

# Exercise 1: Lambda function

1. First think what this code does, try to guess the output and then try the code out to check if you were right!

```
z = range(2, 7)  
map(lambda x: x**2, z)
```

2. Use filter and lambda function to show the even values in the list z

# Exercise 2: Regular expressions

*Preparation step:*

- a) Get the file '**nuc\_sequence.txt**' from the lecture web page.
- b) Save the nucleotide sequence of the previous file in a variable called `s`.  
Hint: You can use `readlines()` function.  
Please note that the output of this function is list, so you need to convert it to string.
- c) Change every letter to upper case.  
Hint: use `upper()` function
- d) Remove newlines (`\n`) characters.  
Hint: use `replace()` function

*Now we can query our gene with regular expressions:*

- 1. Find all the substrings that contain just three 'A' or more: e.g. AAA or AAAAAA.  
Hint: you can use `re.finditer (RE, string)` function to iterate over the string.
- 2. To show the results you can use `.group()` and `.span()` functions.  
Find all the substrings that start with a 'T' and then they have two or three 'C' and then they end with a 'G'.