bd72']
for acc in accs:
    if acc.startswith('a') and not acc.endswith('6'):
        print(acc)
```

```
file1 = open("one.txt", "w")
file2 = open("two.txt", "w")
file3 = open("three.txt", "w")
accs = ['ab56', 'bh84', 'hv76', 'ay93', 'ap97', 'bd72']
for accession in accs:
    if accession.startswith('a'):
        file1.write(accession + "\n")
    elif accession.startswith('b'):
        file2.write(accession + "\n")
    else:
        file3.write(accession + "\n")
```

# Reading and writing files

To work with text file contents you need to first open the file and then read the contents in as a string

```
# open the file
my_file = open("dna.txt")
# read the contents
my_dna = my_file.read()
# calculate the length
dna_length = len(my_dna)
# print the output
print("sequence is " + my_dna + " and length is " + str(dna_length))
```

