

# BC205: Algorithms for Bioinformatics.

## Exercise 1. Introduction and Sequence Analysis

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### 1. Finding palindromes in a sequence using recursion

The function **palindrome\_subsequence** takes a sequence string as an input.

First, a list comprehension is created which generates all the possible 3-mer substrings that are represented in the input sequence. Inside the list comprehension, the conditional `if`, checks if each of these substrings generated in the loop, are equal to its reverse.

If a palindrome substring with length  $> 3$  is found successfully, it is added to a list called **pal\_list**.

If **pal\_list** is not empty, which means that at least one palindrome substring is found, then the function returns **True**, and the palindrome with the maximum length.

Else, if **pal\_list** is empty, no palindromes of length  $> 3$  were found in the input sequence, and the function will return **False**.

In order to cross check the results, we can try different inputs of small sequences and check whether the results are the expected.

```
# create a function that will take a sequence string as input and return
# True or False depending on whether it contains a palindrome substring and if
# so, will also return the longest substring
```

```
def palindrome_subsequence(sequence):
    #scan the sequence string with step = 2 to exclude 2-mers
    pal_list = [sequence[i:j+1] for i in range(len(sequence))
                for j in range(i+2, len(sequence))

                if sequence[i:j+1] == sequence[i:j+1][::-1]]

    # if palindrome substrings were found, return True and the biggest one
    if pal_list:
        p = max(pal_list)
        return True, p
    else:
        # if pal_list is empty, sequence has no palindrome substrings, return False

        return False
```

```
# check for palindromes in the sequence 'TTCAGGTGG'
```

```
print(palindrome_subsequence('TTCAGGTGG'))          # output True, GGTGG
```

```
#check for palindromes in the sequence 'CTGTTATTAATTATTGCAT'
```

```
print(palindrome_subsequence('CTGTTATTAATTATTGCAT')) # ouput TRUE TTATTAATTATT
```

```
#check for palindromes in the sequence 'GGCATCGGATTCGT'
```

```
print(palindrome_subsequence('GGCATCGGATTCGT'))    # output False
```

The output of the given examples, is as we would expect:

- TTCAGGTGG : As we can see, this sequence contains these palindromes > 3 nt GGTGG, 'GTG'
- TTCACGTAAGTGG : Contains the following substrings TGT, 'TAAT', 'ATTA', 'GTTATTAATTATTG', 'TTATTAATTATT', 'TATTAATTAT', 'TAT', 'TTAATT', 'TTATT', 'ATTAATTA'.  
In this sequences there are also many 2nt long palindrome substrings but they were excluded.
- GGCATCGGATTCGT : This sequence does not contain any palindrome subsequence with length bigger than 2, thus the output is False, as expected.

