finger print

```
def solve_fingerprint(seq_list, no_of_col):
  seq_dict=dict()
  for column in range(no_of_col):
   counta,countc,countt,countg=0,0,0,0
   for colseq in seq_list:
    if colseq[column]=='A':
     counta+=1
    elif colseq[column]=='T':
     countt+=1
    elif colseq[column]=='C':
     countc+=1
    elif colseq[column]=='G':
     countg+=1
   seq_dict[column]=[counta,countc,countt,countg]
  display_results(seq_dict)
def display_results(seq_dict):
 print("\tA \tC \tT \tG")
 for key in seq_dict:
   print("\n",*seq_dict[key],sep="\t")
no_of_seq=int(input("Enter the number of sequence: "))
print("Enter all the sequences")
seq_list=[]
for _ in range(no_of_seq):
 seq_list.append(list(map(str, input("").split())))
solve_fingerprint(seq_list,len(seq_list[0]))
Enter the number of sequence: 4
Enter all the sequences
ACTGATG
ATCAGAA
ATAAGCA
AGTTAGC
ACTG
4000
0121
1120
2011
2002
1111
```

pairwise

2101

se2=input("Enter the second sequence::") seq1=list(se1) seq2=list(se2) score=[] def Pairwise_alignment(a,b): gap(a,b) print(a) print(b) value=0 length=len(a) for i in range(0,length): if(a[i]==b[i]): score.append('1') value=value+1 else: score.append('0') print(score) print(value) def gap(a,b): if(len(a)==len(b)):

se1=input("Enter the first sequence::")

In [1]:

In [2]:

```
print()
else:
k=int(input("enter the position to insert::"))
if (len(a)<len(b)):
    a.insert(k,'-')
else:
    b.insert(k,'-')
return(a,b)

Pairwise_alignment(seq1,seq2)

Enter the first sequence::abcdefgh
Enter the second sequence::abddef
enter the position to insert::2
['a', 'b', 'c', 'd', 'e', 'f', 'g', 'h']
['a', 'b', '-', 'd', 'c', 'd', 'e', 'f']
['1', '1', '0', '1', '0', '0', '0', '0']
3
```

Identity

```
se1=input("Enter the first sequence::")
 se2=input("Enter the second sequence::")
 seq1=list(se1)
seq2=list(se2)
 def find_identity(a,b):
   gap(a,b)
   print(a)
   print(b)
   score=0
   length=len(a)
   total_elements=len(a)*len(b)
   for i in range(0,length):
      for j in range(0,length):
         if(a[i]==b[j]):
           score=score+1
   identity=(score/total elements)*100
   print("Matching Score::",score)
   print("Identity of the sequences::",identity)
 def gap(a,b):
   if(len(a)==len(b)):
      print()
   else:
      k=int(input("enter the position to insert gap ::"))
      if (len(a)<len(b)):
         a.insert(k,'-')
      else:
         b.insert(k,'-')
   return(a,b)
    name == " main ":
   find identity(seq1, seq2)
Enter the first sequence::abcdefgh
Enter the second sequence::abdcdefg
['a', 'b', 'c', 'd', 'e', 'f', 'g', 'h']
['a', 'b', 'd', 'c', 'd', 'e', 'f', 'g']
Matching Score:: 8
Identity of the sequences:: 12.5
```

regualar expression

```
def gen_reg_exp(seq_list, no_of_col):
    final_list=[]
    for colnum in range(no_of_col):
        collist=[]
        for colseq in seq_list:
            collist.append(colseq[colnum])
        if len(set(collist))==len(collist):
            final_list.append('x')
        else:
            if len(set(collist))==1:
```

In [3]:

In [4]:

```
final_list.append(collist[0])
         final_list.append(".join(set(collist)))
   display_output(final_list)
def display_output(final_list):
   print(*final_list, sep='-')
if __name__ == '__main__':
   no_of_seq = int(input("Enter the number of sequence: "))
   print("Enter all the sequences")
   seq_list = []
   for _ in range(no_of_seq):
     seq\_list.append(list(map(str, input("").split())))
   gen_reg_exp(seq_list, len(seq_list[0]))
Enter the number of sequence: 4
Enter all the sequences
ADLGAVFALCDRYFQ
SDVGPRSCFCERFYQ
ADLGRTQLRCDRYYQ
ADIGQPHSLCERYFQ
AS-D-LIV-G-x-x-x-LRF-C-ED-R-FY-YF-Q
```

similarity

sequence_one=input("Enter the first sequence: ")

```
sequence_two=input("Enter the second sequence: ")
 how_many=int(input("How many elements for similarity condition?"))
 similarities=[]
 for i in range(0,how_many):
    a=input("Enter an element: ")
    c=int(input("How many elements is it similar to? "))
    similarities.append([])
    similarities[i].append(a)
    for j in range(0,c):
      b=input("What is it similar to? ")
      similarities[i].append(b)
 def compare(o,t,s):
  print(o)
  print(t)
  print(s)
  score=0
  for i in range(len(o)):
    for j in range(len(s)):
      if o[i] in s[j] and t[i] in s[j] and o[i] != t[i]:
         score+=1
  similarity= (score*100)/len(o)
  return similarity
 print(compare(list(sequence_one),list(sequence_two),similarities),"%")
Enter the first sequence: abcdefgh
Enter the second sequence: abdcdefg
How many elements for similarity condition?2
Enter an element: a
How many elements is it similar to? 1
What is it similar to? a
Enter an element: b
How many elements is it similar to? 2
What is it similar to? b
What is it similar to? d
['a', 'b', 'c', 'd', 'e', 'f', 'g', 'h']
['a', 'b', 'd', 'c', 'd', 'e', 'f', 'g']
[['a', 'a'], ['b', 'b', 'd']]
```

percentage matching

In [5]:

0.0 %