



Exercise

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Factorial

the factorial of a positive integer, denoted by $n!$, is the product of all positive integers less than or equal to n :

$$n! = n \times (n - 1) \times (n - 2) \times (n - 3) \times \dots \times \mathbf{3} \times \mathbf{2} \times \mathbf{1}.$$

Recursion

method of solving a problem where the solution depends on solutions to smaller instances of the same problem

In python you could implement it with a function that calls itself.

Recursive Factorial

```
def factorial_rec(n):  
    if n == 0:  
        return 1  
    else:  
        return n * factorial_rec(n-1)
```

Iterative Factorial

```
def factorial(n):  
    for i in range((n-1), 0, -1):  
        n = n * i  
    return n
```

Fibonacci Numbers

the Fibonacci numbers, commonly denoted F_n form a sequence, called the Fibonacci sequence, such that each number is the sum of the two preceding ones, starting from 0 and 1:

$$F_0 = 0, F_1 = 1 \text{ and}$$
$$F_n = F_{n-1} + F_{n-2}$$

Recursive Fibonacci

```
def fibonacci_rec (n):  
    if n == 0 or n == 1:  
        return 1  
    else:  
        return fibonacci_rec(n-1) + fibonacci_rec(n-2)
```

Iterative Fibonacci

```
def fibonacci(n):  
    if n == 0:  
        return n  
    last = 0  
    next = 1  
    for _ in range(1, n+1):  
        previous_last = last  
        last = next  
        next = previous_last + last  
    return next
```


Pigreco - Leibniz Formula

$$\sum_{n=0}^{\infty} \frac{(-1)^n}{2n+1} = 1 - \frac{1}{3} + \frac{1}{5} - \frac{1}{7} + \dots + \frac{(-1)^n}{2n+1} + \dots = \frac{\pi}{4}.$$

Simplified version to calculate the convergence of π

$$\pi = 4/1 - 4/3 + 4/5 - 4/7 + 4/11 \dots$$

```
def calculate_pi(n_terms):  
    numerator= 4.0  
    denominator = 1.0  
    operation = 1.0  
    pi = 0.0  
    for _ in range(n_terms):  
        pi += operation * (numerator/denominator)  
        denominator += 2.0  
        operation *= -1.0  
    return pi
```

Searching

Linear Search

```
def linear_contains(iterable, key):  
    for item in iterable:  
        if item == key:  
            return True  
    return False
```

Binary search

only on sorted data

```
def binary_contains(sorted_iterable, key):
    low = 0
    high = len(sorted_iterable) - 1
    while low <= high:
        mid = (low + high) // 2
        if sorted_iterable[mid] < key:
            low = mid + 1
        elif sorted_iterable[mid] > key:
            low = mid - 1
        else:
            return True
    return False
```

Contacts

```
if __name__=="__main__":

    print('Contacts')
    print('-----')
    contacts={}
    your_choice= 0
    while your_choice != 5:
        menu()
        your_choice=int(input('Select your option: '))
        print('-----')
        if your_choice==1:
            add_contact(contacts)
        elif your_choice==2:
            remove_contact(contacts)
        elif your_choice==3:
            list_contacts(contacts)
        elif your_choice==4:
            search_contact(contacts)
```

```
def menu():  
    print('1 - add a contact')  
    print('2 - remove a contact')  
    print('3 - list all contacts')  
    print('4 - search a contact')  
    print('5 - exit')
```



```
def add_contact(dict):  
    contact=input('Insert name: ')  
    mobile_number=input('Mobile number: ')  
    dict[contact]=mobile_number  
    print('Contact %s with %s number added\n' % (contact,mobile_number))
```

```
def remove_contact(dict):  
    contact = input('Contact to remove: ')  
    if contact in dict.keys():  
        del dict[contact]  
        print('Contact removed')  
    else:  
        print('Contact %s not found' % (contact))
```

```
def list_contacts(dict):  
    if len(dict)==0:  
        print("Nothing here")  
    for x in dict.keys():  
        print("Contats: %s \tNumber: %s" % (x, dict[x]))
```

```
def search_contact(dict):
    contact = input('Insert contact to search: ')
    for name in dict.keys():
        if name == contact:
            print('Contact %s with %s number\n' % (name,dict[name]))
    print("Nothing here")
```

DNA

Deoxyribonucleic acid

Genes are commonly represented in computer software as a sequence of the characters cytosine [C], guanine [G], adenine [A] or thymine [T]. Each letter represents a *nucleotide*, and the combination of three nucleotides is called a *codon*.



```
Nucleotide = {'A': 'A', 'C': 'C', 'G': 'G', 'T': 'T'}  
gene_str = "ACGTGGCTCTCTAACGTACGTACGTACGGGGTTTATATATACCCTAGGACTCCCTTT"
```

```
def string_to_gene(s):  
    gene = []  
    for i in range(0, len(s), 3):  
        if (i + 2) >= len(s):  
            return gene  
        codon = (Nucleotide[s[i]], Nucleotide[s[i+1]], Nucleotide[s[i+2]])  
        gene.append(codon)  
    return gene
```

```
def linear_contains (gene, key_codon):  
    for codon in gene:  
        if codon == key_codon: # only for educational purpose  
            #else return (key_codon in gene) should be enough  
            return True  
    return False
```


binary search works only on sorted gene!

```
def binary_contains(gene, key_codon):  
    low = 0  
    high = len(gene) - 1  
    while low <= high:  
        mid = (low + high) // 2  
        if gene[mid] < key_codon:  
            low = mid + 1  
        elif gene[mid] > key_codon:  
            high = mid - 1  
        else:  
            return True  
    return False
```

```
if __name__ == '__main__':
    gene_str = "ACGTGGCTCTCTAACGTACGTACGTACGGGGTTTATATATACCCTAGGACTCCCTTT"
    my_gene = string_to_gene(gene_str)
    print(my_gene)
    acg = (Nucleotide['A'], Nucleotide['C'], Nucleotide['G'])
    gat = (Nucleotide['G'], Nucleotide['A'], Nucleotide['T'])

    print(linear_contains(my_gene, acg)) # True
    print(linear_contains(my_gene, gat)) # False

    sorted_gene = sorted(my_gene)
    print(binary_contains(sorted_gene, acg)) # True
    print(binary_contains(sorted_gene, gat)) # False
```