



PyQB

Monga

Dictionaries

Sets

Comprehensions

Files

Exercises

# Programming in Python<sup>1</sup>

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## Lecture VI: Dictionaries, sets, comprehensions



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Exercises

- Students: 27
- One triangle: tried by 22, 16 correct solutions
- Triangle kinds: tried by 20, 9 correct solutions
- DNA Hamming: tried by 19, 12 correct solutions
- Newton Sqrt: tried by 17, 13 correct solutions
- 7 students did all the exercises correctly



# Dictionaries

A composite type `dict` that implements a `mapping` between immutable `keys` and `values`.

```
d = {'key': 'foo', 3: 'bar'}
```

```
print(d['key']) # 'foo'  
print(d[3])     # 'bar'  
print(d[2])     # error!
```

Notation is similar to lists/tuples, but `dicts` are not sequences indexed by numbers, you must use only the existing keys (`d.keys()`).

```
if x in d.keys():  
    print(d[x])
```

A sequence of values can be obtained with `d.values`. A sequence of 2-tuples (key, value) with `d.items()`.

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A `set` is a composite object with no duplicate (non mutable) elements. Common set operations are possible.

- Set literals: `{1,2,3}` `set()`
- `{1,2,3}.union({3,5,6})`  
`{1,2,3}.intersection({3,5,6})`



# Comprehensions

**Comprehensions** are a concise way to create lists, sets, maps... It resembles the mathematical notation used for sets

$$A = \{a^2 | a \in \mathbb{N}\}.$$

```
squares = [x**2 for x in range(10)]
```

*# equivalent to:*

```
squares = []  
for x in range(10):  
    squares.append(x**2)
```

*# filtering is possible*

```
odds = [x for x in range(100) if x % 2 != 0]
```

*# with a set*

```
s = {x for x in range(50+1) if x % 5 == 0}
```

*# with a dict*

```
d = {x: x**2 for x in range(10)}
```

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A **file** is an abstraction the operating system uses to preserve data among the execution of programs. Data must be accessed **sequentially**.

- We need commands to ask to the OS to give access to a file (**open**).
- It is easy to read or write data **sequentially**, otherwise you need special commands (**seek**) to move the file “cursor”
- The number of open files is limited ( $\approx$  thousands), thus it is better to **close** files when they are not in use

Files contain bits (normally considered by group of bytes, 8 bits), the interpretation (“format”) is given by the programs which manipulate them. However, “lines of printable characters” (**plain text**) is a rather universal/predefined interpretation, normally the easiest to program.

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# File read access

```
f = open('filename.txt', 'r') # read only

# iterating on a file reads (all) the lines
for i in f:
    print(i)

# End of file already reached, result is ''
f.readline()

f.close()

# File closed, error!
f.readline()
```

To avoid remembering to close explicitly, Python provides the **context manager** syntax.

```
with open('filename.txt', 'r') as f:
    for i in f:
        print(i)
```

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Exercises

- Write a function to compute the complement of a DNA strand: every A becomes a T, every T an A, every C an G, every G an C.
- Apply the function to every line of a file with a DNA sequence
- Write a function that gives the set of (unique) sequences of 10 nucleic acids in a file



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Exercises

- <https://classroom.github.com/a/MhchQHAd>
- <https://classroom.github.com/a/36ITXw1V>