EEB 177 Lecture 8

Thursday Feb 8th, 2020

Preliminaries

- Create a new jupyter notebook and save the file "classwork-Thursday-2-6.ipynb" to your class-assignments directory
- push this to your remote repository

you can write answers to today's exercises in this file.

Topics

end of collections...

for, if, else, while statements

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 diction
my_dict = {"a": "test", "b": 3.14, "c": [1, 2, 3,
4]}
GenomeSize = {"Homo sapiens": 3200.0, "Escherichia coli": 4
A dictionary has no natural order (i.e., the order of key
GenomeSize

{'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
```

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}
```

```
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]: {}

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'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
```

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
```

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]: z_2 = \{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 "d

collections recap

```
# round brackets --> tuple
In [1]: type((1, 2))
Out[1]: tuple
# square brackets --> list
In [2]: type([1, 2])
Out<sup>[2]</sup>: 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 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) \times 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
    print("Its name has " + str(name_length) + " letters")
```

the range function

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

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

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

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)
```

a while loop runs until some condition is met.

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

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 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")
```

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

```
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:
```

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('3
 print(accession)

accs = ['ab56', 'bh84', 'hv76', 'ay93', 'ap97', 'bd72']
for accession in accs:

if accession.startswith('a') or accession.startswith('&
 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)
```