

Day 4

# Last session

- Sorting
- Reading and writing files
- Using *import* to import modules
- Parsing genomic regions
- Obtaining sequences and reverse complement using *ucscgenome* and *str.translate()*.

# This session

- Dictionaries
- Functions
- Scripting
- Rosalind

# Dictionaries

- Dictionaries are *collections*, just like lists but...
- They are indexed by values (*keys*) other than numbers
- They are unordered

# Dictionaries

```
# an empty dictionary
```

```
a = {}
```

```
print(a)
```

```
# a dictionary, indexed by strings
```

```
a = {"one": 1, "two": 2, "three": 3}
```

```
print(a["two"])
```

```
# does a key exist? (the `in` operator)
```

```
print("one" in a)
```

```
print("four" in a)
```

```
# or like this
```

```
if "one" in a:
```

```
    print("a contains 'one'")
```

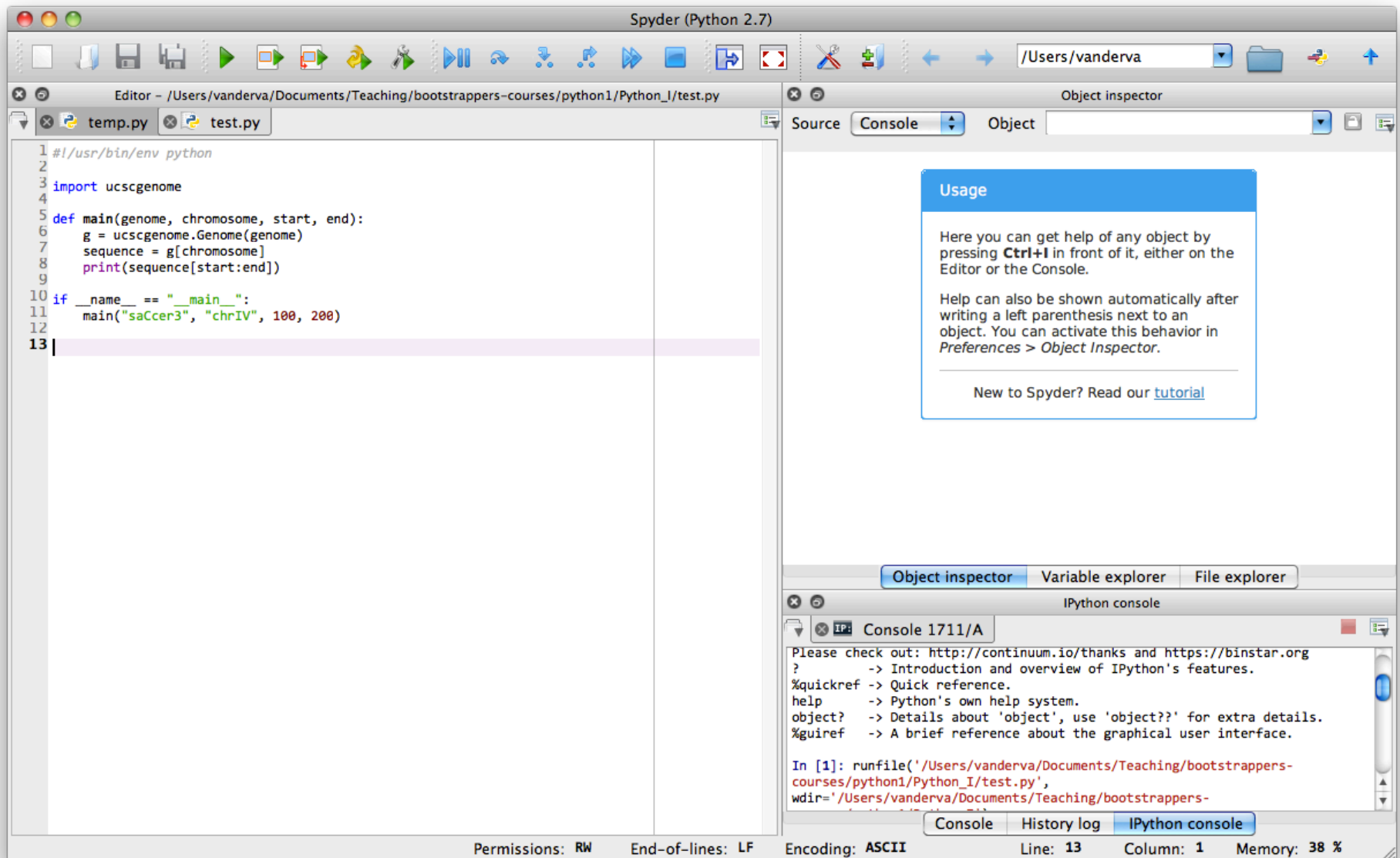
# Functions

- Functions are “predefined blocks of code” that accept parameters and return values
- We have used quite a few functions already.

```
def area(length, width):  
    return length * width
```

```
def mean(numbers): # this defines a function `mean()``  
    mysum = sum(numbers)  
    return float(mysum) / len(numbers)
```

# Spyder IDE



# Scripting

```
#!/usr/bin/env python
```

```
def area(length, width):  
    return length * width
```

```
def mean(numbers): # this defines a function `mean()``  
    mysum = sum(numbers)  
    return float(mysum) / len(numbers)
```

```
def hello():  
    print("Hello World")
```

```
def main(): # this defines a function `main()``  
    x = [1, 2, 3, 4, 5, 6, 7, 8, 9, 10]  
    print(mean(x))  
    print(type(mean(x)))  
    myvariable = hello()  
    print(type(myvariable))  
    print(area(10, 15))
```

```
if __name__ == "__main__":  
    main()
```