CSE 503 Introduction to Computer Science for Non-Majors

Dr. Eric Mikida epmikida@buffalo.edu 208 Capen Hall

Day 13 Dictionaries

Recap

- Mistakes will happen and things will go wrong
- How do we fix it? First we need to figure out what is going wrong.
 - We can use output (print or console.log) to give more information about what is happening during execution.
 - Checking that the output matches with our expectations can help reveal where things have gone wrong.
 - Asserts can also be used to automate the checking, and error messages can give more details on why a failure occurs.

DNA Frequency Exercise from Last Time

Write a function called **dnaFrequency** that takes a single DNA string, and returns a list of 4 lists, one for each base and its count.

For example:

dnaFrequency("ACAGCCTAAG") must return
[["A", 4],["C", 3], ["G",2],["T",1]]
dnaFrequency("TCAGCCTAAG") must return
[["A", 3],["C", 3], ["G",2],["T",2]]

DNA Frequency

1.	<pre>def dnaFrequency(string):</pre>
2.	bases = "ACGT"
3.	1 = []
4.	for b in bases:
5.	<pre>l.append([b, dnaCount(string, b)])</pre>
6.	
7.	return l

- What order do we put the lists in? Why?
 - Now we have to remember this order...

- What order do we put the lists in? Why?
 - Now we have to remember this order...
- How would we access the count for "A", for example?
 - **f** = dnaFrequency("AACTACGGCT")
 - **f[0][1]**
 - That is awkward. What ties that to "A"?
 - What if the order changes?

- What order do we put the lists in? Why?
 - Now we have to remember this order...
- How would we access the count for "A", for example?
 - **f** = dnaFrequency("AACTACGGCT")
 - **f[0][1]**
 - That is awkward. What ties that to "A"?
 - What if the order changes?
- How would we prefer to access the data?

Ordered vs Associative

- So far the collections we've seen (lists and arrays) have been ordered
 - They store a collection of values in a specific order
 - We access elements by their position in the list
 - ie a[0], a[3], a[147]
- Associative collections are different type of collection we can use in both Python and JavaScript
 - These collections associate a *key* with a *value* (called a <key, value> pair)
 - We access elements by their key

Key-Value Pairs in Real Life

<"First Name":"Eric"> <"Occupation":"Lecturer">

<"Siblings":3>

<"UBIT":"epmikida">

<"Last Name":"Mikida">

<"Favorite Number":2>

Key-Value Pairs in DNA Example





<"T": 9>



Python: Dictionary

- In Python, a key-value mapping is called a Dictionary
 - Dictionaries are indexed by *key* (instead of by a position)
 - A dictionary consists of a collection of key:value pairs, with the requirement that keys are unique
 - Strings can be keys, but so can any other value

Python Dictionary

The delimiters used to specific dictionaries are curly braces { }

Python Dictionary

The delimiters used to specific dictionaries are curly braces { } An empty dictionary can be created with a set of braces:

 $d1 = \{\}$

Python Dictionary: Creation

The delimiters used to specific dictionaries are curly braces { } An empty dictionary can be created with a set of braces:

d1 = {}

A dictionary can be given initial key:value pairs by giving it a comma separated list of key:value pairs inside the braces. This is also how dictionaries are printed as output.

d2 = { 'A':6, 'C':3, 'G':1, 'T':2}

Square brackets can be used to add/update/access individual items:

d = {"name":"Eric"}
d["age"] = 32 # Brackets can add a key:value pair
d["age"] = 29 # They can also update an existing pair
print(d["age"]) # ...or just to access a value

Square brackets can be used to add/update/access individual items:

```
d = {"name":"Eric"}
d["age"] = 32 # Brackets can add a key:value pair
d["age"] = 29 # They can also update an existing pair
print(d["age"]) # ...or just to access a value
```

The update function can be used to add/update from another dictionary
 d.update({"age":50, "job":"Lecturer"})

The get function provides a different way to access values

Behave the same if the key exists
print(d["name"]) # Prints "Eric"
print(d.get("name")) # Prints "Eric"

The get function provides a different way to access values

```
# Behave the same if the key exists
print(d["name"])  # Prints "Eric"
print(d.get("name")) # Prints "Eric"
```

```
# Behave different when the key does not exist
print(d["salary"])  # Error! Key not in dictionary
print(d.get("salary"))  # No error, no return value
print(d.get("salary", False)) # Returns false
```

Python Dictionary: Removal

Items can be removed with the **del** keyword, or **pop** function

del d["age"] # Removes "age", returns nothing
d.pop("job") # Removes "job", returns its value
print(d) # Now d is just {"name":"Eric"}

Membership can be tested with in and not in

"name" in d # Would evaluate to True
"age" in d # Would evaluate to False (age was just removed)
"job" not in d # Would evaluate to True

Python Dictionary: Keys, Values, Items

Dictionaries provide access to sequences for keys, values, and pairs
 d = {"Manager":"Sally", "Cashier":"Bob", "Security":"Joel"}
 for k in d.keys(): # Will print out "Manager", "Cashier", etc...
 print(k)

for v in d.values(): # Will print out "Sally", "Bob", "Joel"
 print(v)

for x in d.items(): # Will print out ("Manager", "Sally"), etc...
print(x)

DNA Frequency Revisited

Write a function called **dnaFrequency** that takes a single DNA string, and returns a *dictionary* containing the frequency of each base.

For example:

dnaFrequency("ACAGCCTAAG") must return
{"A":4,"C":3,"G":2,"T":1}

How does this compare to the list version?