

## Dictionaries and Functions

### Lists and Array Reminders

To create a list of items, use the `[]`

```
genes = ['SOD1', 'CDC11', 'YFG1']
print(genes)
print(genes[1])
print(genes[1:]) # everything after slot 1 (incl 1)
print(genes[:1]) # everything before slot 1
print(len(genes))

['SOD1', 'CDC11', 'YFG1']
CDC11
['CDC11', 'YFG1']
['SOD1']
3
```

Can also use negative numbers to start count from the back.

```
>>> print(genes[-1])
YFG
```

### Sets are unordered non-redundant collections of data.

Python also includes a data type for sets. A set is an unordered collection with no duplicate elements. Basic uses include membership testing and eliminating duplicate entries. Set objects also support mathematical operations like union, intersection, difference, and symmetric difference.

```
>>> basket = {'apple', 'orange', 'apple', 'pear', 'orange', 'banana'}
>>> print(basket) # show that duplicates have been removed
{'orange', 'banana', 'pear', 'apple'}
>>> 'orange' in basket # fast membership testing
True
>>> 'crabgrass' in basket
False

>>> # Demonstrate set operations on unique letters from two words
...
>>> a = set('abracadabra')
>>> b = set('alacazam')
>>> a # unique letters in a
{'a', 'r', 'b', 'c', 'd'}
>>> a - b # letters in a but not in b
{'r', 'd', 'b'}
>>> a | b # letters in a or b or both
{'a', 'c', 'r', 'd', 'b', 'm', 'z', 'l'}
```

```
>>> a & b                                # letters in both a and b
{'a', 'c'}
>>> a ^ b                                # letters in a or b but not both
{'r', 'd', 'b', 'm', 'z', 'l'}

>>> a = {x for x in 'abracadabra' if x not in 'abc'}
>>> a
{'r', 'd'}
```

## Some built-in list functions

- `range()` - `range(start, stop[, step])`

```
>>> range(5,10,1)
[5, 6, 7, 8, 9]
>>> range(5,-1,-1)
[5, 4, 3, 2, 1, 0]
```

- `map()` - lets you update a list with a function

```
l = [ 'a', 100, 12/3.3 ]
# " ".join(l) # this throws an error
# " ".join(map(str,l)) # have to cast numbers as string
print( " ".join(map(str,l)) )
l = [1,2,3,4]
squares = map(lambda x: x**2,l)
print(squares)

['a', 100, 3.6363636363636367]
[1, 2, 3, 4]
[1, 4, 9, 16]
```

## Reverse a list

- `reversed()` - iterate in reverse order of an array/string

```
l = ['zzz', 'yyy', 'a']
print(list(reversed(l)))
for n in reversed(l):
    print(n)

['a', 'yyy', 'zzz']
a
yyy
zzz
```

## More array functions

See more details here <https://docs.python.org/3/tutorial/datastructures.html>

- `list.append(x)` - Add an item to the end of the list;
- `list.pop([i])` - Remove the item at the given position in the list;
- `list.extend(L)` - Extend the list by appending all the items in the given list;
- `list.insert(i, x)` - Insert an item at a given position. The first argument is the index of the element before which to insert, so `a.insert(0, x)` inserts at the front of the list, and `a.insert(len(a), x)` is equivalent to `a.append(x)`.
- `list.remove(x)` - Remove the first item from the list whose value is `x`. It is an error if there is no such item. list, and return it. If no index is specified, `a.pop()` removes and returns the last item in the list.
- `list.index(x)` - Return the index in the list of the first item whose value is `x`. It is an error if there is no such item.
- `list.count(x)` - Return the number of times `x` appears in the list.
- `list.sort(cmp=None, key=None, reverse=False)` - Sort the items of the list in place
- `list.reverse()` - Reverse the order of the items in the list

## Sorting Lists

The `LIST.sort()` function on a list or the `sorted(LIST)` <https://docs.python.org/3/howto/sorting.html>

```
#!/usr/bin/env python3
genes = ['SOD1', 'CDC11', 'YFG1']
print(genes)
sort_genes = sorted(genes)
print(sort_genes)
numbers = [141, 7, 90, 3, 13]
print("unsorted", numbers)
numbers.sort()
print("sorted", numbers)
print("reversed", sorted(numbers, reverse=True))

alphanumbers = ['141', '7', '90', '3', '13']
print("Alphanumeric strings", alphanumbers)
print("Alpha sorted numbers", sorted(alphanumbers))
print("Numeric sorted", sorted(alphanumbers, key=int))
```

## Dates and times

See <https://docs.python.org/3/library/datetime.html>

```
from datetime import datetime

dates = ['3-Jan-2016', '4-Mar-2015', '2-Aug-1999', '1-May-2000']
print(dates)
dates.sort()
```

```

print(dates)

#newdates = [ datetime.strptime(d,"%d-%b-%Y") for d in dates ]
newdates = []
for str in dates:
    newdates.append(datetime.strptime(str,'%d-%b-%Y'))
print(newdates)
newdates.sort()
print(newdates)

for n in newdates:
    print(datetime.strftime(n,"%Y-%b-%d")," OR ",
        datetime.strftime(n,"%Y-%m-%d"), " OR ",
        datetime.strftime(n,"%A, %b %d, %Y"), " OR ",
        datetime.strftime(n,"%c")
    )

```

Iterate on Strings/Arrays in the same way

```

lst = [ 'BRCA1','SOD1','PTEN']
for gene in sorted(lst):
    print("gene is",gene)

```

```

DNA='AAAACCGTAG'
for let in DNA:
    print(let)

```

```

for let in reversed(DNA):
    print(let)

```

```

BRCA1
PTEN
SOD1

```

```

A
A
A
...

```

```

G
A
T
...

```

## Dictionaries

Dictionaries allow storing of data associated with a key instead of as an ordered list.

Initialize a dictionary, Dictionaries are key and value pairs

```
things = {}          # an empty dictionary
listofstuff = []     # an empty array
print(things)
things = {'diane': 10, 'jack': 13}
print(things)
things['diane']
things['billy'] = 15 # assign a new key/value pair
# if you have a list of pairs of things
strangerthings = dict([('Will', 12), ('Jim', 44), ('Joyce', 45), ('Eleven',11),('Lucas',10)])
strangerthings['Eleven']

{}
{'diane': 10, 'jack': 13}
10
11
```

## Iterate through a dictionary

Using the for loop and the items() function

```
for key,value in strangerthings.items():
    print("key is", key,"value is",value)

key is Will value is 12
key is Jim value is 44
key is Joyce value is 45
key is Eleven value is 11
key is Lucas value is 10
```

## To test if a key is in a dictionary

It is often the case you make a dictionary of values and you have another list and you want to cross-reference it. But maybe not all the values are in your dictionary.

```
set1 = {'a': 'apple', 'b': 'bear'}
if 'a' in set1:
    print("a is in set1")

if 'z' in set1:
    print("z is in set1")
```

## Functions

These are blocks of code that can be called repeatedly. Simplify tool development.

Might have subroutine to read a sequence file. Or compute a statistic.

Uses indentation just like loops.

```
def ROUTINENAME(ARGUMENTS):  
    CODE HERE
```

These can be used to run a routine that you might do repeatedly.

```
def average(list):  
    count=0  
    sum = 0.0  
    for item in list:  
        count += 1  
        sum += item  
    return sum / count  
  
print(average([100,200,300,150,110,99]))
```

## Read Fasta code part 1

<https://drj11.wordpress.com/2010/02/22/python-getting-fasta-with-itertools-groupby/>

See [https://github.com/biodataprogram/code\\_templates/blob/master/Lists\\_Dictionaries/fasta\\_parser.py](https://github.com/biodataprogram/code_templates/blob/master/Lists_Dictionaries/fasta_parser.py)

```
import itertools  
import sys  
import re  
  
# based on post here  
# https://drj11.wordpress.com/2010/02/22/python-getting-fasta-with-itertools-groupby/  
  
# define what a header looks like in FASTA format  
def isheader(line):  
    return line[0] == '>  
  
  
# this function reads in fasta file and returns pairs of data  
# where the first item is the ID and the second is the sequence  
# it isn't that efficient as it reads it all into memory  
# but this is good enough for our project  
def aspairs(filename):  
    seq_id = ''
```

```

sequence = ''
with open(filename,"r") as f:
    for header,group in itertools.groupby(f, isheader):
        if header:
            line = next(group)
            seq_id = line[1:].split()[0]
        else:
            sequence = ''.join(line.strip() for line in group)
            yield seq_id, sequence

# here is my program
# get the filename from the cmdline
seqfilename = sys.argv[1]
seqs = dict(aspairs(seqfilename))

# iterate through the sequences
n=0
for k,v in seqs.items():
    print( "id is ",k,"seq is",v)
    n += 1

print(n,"sequences")

id is  Q0142 seq is MTGSGTPPSREVNTYYMTMTMTMTMIMIMTMTMNIHFNNNNNNNNINMNSRRMYLFIL*M
id is  Q0143 seq is MGLWISFGTPPSYTYLLIMNHKLLLINNNNLTEVHTYFNININIDKMYIH*

```

## Dictionaries For Unique Lists

Dictionaries are useful ways to generate a unique list

```

dna = 'AAGAGAGGATACA'
bases = {'A':0, 'C':0, 'G':0, 'T':0 }
for l in dna:
    bases[l] += 1

print(bases)

{'A': 7, 'C': 1, 'G': 4, 'T': 1}

```